PCA & LDA



Dimensionality Reduction

- Many modern data domains involve huge numbers of features / dimensions
 - Documents: thousands of words, millions of bigrams
 - Images: thousands to millions of pixels
 - Genomics: thousands of genes, millions of DNA polymorphisms

Why to reduce dimensions?

- High dimensionality has many costs
 - Redundant and irrelevant features degrade performance of some ML algorithms
 - Difficulty in interpretation and visualization
 - Computation may become infeasible
 - what if your algorithm scales as O(n³)?
 - Curse of dimensionality

A Toy Example

	Educations	Jobs opportunities	Safety	Air quality	Entertainment	Connectivity	Foods
City-1	8	7	7	9	8	7	8
City-2	6	8	7	6	8	9	8
City-3	9	10	7	8	8	4	8
City-4	4	4	6	2	8	5	7
City-5	3	6	6	5	8	9	8
City-6	2	7	6	3	8	4	7
City-7	7	8	7	8	8	6	8
City-8	8	3	6	3	8	8	7
City-9	5	9	7	7	8	5	7
City-10	6	8	6	9	8	9	8

Another example

	Educations	Jobs opportunities	Safety	Air quality	Entertainment	Connectivity	Foods
City-1	8	7	7	9	8	7	8
City-2	6	8	7	6	8	9	8
City-3	9	10	7	8	8	4	8
City-4	4	4	6	2	8	5	7
City-5	3	6	6	5	8	9	8
City-6	2	7	6	3	8	4	7
City-7	7	8	7	8	8	6	8
City-8	8	3	6	3	8	8	7
City-9	5	9	7	7	8	5	7
City-10	6	8	6	9	8	9	8

Dimension reduction

■ Two approaches to perform dim. reduction \(\partial^{N}\) \(\partial^{M}\) (M<N)</p>

Feature selection: choosing a subset of all the features

$$[x_1 \ x_2...x_N] \xrightarrow{\text{feature selection}} [x_{i_1} \ x_{i_2}...x_{i_M}]$$

Feature extraction: creating new features by combining existing ones

$$[x_1 \ x_2...x_N] \xrightarrow{\text{feature}} [y_1 \ y_2...y_M] = f([x_{i_1} \ x_{i_2}...x_{i_M}])$$

 In either case, the goal is to find a low-dimensional representation of the data that preserves (most of) the information or structure in the data

Linear feature extraction

- The "optimal" mapping y=f(x) is, in general, a non-linear function whose form is problem-dependent
 - Hence, feature extraction is commonly limited to linear projections y=Wx

$$\begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ x_N \end{bmatrix} \xrightarrow{\text{linear feature extraction}} \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_M \end{bmatrix} = \begin{bmatrix} w_{11} & w_{12} & \cdots & w_{1N} \\ w_{21} & w_{22} & \cdots & w_{2N} \\ \vdots & \vdots & \ddots & \vdots \\ w_{M1} & w_{M2} & & & w_{MN} \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ x_N \end{bmatrix}$$

Models



- Linear methods
 - Principal component analysis (PCA)
 - Multidimensional scaling (MDS)
 - Independent component analysis (ICA)
- Nonlinear methods
 - Kernel PCA
 - Locally linear embedding (LLE)
 - Laplacian eigenmaps (LEM)
 - Semidefinite embedding (SDE)

Statistics Preliminaries

Mean

 Let X₁,X₂, , ... X_n be n observations of a random variable X

$$\mu = \overline{X} = E[X] = \frac{1}{n} \sum_{i=1}^{n} X_i$$

 Mean is a measure of central tendency (others are mode and median)

Standard Deviation

Measure of variability (square root of variance)

$$\sigma = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (X_i - \mu)^2}$$

Statistics Preliminaries

Covariance

A measure of how two variables change together

$$\Sigma_{XY} = \frac{1}{n} \sum_{i=1}^{n} (X_i - \mu_X) (Y_i - \mu_Y)^T$$

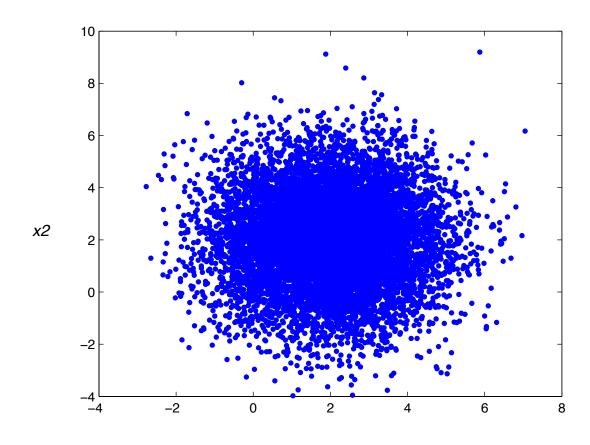
$$\Sigma_{XY} = E[(X - E[X]) (Y - E[Y])]$$

Covariance Matrix

$$\Sigma = \begin{pmatrix} \sigma_{X1}^{2} & \Sigma_{X1X2} & \dots & \Sigma_{X1Xd} \\ \Sigma_{X1X2} & \sigma_{X2}^{2} & \dots & \Sigma_{X2Xd} \\ \vdots & & & \vdots \\ \Sigma_{X1Xd} & \Sigma_{X2Xd} & \dots & \sigma_{Xd}^{2} \end{pmatrix}$$

Statistics Preliminaries – An Example

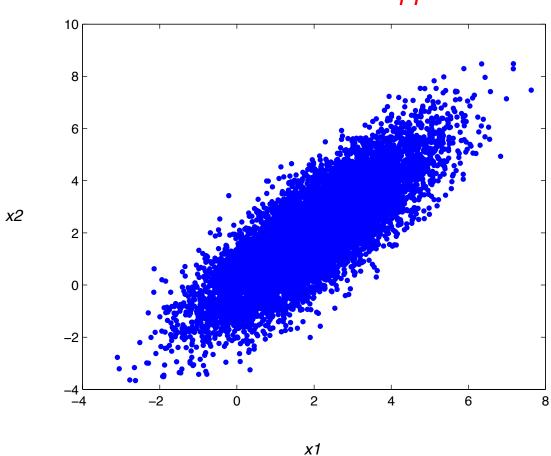
$$\mu = \begin{pmatrix} 2 \\ 3 \end{pmatrix} \qquad \Sigma = \begin{pmatrix} 2 & 0 \\ 0 & 3 \end{pmatrix}$$



PCA – An Example

$$\mu = \begin{pmatrix} 2 \\ 3 \end{pmatrix} \qquad \Sigma = \begin{pmatrix} 2 & 2 \\ 2 & 3 \end{pmatrix}$$

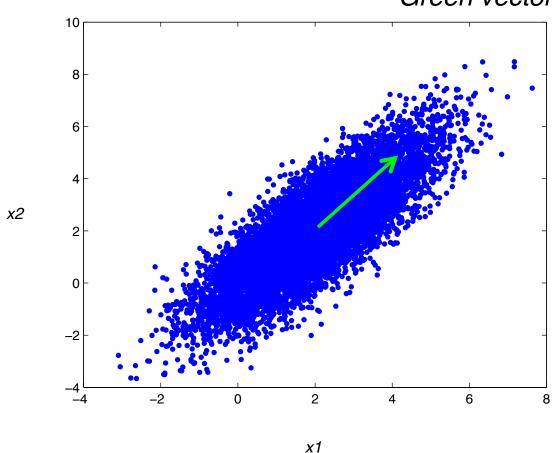
Can you find a vector that would approximate this 2-D space?



PCA - Let's build some intuition

$$\mu = \begin{pmatrix} 2 \\ 3 \end{pmatrix} \qquad \Sigma = \begin{pmatrix} 2 & 2 \\ 2 & 3 \end{pmatrix}$$

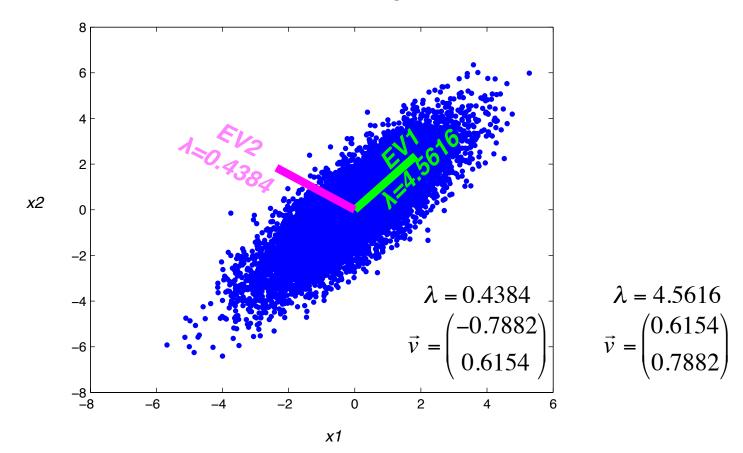
Can you find a vector that would approximate this 2-D space?
Green vector right?



PCA - Let's build some intuition

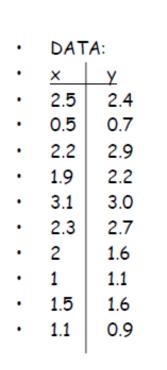
$$\mu = \begin{pmatrix} 0 \\ 0 \end{pmatrix} \qquad \Sigma = \begin{pmatrix} 2 & 2 \\ 2 & 3 \end{pmatrix}$$

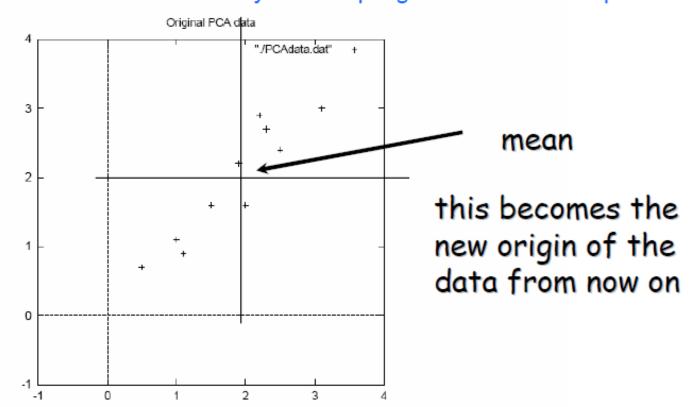
It would be nice to diagonalize the covariance matrix then you have only think about variance Think eigenvectors of covariance matrix



A 2-D Example Step-1

http://kybele.psych.cornell.edu/~edelman/Psych-465-Spring-2003/PCA-tutorial.pdf





A 2-D Example Step-2

Calculate the covariance matrix

Step-3

Calculate the eigenvectors and eigenvalues of the covariance matrix

A 2-D Example

Step-4

Feature Vector

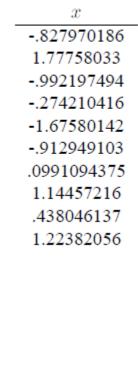
FeatureVector = (eig₁ eig₂ eig₃ ... eig_n) We can either form a feature vector with both of the eigenvectors:

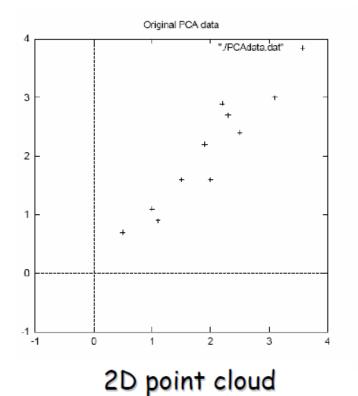
-.677873399 -.735178656 -.735178656 .677873399

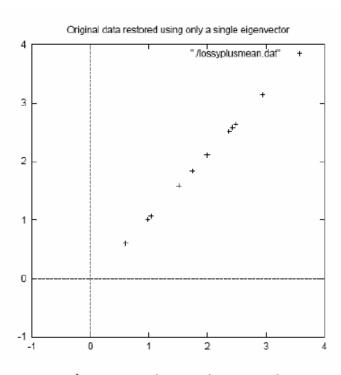
or, we can choose to leave out the smaller, less significant component and only have a single column:

- .677873399 - .735178656

Transformed Data (Single eigenvector)



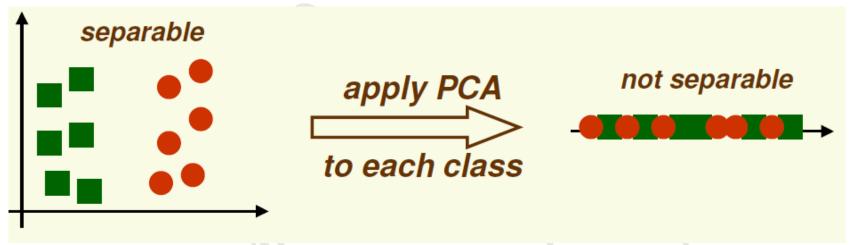




Approximation using one eigenvector basis

Limitations of PCA?

➤ PCA projects the data in lower dimensional space spanned by the directions of maximum variance (most accurate *data*)



 However the directions of maximum variance may be useless for classification (linear...). LDA overcomes this limitation of PCA.

LDA→ Lower dimensions + preserve classification property

Image Source: www.csd.uwo.ca/~oveksler

Bias

- > LDA projects the data in lower dimensional space in such a way that patterns from different classes are well separated.
- ➤ How to find such a line (in 2D), and lower dimensional space in general?

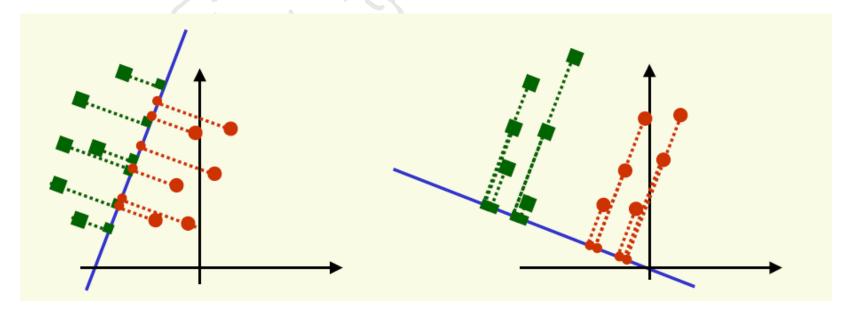
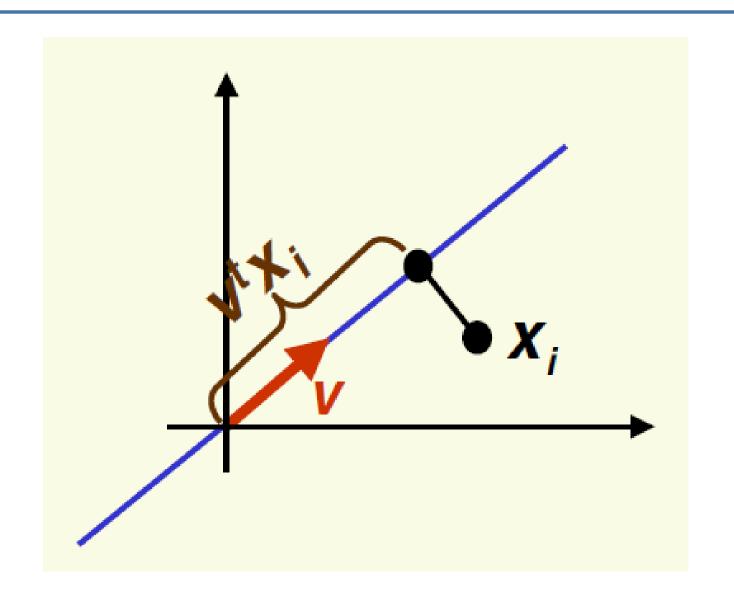


Image Source: www.csd.uwo.ca/~oveksler

Projection of a feature vector on to a space



LDA:

Input; We Rave a binary classification problem, where patterns are coming from a d-dimensional space.

-> n, samples are from class-1.

-> 72 samples are from class-2.

カーナルコニか

Samples are given by vectors

X1, X2, X3 -.., Xn

LDA...

- * Let M1 and M2 be the means of patterns from class-1 and class-2, respectively in original space.
- * Moreover, m, and m2 are mean of batterns after projection.

Then,

$$\tilde{\mu}_{1} = \frac{1}{n_{1}} \sum_{x_{i} \in c_{1}} V^{T} x_{i} = V^{T} \left(\frac{1}{n_{1}} \sum_{x_{i}} x_{i} \right) = V^{T} \mu_{1}$$

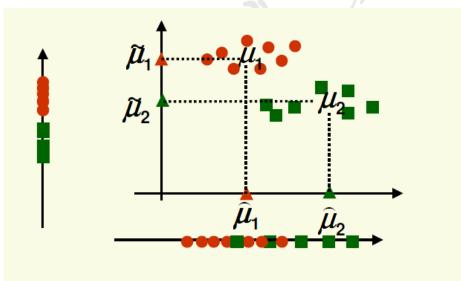
Illy RZ= VTR2

LDA..

Our aim is to have the patterns of these two classes away from each other after projection.

So maximizing distance between their means may be a good measure:

 \rightarrow The larger $|\widetilde{M}_1-\widetilde{M}_2|$, the better is separation.



*The vertical axis
is a better dire ction

* However,

|M_-M_2| > |M_-M_2|

LDA

Therefore, we are missing something. Yes, we are not considering variance of the classes.

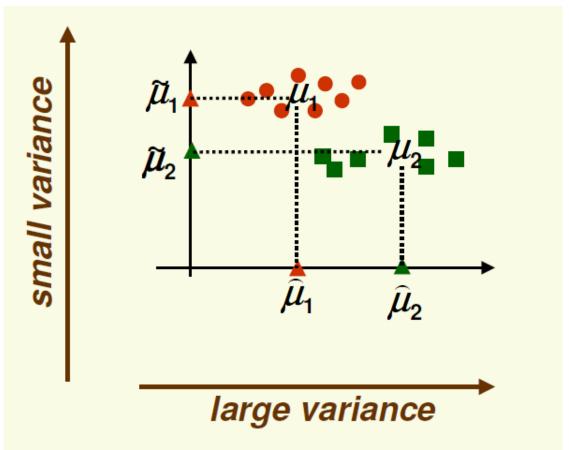


Image Source: www.csd.uwo.ca/~oveksler

LDA...

Define the Scatters as follows:

$$\tilde{S}_{i}^{2} = \sum_{i \in C_{1}} (4i - \tilde{\mu}_{i})^{2}$$
 for class-1

and
$$\widetilde{S}_2 = \sum_{\forall i \in C_2} (\forall i - \widetilde{M}_2)^2$$
 for class-2

where, Yi = VTXi

Then, we need to Maximize $|\tilde{M}_1 - \tilde{M}_2|$ and minimize \tilde{S}_1^2 and \tilde{S}_2^2

Objective function

want projected means are far from each other

$$J(\mathbf{v}) = \frac{(\tilde{\mu}_1 - \tilde{\mu}_2)^2}{\tilde{\mathbf{s}}_1^2 + \tilde{\mathbf{s}}_2^2}$$

want scatter in class 1 is as small as possible, i.e. samples of class 1 cluster around the projected mean $\tilde{\mu}_1$

want scatter in class 2 is as small as possible, i.e. samples of class 2 cluster around the projected mean $\tilde{\mu}_2$

LDA

There fore, Our objective function can be rewritten as Now SBV = > SWV Eigenvalul (52¹23) V = XV Problem

Simplification

$$\left(S_{w}^{-1}S_{B}V=\lambda V\right)$$

* But SBX for any vector X, and (Mi-Mz)
are parallel, i.e.

$$S_{B}X = (M_1 - M_2)(M_1 - M_2)^{T}X = \alpha(M_1 - M_2)$$

$$\longrightarrow \alpha$$

Hences

$$V = S_{w_1}^{-1}(M_1 - M_2) \qquad \qquad (*)$$

Summary: From X1, X2, ..., Xn, calculate Swand (M-M2) => Then & gives you direction of the line.

Example

Consider the following data

$$C_1: [C_{1,2}), (2,3), (3,3), (4,5), (5,5)]^T$$
 $C_2: [C_{1,0}), (2,1), (3,1), (3,2), (5,3), (6,5)]^T$

Then $M = [3 \ 3 \cdot 6]^T$ and $M_2 = [3 \cdot 3 \ 2]^T$

Also,

 $S_1 = 4 * Cov(c_1) = \begin{bmatrix} 10 & 8 \\ 8 & 7 \cdot 2 \end{bmatrix}$
 $S_2 = 4 * Cov(c_2) = \begin{bmatrix} 17 \cdot 3 & 16 \\ 16 & 16 \end{bmatrix}$
 $S_W = S_1 + S_2 = \begin{bmatrix} 27 \cdot 3 & 24 \\ 24 & 23 \cdot 2 \end{bmatrix}$

Example...

Hence,
$$S\bar{w}^{1} = \begin{bmatrix} .39 & -.41 \\ -.41 & .47 \end{bmatrix}$$

Therefore, the optimal line direction V

$$V = S_{w}^{-1}(M_{-}M_{2}) = \begin{bmatrix} -0.79\\ 0.89 \end{bmatrix}$$

