# Data Mining: W4240 Dimensionality Reduction I

Lecture 23

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#### Outline

- 1. High dimensional data
- 2. Principal components analysis (PCA) overview
- 3. (Review: eigenvalues and eigenvectors)
- 4. PCA computation

Next time: more PCA, doing PCA with R

The Curse of Dimensionality: "There are multiple phenomena referred to by this name in domains such as numerical analysis, sampling, combinatorics, machine learning, data mining and databases. The common theme of these problems is that when the dimensionality increases, the volume of the space increases so fast that the available data becomes sparse. This sparsity is problematic for any method that requires statistical significance. In order to obtain a statistically sound and reliable result, the amount of data needed to support the result often grows exponentially with the dimensionality." (Wikipedia)

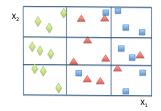
Example: predict the class given covariates



Pretty easy when data is in 1 dimension (9 samples gives us a good enough predictor)

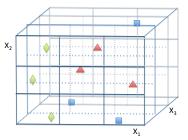
*Example:* predict the class given covariates; same data in 2 dimensions





In 2 dimensions, we are much less sure about our predictions (9 for 9 regions...) or we need more data ( $3 \times 3^2 = 27$  observations for same sureity)

*Example:* predict the class given covariates; original data in 3 dimensions



In 3 dimensions, we only have data in 1/3 of our regions (9 in 27 regions...). For same predictive power, we need  $3 \times 3^3 = 81$ 

#### Implications:

- ▶ Data required for prediction grows like  $n^d$  (prediction methods fail)
- Describing and storing the data becomes increasingly difficult (summarizing data hard)

#### Goals:

- Summarize data compactly
- Reduce dimensions as preprocessing for other methods

#### How do we beat the curse?

**Problem:** want to describe D&D characters

#### Attributes:

- Strength
- Dexterity
- Constitution
- ▶ Intelligence
- Wisdom
- Charisma

Each can take values in  $\{1, 2, ..., 20\}$ . Do I really need 64 million  $(20^6)$  different values to describe characters well?

#### Possible character A:

- ► Strength = 18
- ► Dexterity = 14
- ► Constitution =16
- ▶ Intelligence = 10
- ▶ Wisdom = 11
- ► Charisma = 9

#### Possible character B:

- ▶ Strength = 18
- ► Dexterity = 18
- ► Constitution = 6
- ► Intelligence = 4
- ▶ Wisdom = 19
- ▶ Charisma = 9

Would you expect to see both of these characters?

Can describe set of possible characters pretty well:

- mage
- fighter
- rogue
- cleric
- specialized subtypes: rangers, monks, bards, etc

64 million possible builds probably closer to 10,000... and can even summarize more efficiently with a small set of classes (clusters)

**Clustering** can be used to efficiently summarize large data sets, but it does not help for preprocessing data for statistical methods

**Problem:** want to describe student performance and compare students

#### Data:

- scores on five homeworks
- midterm score
- final score

How can I summarize data for a student in a way that will be useful for comparisons?

#### If data truly high dimensional:

- scores for each of 7 items have low correlation
- need all items to summarize student (good on HWs 1,2,4, middling on HW 3 and midterm, abysmal on HW 5, low on final)
- no good way to compare students

But we would not expect to see a student like the one above. Why?

Any suggestions for summarizing student performance in a way that is useful for comparing students? Why would this be reasonable?

I have some (randomized) data. Let's look at it in R.

Does it agree with your prediction about summary?

This approach is called **dimensionality reduction**. We will be studying this for the next two classes.

Data with d dimensions:

$$\mathbf{x} = \begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ \vdots \\ x_d \end{bmatrix}$$

 $x_j$  has many names: covariate, feature, input, or independent variable.

Assumptions for today:

- **>** all features are continuous-valued  $(\mathbf{x} \in \mathbb{R}^d)$
- ▶ *d* is large (10 or 100 or more)

Data with *n* observations and *d* dimensions:

$$\mathbf{X} = \begin{bmatrix} x_{11} & \dots & x_{1d} \\ x_{21} & \dots & x_{2d} \\ \vdots & & \vdots \\ x_{n1} & \dots & x_{nd} \end{bmatrix} = \begin{bmatrix} \mathbf{x}_1^T \\ \mathbf{x}_2^T \\ \vdots \\ \mathbf{x}_n^T \end{bmatrix}$$

Here  $\mathbf{x}^T$  means the transpose of  $\mathbf{x}$ .

Today, **bold** means matrix or vector, and *plain* means scalar.

#### Two approaches:

► **Feature Selection:** choose a subset of features for prediction task

$$[x_1, x_2, \dots, x_d] \rightarrow [x_{i1}, x_{i2}, \dots, x_{iK}]$$

But you need to know your prediction task before you do feature selectionn.

► **Feature Extraction:** create new features by combining existing ones

$$[x_1, x_2, \dots, x_d] \rightarrow [f_1(x_{11}, x_{12}, \dots, x_{1d}), \dots, f_K(x_{K1}, x_{K2}, \dots, x_{Kd})]$$
  
=  $[y_1, \dots, y_K]$ 

Feature extraction is often a data preprocessing step since you do not need to know which prediction methods you will use.

Today, we will look at linear feature extraction

$$\begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ \vdots \\ x_d \end{bmatrix} \rightarrow \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_K \end{bmatrix} = \begin{bmatrix} w_{11} & w_{12} & \dots & \dots & w_{1d} \\ w_{21} & w_{22} & \dots & \dots & w_{2d} \\ \vdots & & & & \vdots \\ w_{K1} & w_{K2} & \dots & \dots & w_{Kd} \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ \vdots \\ \vdots \\ x_d \end{bmatrix}$$

Terminology:  $y_{i1}, \ldots, y_{iK}$  are the scores,  $\mathbf{w}_1, \ldots, \mathbf{w}_K$  are the loadings.

#### Linear Feature Extraction

Student score averages are a form of linear feature extraction:

$$y = [0.067, 0.067, 0.133, 0.067, 0.067, 0.25, 0.35] \begin{bmatrix} HW1 \\ HW2 \\ HW3 \\ HW4 \\ HW5 \\ Midterm \\ Final \end{bmatrix}$$

The student score average is the *score* and the assignment weights are the *loadings* 

#### Linear Feature Extraction

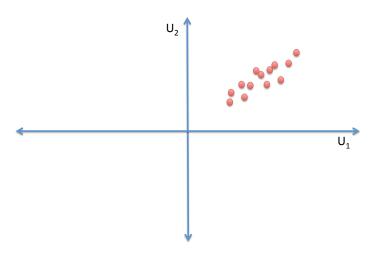
How can we find a good set of loadings and scores for a general data set?

Principal components analysis (PCA): a covariance matrix singular value decomposition method for unsupervised data

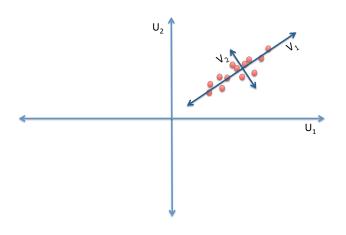
Principal components: a set of linearly uncorrelated variables—these will be the loadings

Multiply loadings with original data to get scores!

Basic idea: have a dataset



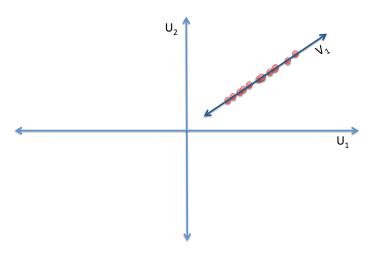
Basic idea: given dataset, want to find rotation (linear transformation) that best describes data



$$y_1 = w_{11}x_1 + w_{12}x_2, \ y_2 = w_{21}x_1 + w_{22}x_2$$

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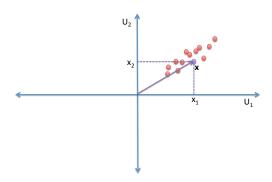
Basic idea: given linear transformation, we can throw out the less descriptive dimensions and still have a decent representation



Mathematically, how do we do this?

Simple case:  $\mathbf{x} \in \mathbb{R}^2$ , and we want a good projection  $y \in \mathbb{R}$ 

- ightharpoonup make  $x_1, x_2$  scalars
- ightharpoonup make the axes vectors,  $\mathbf{u}_1, \mathbf{u}_2$
- $\mathbf{x} = x_1 \mathbf{u}_1 + x_2 \mathbf{u}_2$



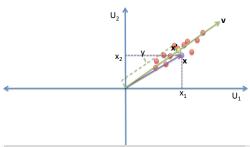
Mathematically, how do we do this?

Simple case:  $\mathbf{x} \in \mathbb{R}^2$ , and we want a good projection  $y \in \mathbb{R}$ 

- now suppose we make a new direction vector v
- ▶ let  $\mathbf{w}_1$  be the linear transformation of  $\mathbf{u}_1$ ,  $\mathbf{u}_2$  into  $\mathbf{v}$ :

$$\mathbf{v} = w_{11}\mathbf{u}_1 + w_{12}\mathbf{u}_2$$

- we can project **x** onto **v** to make  $\mathbf{x}' = y\mathbf{w}_1$
- $\blacktriangleright$  two coordinates  $x_1, x_2$  get turned into one, y



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#### To find best $\mathbf{w}_1$ :

- "closeness" is based on squared error between original points and new points
- usual notion of distance is Euclidean:

$$d(\mathbf{x}_i, \mathbf{x}_i') = \sqrt{\sum_{j=1}^d \left(x_{ij} - x_{ij}
ight)^2}$$

- ▶ we will measure distance as Euclidean distance squared → big deviations heavily penalized, smaller deviations less so
- objective is to minimize squared errors

$$\hat{\mathbf{w}_1} = \arg\min_{\mathbf{w}_1} \left\{ \sum_{i=1}^n ||\mathbf{x}_i - \mathbf{x}_i'||_2^2 = \sum_{i=1}^n \sum_{j=1}^d \left(\mathbf{x}_{ij} - \mathbf{x}_{ij}'\right)^2 \right\}$$

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#### To find best $\mathbf{w}_1$ :

also have constraint:

$$||\mathbf{w}_1||_2 = \sqrt{\sum_{j=1}^d w_{ij}^2} = 1$$

this means one step in old coordinates  ${\bf u}$  is equal to one step in new coordinates  ${\bf v}$  (this is called the L2 norm... we will use this and the L1 norm  $||{\bf x}|| = \sum_{j=1}^d |x_j|$  a lot in this class)

To find the best single feature  $\mathbf{w}_1$ :

- 1. center the data (subtract mean)
- 2. find best single feature,  $\mathbf{w}_1$  by

$$\begin{split} \hat{\mathbf{w}}_1 &= \arg \min_{\mathbf{w}_1 : ||\mathbf{w}_1||_2 = 1} \sum_{i=1}^n \sum_{j=1}^d \left( x_{ij} - x_{ij}' \right)_2^2 \\ &= \arg \min_{\mathbf{w}_1 : ||\mathbf{w}_1||_2 = 1} \sum_{i=1}^n \sum_{j=1}^d \left( x_{ij} - y_i w_{1j} \right)^2 \\ &= \arg \min_{\mathbf{w}_1 : ||\mathbf{w}_1||_2 = 1} \sum_{i=1}^n \sum_{j=1}^d \left( x_{ij} - w_{1j} \mathbf{x}_i^T \mathbf{w}_1 \right)^2 \\ &= \dots \qquad (\mathrm{matrix \ algebra}) \\ &= \arg \max_{\mathbf{w}_1 : ||\mathbf{w}_1||_2 = 1} \left\{ \mathbf{w}_1^T \mathbf{X}^T \mathbf{X} \mathbf{w}_1 = \frac{\mathbf{w}_1^T \left( \frac{1}{n} \mathbf{X}^T \mathbf{X} \right) \mathbf{w}_1}{\mathbf{w}_1^T \mathbf{w}_1} \right\} \end{split}$$

Fun fact:  $\frac{1}{n}\mathbf{X}^T\mathbf{X} = \frac{1}{n}\sum_{i=1}^n \mathbf{x}_i \mathbf{x}_i^T$  is the maximum likelihood estimate of the covariance matrix for  $[x_1, \dots, x_d]^T$ 

Another fun fact:

$$\frac{\mathbf{w}_{1}^{T}\left(\frac{1}{n}\mathbf{X}^{T}\mathbf{X}\right)\mathbf{w}_{1}}{\mathbf{w}_{1}^{T}\mathbf{w}_{1}}$$

is called a Rayleigh quotient

Since  $\frac{1}{n}\mathbf{X}^T\mathbf{X}$  is symmetric:

- ▶ the Rayleigh quotient has a value that is equal to the largest eigenvalue of  $\frac{1}{n}\mathbf{X}^T\mathbf{X}$
- ▶ the value of **w**<sub>1</sub> that maximizes the Rayleigh quotient is the is the eigenvector associated with the largest eigenvalue

To get the next best feature,  $\mathbf{w}_2$ :

1. re-center the data by subtracting off the projection  $\mathbf{x}_i'$  from  $\mathbf{x}_i$ :

$$\tilde{\textbf{X}}_1 = \textbf{X} - \textbf{X} \textbf{w}_1 \textbf{w}_1^T$$

2. minimize squared error as we did previously:

$$\hat{\mathbf{w}}_2 = \arg\min_{\mathbf{w}_2 : ||\mathbf{w}_2||_2 = 1} \mathbf{w}_2^T \tilde{\mathbf{X}}_1^T \tilde{\mathbf{X}}_1 \mathbf{w}_2$$

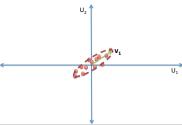
This process can be repeated until we have constructed rank(X) features

There is actually a nicer geometric interpretation.

objective is to minimize squared errors

$$\hat{\mathbf{w}}_1 = \arg\min_{\mathbf{w}_1 : ||\mathbf{w}_1||_2 = 1} \sum_{i=1}^n \sum_{j=1}^d (\mathbf{x}_{ij} - \mathbf{x}'_{ij})_2^2$$

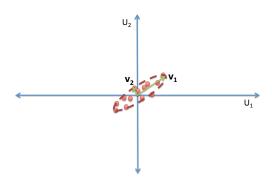
- center data
- ▶ fit a multivariate Gaussian distribution fit to the data
- $\blacktriangleright$  Gaussian has mean 0, covariance  $\Sigma$
- ightharpoonup 
  igh



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This generalizes to multiple dimensions

- suppose we have d original dimensions and we would like K new ones
- ▶ the optimal vectors have the same direction as the K ellipse directions with maximum variance



How do we find the K most descriptive directions?

- center everything to have mean 0
- fit a multivariate Gaussian distribution to the data
- estimate the covariance matrix,

$$\hat{\Sigma} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i \mathbf{x}_i^T$$

- ▶ find the eigenvalues and eigenvectors of the covariance matrix
- ▶ the *K* most descriptive directions are the eigenvectors associated with the *K* largest eigenvalues

Matrices can be used to describe transformations:

- ightharpoonup x 
  ightarrow Ax is a transformation of x
- Examples (look at these graphically):

$$\mathbf{x} = (1,1), (1,0), (1,-1), (0,1), (-1,1), (-1,0), (-1,1), (0,-1)$$

Matrices can be used to describe transformations:

 under transformations, some of the original directions may be preserved

$$\mathbf{A}\mathbf{x} = \lambda \mathbf{x}$$

- ▶ the direction **x** is an eigenvector of **A**
- the scalar  $\lambda$  is an eigenvalue **A**
- the eigenvalue describes the magnitude of the change in x under A

Wikipedia actually has a nice .gif demonstrating this

To find the eigenvectors of **A**, find the roots of the polynomial

$$\det\left(\mathbf{A}-\lambda I\right)=0$$

Let's do this with our examples:

We can use the eigenvalues to find the eigenvectors

- start with eigenvalue  $\lambda_i$
- solve the set of linear equations

$$\mathbf{A}\mathbf{x} = \lambda_i \mathbf{x}$$

**x** is the eigenvector associated with  $\lambda_i$ 

Let's do this with our examples:

#### Back to PCA...

- 1. center the data
- 2. compute the covariance matrix  $\frac{1}{n} \mathbf{X}^T \mathbf{X}$
- 3. compute the eigenvectors of  $\frac{1}{n}\mathbf{X}^T\mathbf{X}$  along with their eigenvalues to get loadings
- 4. for the eigenvectors with the K largest eigenvalues, make factor scores by setting

$$\mathbf{y}_{i} = \begin{bmatrix} y_{i1} \\ y_{i2} \\ \vdots \\ y_{iK} \end{bmatrix} = \begin{bmatrix} w_{11} & w_{12} & \dots & w_{1d} \\ w_{21} & w_{22} & \dots & w_{2d} \\ \vdots & & & \vdots \\ w_{K1} & w_{K2} & \dots & w_{Kd} \end{bmatrix} \begin{bmatrix} x_{i1} \\ x_{i2} \\ x_{i3} \\ \vdots \\ \vdots \\ x_{id} \end{bmatrix}$$

- 5. do computations in the (smaller) **y** space, the space of factor scores
- 6. transform results back to original space (if needed)

$$\mathbf{y}_{i} = \begin{bmatrix} y_{i1} \\ y_{i2} \\ \vdots \\ y_{iK} \end{bmatrix} = \begin{bmatrix} w_{11} & w_{12} & \dots & w_{1d} \\ w_{21} & w_{22} & \dots & w_{2d} \\ \vdots & & & \vdots \\ w_{K1} & w_{K2} & \dots & w_{Kd} \end{bmatrix} \begin{bmatrix} x_{i1} \\ x_{i2} \\ x_{i3} \\ \vdots \\ \vdots \\ x_{id} \end{bmatrix}$$

Let's do an example:

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

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

$$\det(\frac{1}{n}\mathbf{X}^{\mathsf{T}}\mathbf{X} - \lambda I) =$$

$$[\lambda_1, \lambda_2] =$$

Let's do an example:

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

$$[\mathbf{w}_1, \mathbf{w}_2] =$$

$$[\mathbf{y}_1,\mathbf{y}_2] =$$

### PCA: Computational Underpinnings

So how do we get these values? Matrix factorization.

- ▶ can compute eigenvectors/eigenvalues of X<sup>T</sup>X through spectral decomposition
- ► can use singular value decomposition of X
  - more computationally stable than first method
  - ▶ better when d > n
  - we won't get into the details

#### Question of the Day

I have found that giving you some simple practice questions helps improve test performance. At the beginning of next class, I will have one of you work this out on the board (voluntary basis).

Find the principal components for the following dataset:

$$\mathbf{X} = \begin{bmatrix} -6 & -4 \\ -2 & 3 \\ 2 & -3 \\ 6 & 4 \end{bmatrix}$$