### **Week 11: PCA What is Principal Component Analysis and SVD**

#### **Higher Dimensional Data**



(Each point is a single cell (observation), we are working in 'gene space'





#### $N=2$   $N=3$   $N=4$

#### **PCA: A rotation**

Transforms data to a new coordinate system where each axes captures maximum possible variance. We are now in 'component space'.



#### Multidimensional Gaussian with its 2 PC's marked





(Note: Variables covary which gives us information)

# **Linear Regression (OLS) vs PC**

 $y \sim x$  vs.  $x \sim y$ 





<https://shankarmsy.github.io/posts/pca-vs-lr> - Know Thy Data

PCA vs Linear Regression



### **What about covariance across this line?**

PCA vs Linear Regression



Maximal Variance along this line (minimal orthogonal to this) No covariance left along this line (if there is non zero covariance, we can still rotate more to get a better line of best fit, we have not gotten maximal variance along line yet)

![](_page_4_Figure_6.jpeg)

#### **What is an Eigen vector?**

![](_page_5_Picture_1.jpeg)

![](_page_5_Picture_2.jpeg)

https://kids.kiddle.co/Eigenvalues\_and\_eigenvectors

Eigen-vectors represent the axes of our new coordinate space. They will no longer change direction in a linear transformation

The associated Eigen Values tell us how much variance that particular eigen vector captures.

![](_page_5_Picture_7.jpeg)

![](_page_5_Picture_8.jpeg)

![](_page_5_Picture_9.jpeg)

Eigen vectors of a covariance matrix provide the direction along which variance varies the most. This is exactly what we want!

## **Eigen Vectors and Covariance Matrices**

P dimensional data (for ex: p genes)

#### $\Sigma = W \Lambda W^T$

Data Matrix: **X** (n points in p dimensions) (n cells and p genes) (Eigendecomposition)

$$
\Sigma = \begin{bmatrix}\n\text{cov}(1,1) & \text{cov}(1,2) & \dots & \text{cov}(1,p) \\
\text{cov}(2,1) & \text{cov}(2,2) & \dots & \text{cov}(2,p) \\
\vdots & \vdots & \ddots & \vdots \\
\text{cov}(p,1) & \text{cov}(p,2) & \dots & \text{cov}(p,p)\n\end{bmatrix}
$$

Finds eigen values **(W)** with zero covariances

![](_page_6_Figure_7.jpeg)

 $\lambda_i$  is eigen value

![](_page_6_Picture_9.jpeg)

### **Single Value Decomposition and rank**

Assume only n observations, but p variables where  $n < p$ . Can we fit it?

rank of X  $r \leq \min(n, p)$ 

Can apply SVD only on cantered data

$$
x^*_{ij} = x_{ij} - \bar{\mathbf{x}}_j
$$

Can decompose it now as:

$$
\mathbf{X}^* = \mathbf{U} \mathbf{S} \mathbf{W}^\top
$$

![](_page_7_Picture_7.jpeg)

Captures the r principal components, eigen vectors of our covariance matrix

### **Plotting data in our lower dimensional space**

Assume q PC's are kept

![](_page_8_Figure_2.jpeg)

#### Data in the new component space

![](_page_8_Figure_4.jpeg)

![](_page_8_Figure_5.jpeg)

#### We are rotating the data by our new vectors in **W**

Gene 1

![](_page_8_Picture_8.jpeg)