# COMP 7180 Quantitative Methods for Data Analytics and Artificial Intelligence

Lecture 4: Dimensionality Reduction (Feature Extraction) – Part I

# Dimensionality

- An object can be described by a set of characters
- Mathematically, an object can be defined as one point in the vector space
  - Each dimension of the vector space is used to describe one character of the object
  - Example: a pixel in an image/video

# How High the dimensionality could be?

 A small gray image with the resolution 100×100 is represented as a 10,000dimensional vector in the pixel space

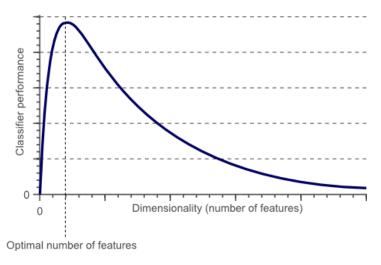
The movie "Kung Fu Panda 3": consider each pixel value as a dimension, the total dimension of this data will be 1280x720x25x60x120x3 = 500,000,000,000 !!!





# **Curse of Dimensionality**

 From a theoretical point of view, increasing the number of features should lead to better performance. However ...



- In practice, the inclusion of more features leads to worse performance (i.e., curse of dimensionality)
  - High computational cost
  - Redundant information

# **Dimensionality Reduction**

### Motivation

- Overcome the curse of dimensionality
- The intrinsic dimension may be small
- Visualization: projection of high-dimensional data onto 2D or 3D
- Data compression: efficient storage and retrieval
- Noise removal: positive effect on query accuracy

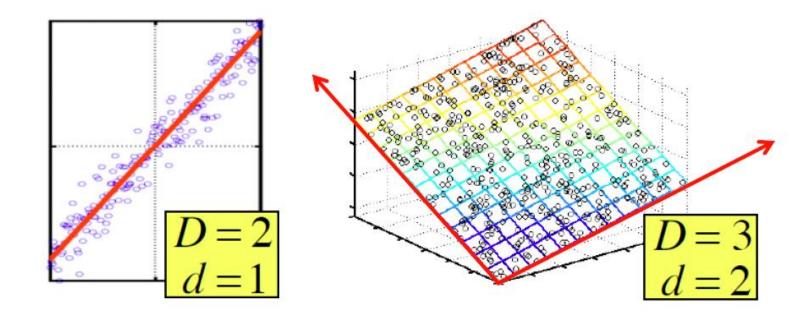
### Definition

 Generate a lower dimensional equivalence to the original highdimensional feature space while capturing essentials of original data according to some criteria

# Applications

 Face recognition, handwritten digit recognition, text summarization, image retrieval, movie editing, protein classification, ...

# **Dimensionality Reduction**



- Assumption: Data lies on or near a low d-dimensional subspace
- Axes of this subspace are effective representation of the data

# **Dimensionality Reduction**

# Compress / reduce dimensionality:

$\mathbf{day}$	We	$\operatorname{Th}$	$\mathbf{F}$ r	$\mathbf{Sa}$	Su
customer	7/10/96	7/11/96	7/12/96	7/13/96	7/14/96
ABC Inc.	1	1	1	0	0
DEF Ltd.	2	2	2	0	0
GHI Inc.	1	1	1	0	0
KLM Co.	5	5	5	0	0
${f Smith}$	0	0	0	2	2
Johnson	0	0	0	3	3
${f Thompson}$	0	0	0	1	1

The above matrix is really "2-dimensional." All rows can be reconstructed by scaling [1 1 1 0 0] or [0 0 0 1 1]

# Rank of a Matrix

- Q: What is rank of a matrix A?
- A: Number of linearly independent columns of A
- For example:

   Matrix  $\mathbf{A} = \begin{bmatrix} 1 & 2 & 1 \\ -2 & -3 & 1 \\ 3 & 5 & 0 \end{bmatrix}$  has rank  $\mathbf{r} = \mathbf{2}$ 
  - >Why? The first two rows are linearly independent, but all three rows are linearly dependent.
- Why do we care about low rank?
  - We can write A as two "basis" vectors: [1 2 1] [-2 -3 1]
  - And new coordinates of : [1 0] [0 1] [1 -1]

# Mathematic Definition of Dimensionality Reduction

Given the high-dimensional data point

$$\mathbf{x} = (x_1, x_2, \cdots, x_D)^T$$

Find a compact representation

$$\mathbf{y} = (y_1, y_2, \dots, y_d)^T \qquad d \le D$$

Construct the transformation function to capture essentials in the original

$$\Phi: \mathbf{x} \to \mathbf{y}$$



$$\rightarrow [32 \ 79 \ 54 \ ... \ ..]^T$$

# Objectives of Dimensionality Reduction

- Generate a lower dimensional equivalence to the original highdimensional feature space while capturing essentials of original data according to some criteria
- Information preserving (unsupervised)
  - We would like to retain as much information (data variance/distance) as possible
  - Principal component analysis (PCA)
- Classification (supervised)
  - We would like to maximize the separation among classes
  - Linear discriminant analysis (LDA)

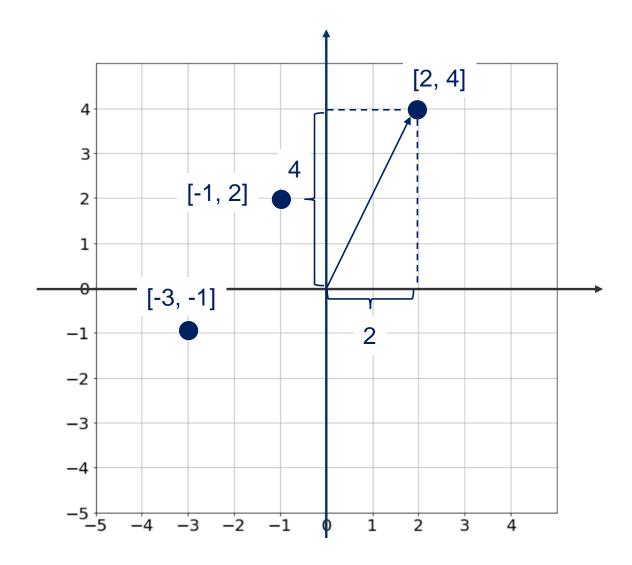
Principal Component Analysis (PCA)

# What is PCA

- Principal component analysis (PCA)
  - A classic linear dimensionality reduction method (Pearson, 1901;
     Hotelling, 1930)
  - Reduce the dimensionality of a data set by finding a new set of projection directions (coordinates), smaller than the original set of directions (coordinates)
  - Preserve most of the samples' information
    - > Directions that capture maximum variance in data

# **Projection**

- Vector projection
  - Dot/inner product of two vectors
  - $\mathbf{a} = [a_1, a_2]^T$ ,  $\mathbf{b} = [b_1, b_2]$
  - $\mathbf{a}^{\mathsf{T}}\mathbf{b} = a_1b_1 + a_2b_2 = \|\mathbf{a}\|\|\mathbf{b}\|\cos\theta$
- Projection on "standard coordinate system"
  - Vector  $[2, 4]^T$  projection on the x-axis is the dot production between [2, 4] and [1, 0]: 2\*1 + 4\*0 = 2
  - Vector  $[2, 4]^T$  projection on the y-axis is the dot production between [2, 4] and [0, 1]: 2\*0 + 4\*1 = 4



# Projection on other directions

- Project on the direction  $\mathbf{u} = \begin{bmatrix} \frac{1}{\sqrt{2}} \end{bmatrix}$
- Project  $[2, 4]^T$  on direction **u**:

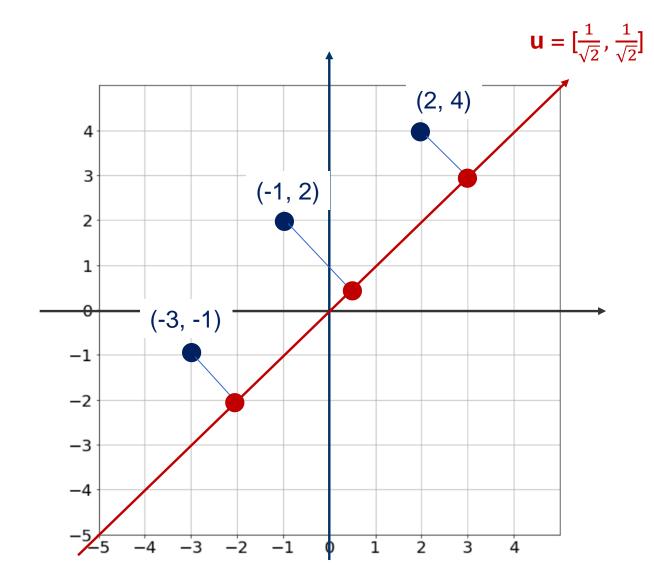
$$2\frac{1}{\sqrt{2}} + 4\frac{1}{\sqrt{2}} = \frac{6}{\sqrt{2}}$$

• Project  $[-1, 2]^T$  on direction **u**:

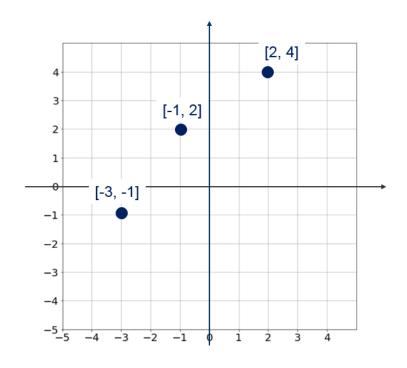
$$-1\frac{1}{\sqrt{2}} + 2\frac{1}{\sqrt{2}} = \frac{1}{\sqrt{2}}$$

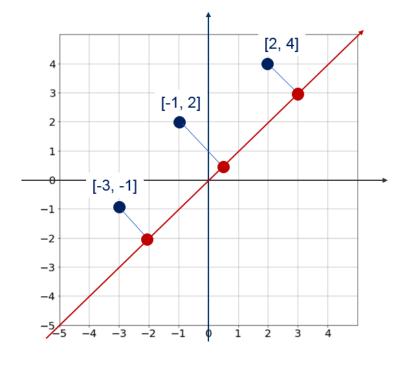
Project [-3, -1]<sup>T</sup> on direction **u**:  

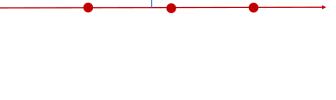
$$-3\frac{1}{\sqrt{2}} + (-1)\frac{1}{\sqrt{2}} = -\frac{4}{\sqrt{2}}$$



# Projection for Dimensionality Reduction







Data Points in 2D

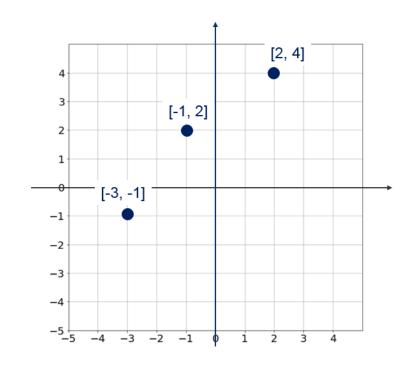
$$\mathbf{X} = \begin{bmatrix} 2 & -1 & -3 \\ 4 & 2 & -1 \end{bmatrix}$$

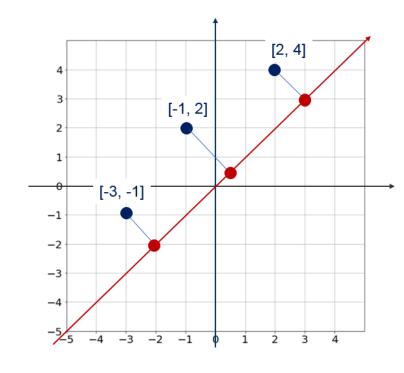
$$\mathbf{u}^{\mathsf{T}}\mathbf{X} = \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \end{bmatrix} \begin{bmatrix} 2 & -1 & -3 \\ 4 & 2 & -1 \end{bmatrix} = \begin{bmatrix} \frac{6}{\sqrt{2}} & \frac{1}{\sqrt{2}} & -\frac{4}{\sqrt{2}} \end{bmatrix}$$

Data Points in 1D

$$\mathbf{Z} = \begin{bmatrix} \frac{6}{\sqrt{2}} & \frac{1}{\sqrt{2}} & -\frac{4}{\sqrt{2}} \end{bmatrix}$$

# Projection for Dimensionality Reduction





This process projects 2 dimensional data to 1 dimensional data (i.e., dimensionality reduction).

Data Points in 2D

Projection onto 1D

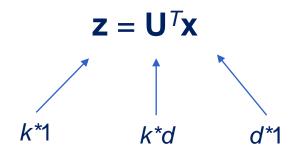
Data Points in 1D

$$\mathbf{X} = \begin{bmatrix} 2 & -1 & -3 \\ 4 & 2 & -1 \end{bmatrix} \qquad \mathbf{u}^{\mathsf{T}} \mathbf{X} = \begin{bmatrix} \frac{1}{\sqrt{2}} & \frac{1}{\sqrt{2}} \end{bmatrix} \begin{bmatrix} 2 & -1 & -3 \\ 4 & 2 & -1 \end{bmatrix} = \begin{bmatrix} \frac{6}{\sqrt{2}} & \frac{1}{\sqrt{2}} & -\frac{4}{\sqrt{2}} \end{bmatrix} \qquad \mathbf{Z} = \begin{bmatrix} \frac{6}{\sqrt{2}} & \frac{1}{\sqrt{2}} & -\frac{4}{\sqrt{2}} \end{bmatrix}$$

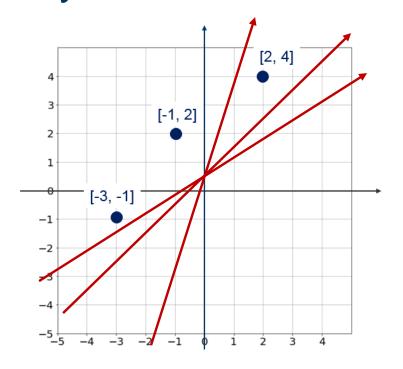
$$\mathbf{Z} = \begin{bmatrix} \frac{6}{\sqrt{2}} & \frac{1}{\sqrt{2}} & -\frac{4}{\sqrt{2}} \end{bmatrix}$$

# **Linear Dimensionality Reduction**

- A projection matrix  $\mathbf{U} = [\mathbf{u}_1 \ \mathbf{u}_2 \ \dots \ \mathbf{u}_k]$  of size  $d^*k$  defines k linear projection directions.
- Each column u<sub>k</sub> in U denotes a linear project direction for d dimensional data (assume k < d)</li>
- Then projection matrix U can be used to transform a high dimensional sample x into a low dimensional sample z by:



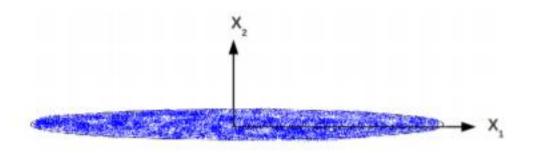
# **Linear Dimensionality Reduction**



There are infinite ways to project the data **X**.

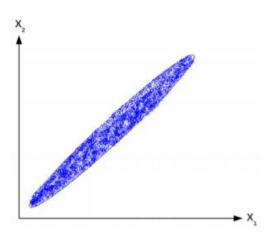
- How do we learn the "best" projection matrix U?
- What criteria should we optimize for learning U?
- Principle Component Analysis (PCA) is an algorithm for doing this.

# PCA as Maximizing Variance: A Simple Illustration



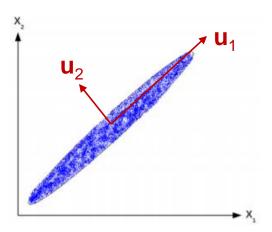
- Consider this two dimensional data
- Each data sample **x** is represented by 2 features  $[x_1, x_2]^T$
- Considering ignoring the feature x<sub>2</sub> for each data sample
- Each 2-dimensional data sample  $\mathbf{x}$  now becomes one-dimensional [ $x_1$ ]
- Are we losing much information by simply removing  $x_2$ ?
  - No. Most of the data spread is along  $x_1$  (very little variance along  $x_2$ )

# PCA as Maximizing Variance: A Simple Illustration



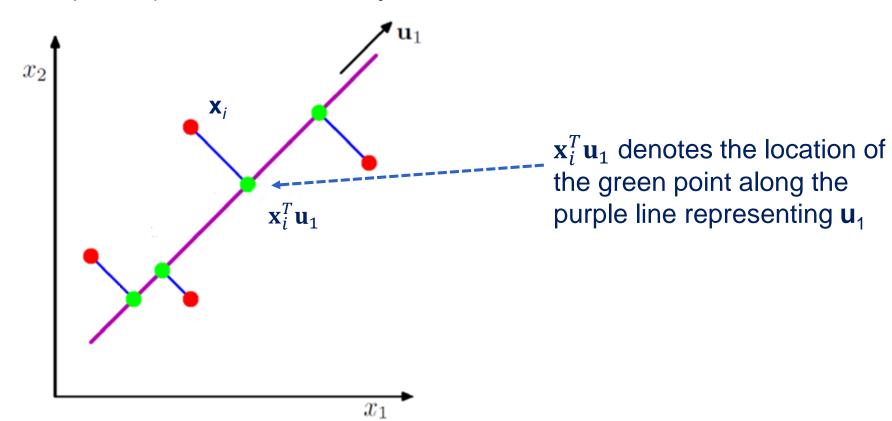
- Consider this two dimensional data
- Each data sample **x** is represented by 2 features  $[x_1, x_2]^T$
- Considering ignoring the feature x<sub>2</sub> for each data sample
- Each 2-dimensional data sample  $\mathbf{x}$  now becomes one-dimensional [ $x_1$ ]
- Are we losing much information by simply removing  $x_2$ ?
  - Yes. This data has substantial variance along both features.

# PCA as Maximizing Variance: A Simple Illustration



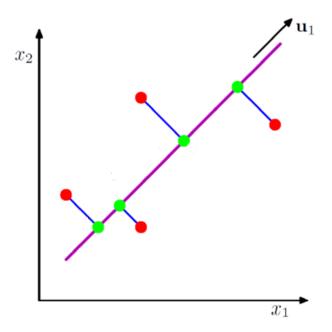
- Now consider we project the data into another two directions u<sub>1</sub>, u<sub>2</sub>
- Each data sample **x** is represented by 2 features  $[z_1, z_2]^T$
- Considering ignoring the feature z<sub>2</sub> for each data sample
- Each 2-dimensional data sample  $\mathbf{x}$  now becomes one-dimensional [ $z_1$ ]
- Are we losing much information by simply removing  $z_2$ ?
  - No. Most of the data spread is along  $z_1$  (very little variance along  $z_2$ )

Projecting  $\mathbf{x}_i$  (a *d*-dimensional feature vector) to a one-dimensional vector  $\mathbf{z}_i$  by  $\mathbf{u}_1$ :  $z_i = \mathbf{u}_1^T \mathbf{x}_i = \mathbf{x}_i^T \mathbf{u}_1$ 



- Projecting  $\mathbf{x}_i$  (a *d*-dimensional feature vector) to a one-dimensional vector  $\mathbf{z}_i$  by  $\mathbf{u}_1$ :  $z_i = \mathbf{u}_1^T \mathbf{x}_i = \mathbf{x}_i^T \mathbf{u}_1$
- Therefore, the mean of projections of all data (i.e., "center" of the green points ) can be computed as

$$\frac{\sum_{i=1}^{n} \mathbf{x}_{i}^{T} \mathbf{u}_{1}}{n} = \frac{\sum_{i=1}^{n} \mathbf{x}_{i}^{T}}{n} \mathbf{u}_{1} = \overline{\mathbf{x}}^{T} \mathbf{u}_{1}$$



 $\bar{\mathbf{x}}$  is the mean feature vector  $\bar{\mathbf{x}} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i$ 

 Variance of the projected data (i.e., "spread" of the green points)

$$\frac{\sum_{i=1}^{n} (\mathbf{x}_{i}^{T} \mathbf{u}_{1} - \bar{\mathbf{x}}^{T} \mathbf{u}_{1})^{2}}{n} = \frac{\sum_{i=1}^{n} ((\mathbf{x}_{i}^{T} - \bar{\mathbf{x}}^{T}) \mathbf{u}_{1})^{2}}{n}$$

Variance of the projected data

$$\frac{\sum_{i=1}^{n}((\mathbf{x}_i^T - \bar{\mathbf{x}}^T)\mathbf{u}_1)^2}{n} = \mathbf{u}_1^T \frac{\sum_{i=1}^{n}(\mathbf{x}_i - \bar{\mathbf{x}})(\mathbf{x}_i^T - \bar{\mathbf{x}}^T)}{n} \mathbf{u}_1$$

Let 
$$\mathbf{S} = \frac{\sum_{i=1}^n (\mathbf{x}_i - \bar{\mathbf{x}}) (\mathbf{x}_i^T - \bar{\mathbf{x}}^T)}{n}$$
, the variance of the projected data is 
$$\mathbf{u}_1^T \mathbf{S} \mathbf{u}_1$$

• **S** is the  $d^*d$  data covariance matrix. If data is already centered (i.e.,  $\bar{\mathbf{x}} = 0$ ), then  $\mathbf{S} = \frac{\sum_{i=1}^{n} (\mathbf{x}_i)(\mathbf{x}_i^T)}{n} = \frac{1}{n} \mathbf{X} \mathbf{X}^T$ 



- Objective: We want  $\mathbf{u}_1$  that the variance of the project data is maximized  $\max_{\mathbf{u}_1^T} \mathbf{S} \mathbf{u}_1$
- To prevent trivial solution (max variance = infinite), assume  $\|\mathbf{u}_1\|_2 = \sqrt{\mathbf{u}_1^T \mathbf{u}_1} = 1$ . Therefore  $\mathbf{u}_1^T \mathbf{u}_1 = 1$
- Therefore, u<sub>1</sub> can be obtained by solving the following optimization problem

$$\max_{\mathbf{u}_1} \mathbf{u}_1^T \mathbf{S} \mathbf{u}_1 + \lambda_1 (1 - \mathbf{u}_1^T \mathbf{u}_1)$$

$$\lambda_1 \text{ is a Lagrange multiplier}$$

- The objective:  $\max_{\mathbf{u}_1} \mathbf{u}_1^T \mathbf{S} \mathbf{u}_1 + \lambda_1 (1 \mathbf{u}_1^T \mathbf{u}_1)$
- Obtaining the optimal solution by taking the derivative with respect to u<sub>1</sub> and setting to zero

$$\mathbf{S}\mathbf{u}_1 = \lambda_1 \mathbf{u}_1$$

- Thus  $\mathbf{u}_1$  is an eigenvector of **S** (with corresponding eigenvalue  $\lambda_1$ )
- **S** is a *d*\**d* matrix, there are *d* possible eigenvectors, which ones to take?

• Note that the constraint  $\mathbf{u}_1^T \mathbf{u}_1 = 1$ , the variance of the projected data is

$$\mathbf{u}_1^T \mathbf{S} \mathbf{u}_1 = \mathbf{u}_1^T \lambda_1 \mathbf{u}_1 = \lambda_1 \mathbf{u}_1^T \mathbf{u}_1 = \lambda_1$$

• Therefore, variance is maximized when  $\mathbf{u}_1$  is the (top) eigenvector with largest eigenvalue.

Other directions can also be found similarly (with each being orthogonal to all previous ones)

• Question: What is u<sub>2</sub>?

$$\max_{\mathbf{u}_{2}} \mathbf{u}_{2}^{T} \mathbf{S} \mathbf{u}_{2}$$

$$s. t. \mathbf{u}_{2}^{T} \mathbf{u}_{2} = 1, \mathbf{u}_{2}^{T} \mathbf{u}_{1} = 0$$

$$\max_{\mathbf{u}_{2}} \mathbf{u}_{2}^{T} \mathbf{S} \mathbf{u}_{2} - \lambda (\mathbf{u}_{2}^{T} \mathbf{u}_{2} - 1) - \phi \mathbf{u}_{2}^{T} \mathbf{u}_{1}$$

$$\frac{\partial}{\partial \mathbf{u}_{2}} (\mathbf{u}_{2}^{T} \mathbf{S} \mathbf{u}_{2} - \lambda (\mathbf{u}_{2}^{T} \mathbf{u}_{2} - 1) - \phi \mathbf{u}_{2}^{T} \mathbf{u}_{1}) = 0$$

• Question: What is  $\mathbf{u}_2$ ?

$$\frac{\partial}{\partial \mathbf{u}_2} (\mathbf{u}_2^T \mathbf{S} \mathbf{u}_2 - \lambda (\mathbf{u}_2^T \mathbf{u}_2 - 1) - \phi \mathbf{u}_2^T \mathbf{u}_1) = 0$$

$$2\mathbf{S}\mathbf{u}_2 - 2\lambda\mathbf{u}_2 - \phi\mathbf{u}_1 = 0$$

$$\phi = 0$$
?

$$\mathbf{S}\mathbf{u}_2 = \lambda \mathbf{u}_2$$

 $\mathbf{u}_2$  is the eigenvector with the second largest eigenvalue.

To show that  $\phi = 0$ , we multiply  $\mathbf{u}_1^T$  on both side, and we have:

$$\mathbf{u}_{1}^{T}(2\mathbf{S}\mathbf{u}_{2} - 2\lambda\mathbf{u}_{2} - \phi\mathbf{u}_{1}) = \mathbf{u}_{1}^{T}0 = 0$$

$$\mathbf{\mathbf{U}}_{1}^{T}\mathbf{S}\mathbf{u}_{2} - 2\lambda\mathbf{u}_{1}^{T}\mathbf{u}_{2} - \phi\mathbf{u}_{1}^{T}\mathbf{u}_{1} = 0$$

$$\mathbf{\mathbf{U}}_{1}^{T}\mathbf{S}^{T}\mathbf{u}_{2} - 0 - \phi\mathbf{u}_{1}^{T}\mathbf{u}_{1} = 0$$

$$\mathbf{\mathbf{U}}_{1}^{T}\mathbf{U}_{1} = 0$$

$$\mathbf{\mathbf{U}}_{1}^{T}\mathbf{u}_{2} - \phi\mathbf{u}_{1}^{T}\mathbf{u}_{1} = 0$$

$$\mathbf{\mathbf{U}}_{1}^{T}\mathbf{u}_{1} = 0$$

$$\mathbf{\mathbf{U}}_{1}^{T}\mathbf{\mathbf{U}}_{1} = 0$$

# Steps of Principle Component Analysis

- Center the data (subtract the mean  $\bar{\mathbf{x}} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i$  from each data point) to get  $\mathbf{X}_c$
- Compute the covariance matrix S using the centered data as

$$\mathbf{S} = \frac{1}{n} \mathbf{X}_c \mathbf{X}_c^T$$

- Do an eigen-decomposition of the covariance matrix S
- Take first k leading eigenvectors  $\{\mathbf{u}_1, \dots, \mathbf{u}_k\}$  with k largest eigenvalue  $\{\lambda_1, \dots, \lambda_k\}$
- The final k dimensional representation of data is obtained by

$$\mathbf{Z} = \mathbf{U}^{\mathsf{T}} \mathbf{X}_{c}$$

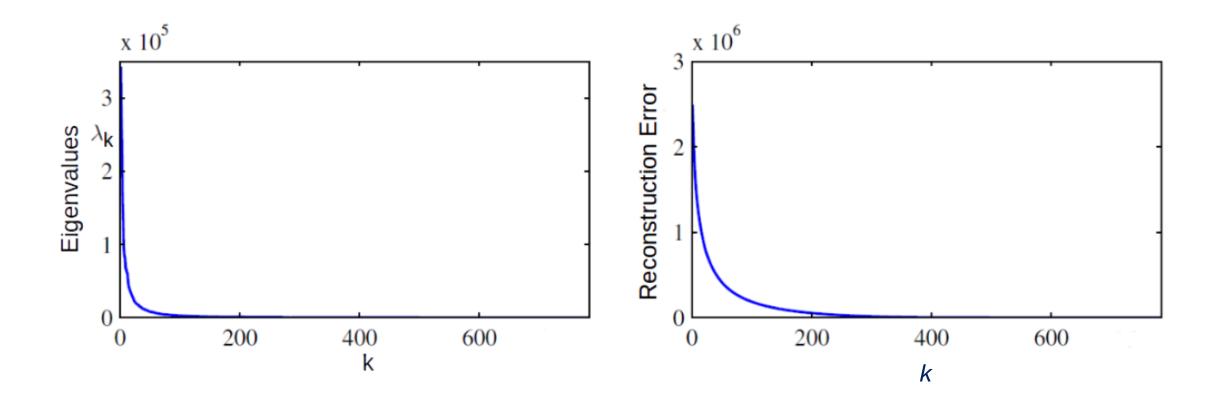
# How many Principal Components to Use?

• Eigenvalue  $\lambda_i$  measures the variance captured by the corresponding projection direction  $\mathbf{u}_i$ 

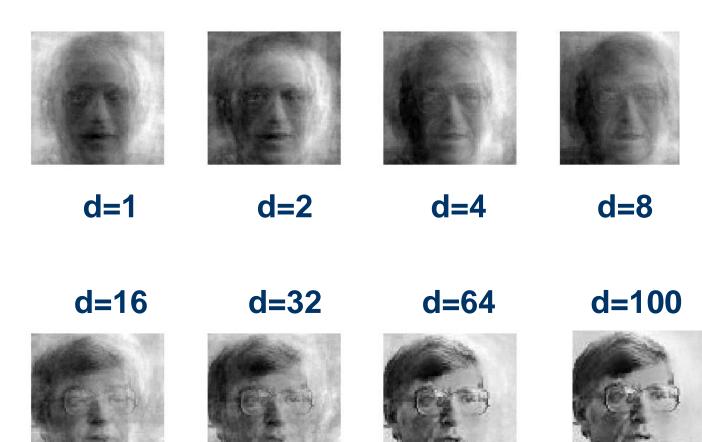
$$\mathbf{u}_i^T \mathbf{S} \mathbf{u}_i = \mathbf{u}_i^T \lambda_i \mathbf{u}_i = \lambda_i \mathbf{u}_i^T \mathbf{u}_i = \lambda_i$$

- The "left-over" variance will therefore be  $\sum_{i=k+1}^{d} \lambda_i$
- Can choose k by looking at what fraction of variance is captured by the first k projection directions:  $\mathbf{u}_1, \mathbf{u}_2, \dots \mathbf{u}_k$
- Another direct way is to look at the spectrum of the eigenvalues plot, or the plot of reconstruction error vs k

# How many Principal Components to Use?



# PCA for image compression



$$z = U^T x$$

$$\overline{x} = U z$$

$$\overline{x} = U U^T x$$

# Original Image 64\*64



# References and Acknowledgement

- Korn F, Jagadish H V, Faloutsos C. Efficiently supporting ad hoc queries in large datasets of time sequences[J]. Acm Sigmod Record, 1997, 26(2): 289-300.
- Zhang D, Zou L, Zhou X, et al. Integrating feature selection and feature extraction methods with deep learning to predict clinical outcome of breast cancer[J]. Ieee Access, 2018, 6: 28936-28944.
- Slide from Maria-Florina Balcan, Carnegie Mellon University, Advanced Introduction to Machine Learning. https://www.cs.cmu.edu/~10715-f18/lectures/pca-2018.pdf
- Slide from Piyush Rai, Indian Institute of Technology Kanpur, Probabilistic Machine Learning (CS772A). https://www.cse.iitk.ac.in/users/piyush/courses/pml\_winter16/slides\_lec10.pdf
- https://slideplayer.com/slide/14520684/
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