



## **Reference (3) Speech Materials**

Jeff (CHI-HSUAN HO)

**The subsequent slides only showcase a few parts of the work materials and results.**



# **Multi-Omics Factor Analysis (MOFA)**

## **A Framework for Unsupervised Integration of Multi-omics Data Sets**

Jeff (CHI-HSUAN HO)

## Methods

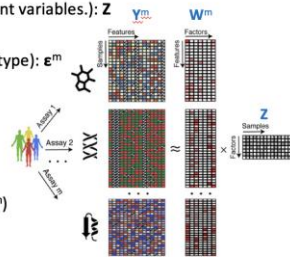
### • MOFA models

- Factor Analysis: Reduce the dimensionality of a (big) dataset into a small set of variables which are easier to interpret and visualize.
- Sample size:  $N$  ( $n$ )
- Feature numbers:  $D$  ( $d$ )
- Data matrices, view:  $M$  ( $m$ )
- Factor matrix (matrix that contains the low-dimensional latent variables.):  $Z$
- Loading matrix, weight matrix:  $W^m$
- Residual noise (its form depends on the specific of the data type):  $\epsilon^m$
- The expectation of  $x$  under the distribution  $q$ . ( $E_q[x]$ ):  $\langle x \rangle$

### • Model:

$$Y^m = ZW^m + \epsilon^m$$

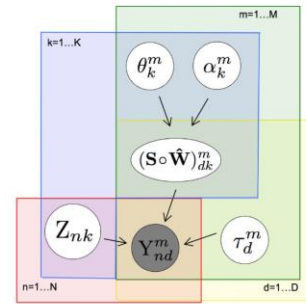
- A Bayesian framework is used. (Place prior dist. on  $Z$ ,  $W^m$ ,  $\epsilon^m$ )



## Methods

### • Likelihood $\times$ Prior (Gaussian data)

$$p(Y, \hat{W}, S, Z, \Theta, \alpha, \tau) = \prod_{m=1}^M \prod_{n=1}^N \prod_{d=1}^{D_m} \mathcal{N} \left( y_{nd}^m \mid \sum_{k=1}^K s_{dk}^m \hat{w}_{dk}^m z_{nk}, 1/\tau_d \right) \prod_{m=1}^M \prod_{d=1}^{D_m} \prod_{k=1}^K \mathcal{N}(\hat{w}_{dk}^m \mid 0, 1/\alpha_k^m) \text{Ber}(s_{dk}^m \mid \theta_k^m) \prod_{n=1}^N \prod_{k=1}^K \mathcal{N}(z_{nk} \mid 0, 1) \prod_{m=1}^M \prod_{k=1}^K \text{Beta}(\theta_k^m \mid a_0^m, b_0^m) \prod_{m=1}^M \prod_{k=1}^K \mathcal{G}(\alpha_k^m \mid a_0^m, b_0^m) \prod_{m=1}^M \prod_{d=1}^{D_m} \mathcal{G}(\tau_d^m \mid a_0^m, b_0^m)$$

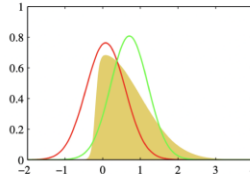


## Methods

### • Posterior – model Inference for Gaussian data: Variational Bayes

- Goal: In the set of the tractable distribution for  $q(X)$  (or  $q$ ), maximize  $\mathcal{L}(q)$  (or minimize  $KL(q||p)$ ) as possible so that  $q(X)$  (or  $q$ ) can provide a good approximation to  $p(Y|X)$ .

- Find the general form for the set of the tractable distribution for  $q(X)$  (or  $q$ ).



- Mean-field theory assumption

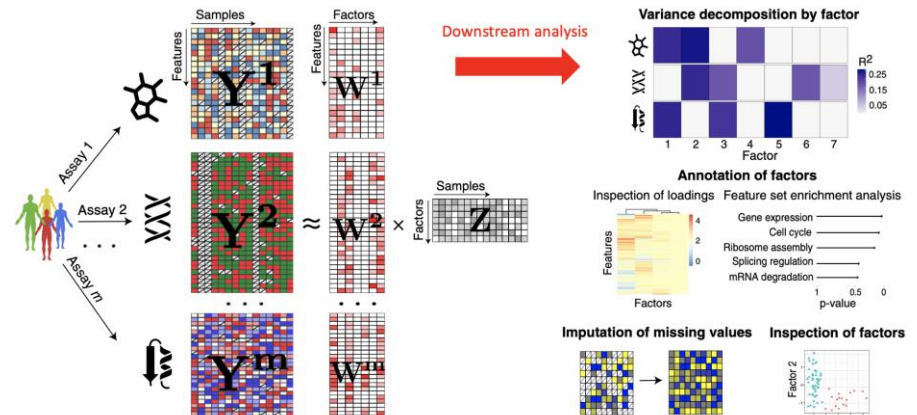
- Partition the elements of  $X$  into  $M$  disjoint groups, then  $q(X) = \prod_{i=1}^M q_i(x_i)$

- Maximize  $\mathcal{L}(q)$  with respect to each of the latent variables and parameters in turn.

- Dissect out the dependence on one of the latent variables and parameters  $q_j(X_j)$ . (Keep the  $\{q_{i \neq j}\}$  fixed, maximize  $\mathcal{L}(q)$  respect to  $q_j(X_j)$ ).

$$\mathcal{L}(q) = \int \prod_{i=1}^M q_i(x_i) \{ \ln p(Y, X) - \sum_{i=1}^M \ln q_i(x_i) \} dX = \int q_j \left\{ \int \ln p(Y, X) \prod_{i \neq j} q_i(x_i) dx_i \right\} dx_j - \int q_j \ln q_j dx_j + \text{const} = \int q_j \ln \tilde{p}(Y, x_j) dx_j - \int q_j \ln q_j dx_j + \text{const} = -KL(q_j(x_j) || \tilde{p}(Y, x_j)), \text{ where } \ln \tilde{p}(Y, x_j) = E_{q_{i \neq j}}[\ln p(Y, X)] + \text{const}.$$

## Methods



<https://github.com/jeff665547/SkillShare-MOFA/raw/master/MOFA.pdf>

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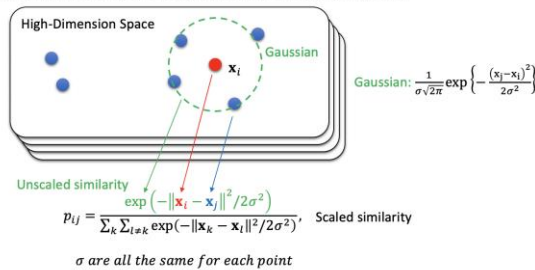


# **Dimension Reduction Techniques From t-SNE to UMAP**

Jeff (CHI-HSUAN HO)

## t-Distributed Stochastic Neighbor Embedding (t-SNE)

- Measure how close between two different high-dimensional objects.



- Similar (closer) points in the high-D space  $\rightarrow$  Large  $p_{ij}$ .

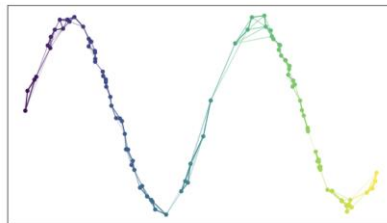
## t-Distributed Stochastic Neighbor Embedding (t-SNE)

- Gradient of Kullback-Leibler divergences:  $C = KL(P||Q) = \sum_i \sum_j p_{ij} \log \frac{p_{ij}}{q_{ij}}$ 
  - Let  $d_{ij} = \|y_i - y_j\|$ ,  $Z_i = \sum_l (1 + d_{il}^2)^{-1}$
  - $\frac{\partial C}{\partial y_i} = \sum_j \left( \frac{\partial C}{\partial d_{ij}} \cdot \frac{\partial d_{ij}}{\partial y_i} + \frac{\partial C}{\partial d_{ji}} \cdot \frac{\partial d_{ji}}{\partial y_i} \right) = \sum_j \left( \frac{\partial C}{\partial d_{ij}} + \frac{\partial C}{\partial d_{ji}} \right) \cdot \frac{y_i - y_j}{d_{ij}} = 2 \sum_j \frac{\partial C}{\partial d_{ij}} \cdot \frac{y_i - y_j}{d_{ij}}$
  - $\frac{\partial C}{\partial d_{ij}} = -\sum_{k \neq i} p_{ik} \frac{\partial}{\partial d_{ij}} \log(q_{ik}) = -\sum_{k \neq i} p_{ik} \frac{\partial}{\partial d_{ij}} (\log(q_{ik} Z_i) - \log Z_i)$ 
    - $= -\sum_{k \neq i} p_{ik} \left( \frac{1}{q_{ik} Z_i} \frac{\partial}{\partial d_{ij}} (1 + d_{ik}^2)^{-1} - \frac{1}{Z_i} \frac{\partial Z_i}{\partial d_{ij}} \right)$ ,  $\frac{\partial}{\partial d_{ij}} (1 + d_{ik}^2)^{-1}$ ,  $\frac{\partial}{\partial d_{ij}} Z_i$  only have value when  $k, l = j$
    - $= p_{ij} \left( \frac{1}{q_{ij} Z_i} 2d_{ij} (1 + d_{ij}^2)^{-2} \right) + \sum_{k \neq i} \frac{-p_{ik}}{Z_i} (1 + d_{ij}^2)^{-2} 2d_{ij}$
    - $= 2d_{ij} \left( p_{ij} \frac{(1 + d_{ij}^2)^{-2}}{q_{ij} Z_i} - \frac{(1 + d_{ij}^2)^{-2}}{Z_i} \sum_{k \neq i} p_{ik} \right) = 2d_{ij} (p_{ij} (1 + d_{ij}^2)^{-1} - q_{ij} (1 + d_{ij}^2)^{-1} \sum_{k \neq i} p_{ik})$
    - $= 2d_{ij} (p_{ij} (1 + d_{ij}^2)^{-1} - q_{ij} (1 + d_{ij}^2)^{-1}) = 2d_{ij} (p_{ij} - q_{ij}) (1 + d_{ij}^2)^{-1}$
  - $\frac{\partial C}{\partial y_i} = 4 \sum_j (p_{ij} - q_{ij}) \cdot (1 + \|y_i - y_j\|^2)^{-1} \cdot (y_i - y_j)$

## Uniform Manifold Approximation and Projection (UMAP)

A computation view of UMAP

- In the High-D space (manifold):



- Consider the symmetric matrix  $B = A + A^T - A \circ A^T$ , where  $\circ$  is the Hadamard (or pointwise) product.
- In a meaningful way, the element in  $B$  represents  $\mu_A(x) + \mu_B(x) - \mu_A(x) \cdot \mu_B(x)$
- $B$  is also the undirected weighted matrix.

## Uniform Manifold Approximation and Projection (UMAP)

- Gradient of Cross entropy

- Let  $d_{ij} = \|y_i - y_j\|_2$ 

$$v_{ij} = \frac{1}{(1 + ad_{ij}^{2b})}, \quad 1 - v_{ij} = \frac{ad_{ij}^{2b}}{(1 + ad_{ij}^{2b})}, \quad \frac{\partial v_{ij}}{\partial d_{ij}} = -\frac{2abd_{ij}^{2b-1}}{(1 + ad_{ij}^{2b})^2}$$

$$\begin{aligned} \frac{\partial C}{\partial y_i} &= \sum_j \left[ -\frac{w_{ij}}{v_{ij}} \frac{\partial v_{ij}}{\partial d_{ij}} + \frac{1 - w_{ij}}{1 - v_{ij}} \frac{\partial v_{ij}}{\partial d_{ij}} \right] \frac{\partial d_{ij}}{\partial y_i} \\ &= \sum_j \left[ \left( -w_{ij} (1 + ad_{ij}^{2b}) + \frac{(1 - w_{ij})(1 + ad_{ij}^{2b})}{ad_{ij}^{2b}} \right) \frac{\partial v_{ij}}{\partial d_{ij}} \right] \frac{\partial d_{ij}}{\partial y_i} \\ &= \sum_j \left[ \frac{2abd_{ij}^{2(b-1)}}{1 + ad_{ij}^{2b}} \cdot w_{ij} \cdot d_{ij} \cdot \frac{(y_i - y_j)}{d_{ij}} \right. \\ &\quad \left. - \frac{2b}{d_{ij}^2 (1 + ad_{ij}^{2b})} \cdot (1 - w_{ij}) \cdot d_{ij} \cdot \frac{(y_i - y_j)}{d_{ij}} \right] \end{aligned}$$

Attractive force

Repulsive force

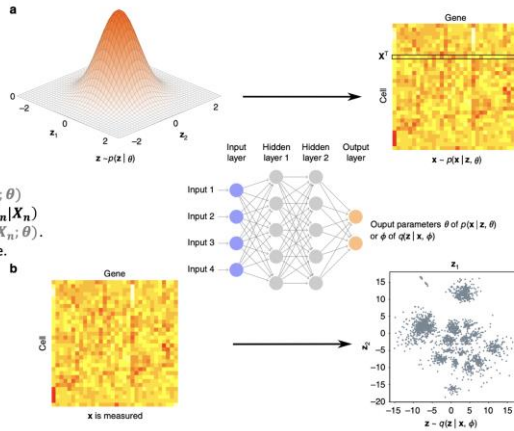
<https://github.com/jeff665547/SkillShare-UMAP-and-t-SNE/raw/master/From%20t-SNE%20to%20UMAP.pdf>



# **Interpretable Dimensionality Reduction of Single Cell Transcriptome Data with Deep Generative Models (SCVIS)**

Jeff (CHI-HSUAN HO)

## SCVIS



- Interest: Find  $P(Z_n|X_n; \theta)$   
→ Find an optimal  $q(Z_n|X_n)$   
to approximate  $P(Z_n|X_n; \theta)$ .  
→ Variational Inference.

## SCVIS

- Regularizer:

$$\sum_i \mathbb{KL}(p_{\cdot|i} || q_{\cdot|i}) = \sum_{i=1}^N \sum_{j=1, j \neq i}^N p_{ji} \log \frac{p_{ji}}{q_{ji}}$$

- Final objective function:

$$\arg \min_{\theta, \phi} \left( - \sum_{n=1}^N \text{ELBO}_n + \alpha \sum_{n=1}^N \mathbb{KL}(p_{\cdot|n} || q_{\cdot|n}) \right)$$

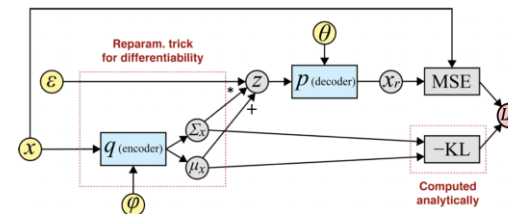
- $\alpha$  is set to the dimensionality of the input high-dimensional data.  
(ELBO scales with the dimensionality of the input data)

## Variational Autoencoder (VAE)

- Relationship with the Autoencoder

$$\begin{aligned} \text{Cost function: } L_b(q(Z_n|X_n; \phi), \theta) &= E_{z \sim q_\phi}[\log P(X_n|Z_n; \theta)] - \text{KL}[q(Z_n|X_n; \phi) || P(Z_n)] \\ &= E_{z \sim N(0,1)}[\log P(X_n|g(\varepsilon, X_n, \phi); \theta)] - \text{KL}[q(Z_n|X_n; \phi) || P(Z_n)] \end{aligned}$$

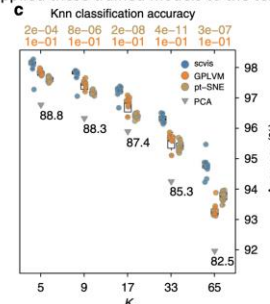
- $q(Z_n|X_n; \phi)$ : Encoder,  $P(X_n|g(\varepsilon, X_n, \phi); \theta)$ : Decoder,  $\mu_\phi(X_n) = \mu_x$ ,  $\sigma_\phi(X_n) = \Sigma_x$
- $E_{z \sim N(0,1)}[\log P(X_n|g(\varepsilon, X_n, \phi); \theta)]$ : MSE, (Reconstruction Error), it based on the encoder and the decoder.
- $\text{KL}[q(Z_n|X_n; \phi) || P(Z_n)]$  only based on the encoder.  
(A penalty for  $\Sigma_x$  (if  $\Sigma_x$  close to 0 → autoencoder), we want it to close 1 (same as prior).)



## Simulation

- Model performance on the new data embedding (pt-SNE, GPLVM, PCA, and SCVIS)

- Train SCVIS, GPLVM, pt-SNE and KNN classifiers on the original 2200 simulated data.
- Applied these trained models to the tenfold dataset (22000 simulated data.)



- SCVIS performs significantly better for different Ks.
- For a larger K, SCVIS assigns the outliers to the six genuine clusters (performance decrease).
- PCA is worse than other methods.

<https://github.com/jeff665547/SkillShare-SCVIS/raw/master/SCVIS%20A%20VAE-based%20approach.pdf>