UVA CS 6316/4501 - Fall 2016 Machine Learning

Lecture 18: Principal Component Analysis (PCA)

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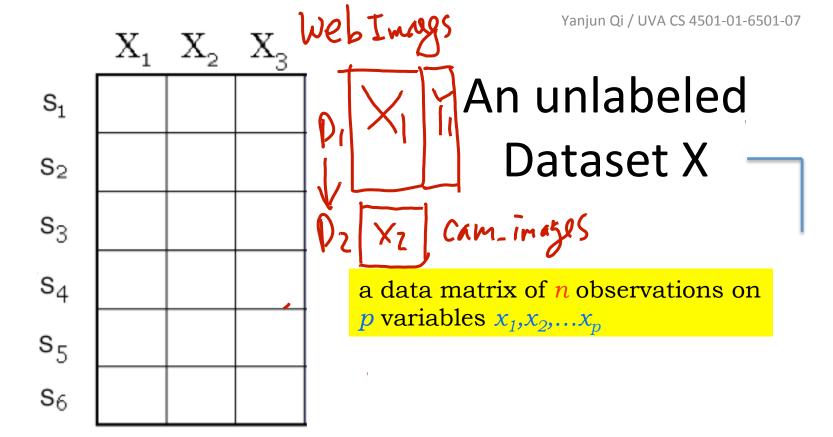
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next after classification? Kmeans + GMM + Hierarchical Spectral WORKING Clustering **KMeans** GMM Basic number of categories **PCA** YES known clustering <10K Randomized NO samples Isomap **PCA** <10K looking Spectral samples YES Embedding WORKING LLE WORKING MiniBatch MeanShift KMeans <10K dimensionality kernel samples **VBGMM** approximation tough reduction predicting structure luck **Bayes- Net HMM**

Where are we? Five major sections of this course

- ☐ Regression (supervised)
- ☐ Classification (supervised)
 - ☐ Feature selection
- Unsupervised models
- Dimension Reduction (PCA)
 - ☐ Clustering (K-means, GMM/EM, Hierarchical)
 - Learning theory
 - ☐ Graphical models



Unsupervised learning = learning from raw (unlabeled, unannotated, etc) data, as opposed to supervised data where a classification/regression label of examples is given

- Data/points/instances/examples/samples/records: [rows]
- Features/attributes/dimensions/independent variables/covariates/predictors/regressors: [columns]

Today

 Dimensionality Reduction (unsupervised) with Principal Components Analysis (PCA)



- Review of eigenvalue, eigenvector
- □ How to project samples into a line capturing the variation of the whole dataset
 ➡ Eigenvector / Eigenvalue of covariance matrix
- PCA for dimension reduction
- □ Eigenface → PCA for face recognition

Review: Mean and Variance

Variance:

$$Var(X) = E((X - \mu)^2)$$

– Discrete RVs:

$$V(X) = \sum_{v_i} (v_i - \mu)^2 P(X = v_i)$$

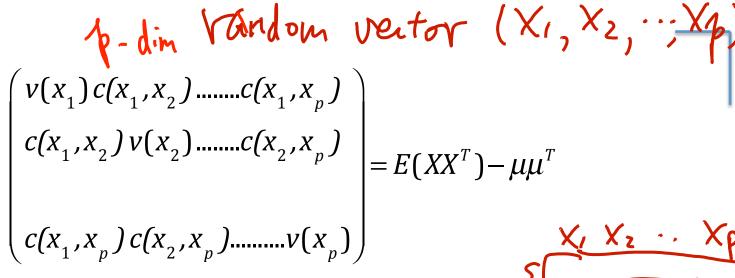
– Continuous RVs:

$$V(X) = \int_{-\infty}^{+\infty} (x - \mu)^2 f(x) dx$$

Covariance:

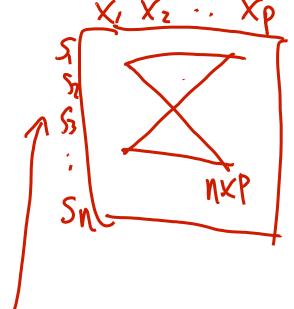
$$Cov(X,Y) = E((X - \mu_{X})(Y - \mu_{Y})) = E(XY) - \mu_{X}\mu_{Y}$$

Review: Covariance matrix



If data is centered, sample covariance matrix is obtained as:

$$C = \frac{1}{n-1} X^T X$$



Review: Eigenvector / Eigenvalue

• The eigenvalues λ_i are found by solving the equation $\mathbf{C}\mathbf{u} = \lambda \mathbf{u}$

$$det(C-\lambda I)=0$$

• Eigenvectors are columns of the matrix U such that

$$C = U D U^T$$

Where

$$O = \begin{bmatrix} 0 & \lambda_2 & \dots & 0 \\ 0 & & & \\ 0 & & & \lambda \end{bmatrix}$$

u≠0

Review: Eigenvalue, e.g.

Let us take two variables with covariance c>0

•
$$\mathbf{c} = \begin{pmatrix} 1 & c \\ c & 1 \end{pmatrix}$$
 $\mathbf{c} - \lambda \mathbf{I} = \begin{pmatrix} 1 - \lambda & c \\ c & 1 - \lambda \end{pmatrix}$

$$det(\mathbf{C}-\lambda \mathbf{I}) = (1-\lambda)^2 - c^2$$

$$\mathbf{C} \mathbf{U} = \lambda \mathbf{U}$$

$$\mathbf{U} \neq \mathbf{0}$$

• Solving this we find $\lambda_1 = 1 + c$

$$\lambda_2 = 1 - c < \lambda_1$$

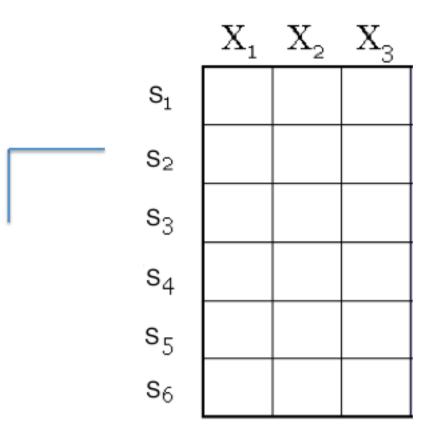
Review: Eigenvector, e.g.

Any eigenvector U satisfies the condition

$$\mathbf{U} = \begin{pmatrix} a_1 \\ a_2 \end{pmatrix} \quad \mathbf{C}\mathbf{U} = \begin{pmatrix} 1 & c \\ c & 1 \end{pmatrix} \begin{pmatrix} a_1 \\ a_2 \end{pmatrix} = \begin{pmatrix} a_1 + ca_2 \\ ca_1 + a_2 \end{pmatrix} = \begin{pmatrix} \lambda a_1 \\ \lambda a_2 \end{pmatrix} \quad \text{with 2}$$
Solving we find
$$\mathbf{U}_1 = \begin{pmatrix} 1/\sqrt{2} \\ 1/\sqrt{2} \end{pmatrix} \cdot \mathbf{U}_2 = \begin{pmatrix} 1/\sqrt{2} \\ -1/\sqrt{2} \end{pmatrix}$$

Today

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An unlabeled Dataset X

a data matrix of n observations on p variables $x_1, x_2, ..., x_p$

- Data/points/instances/examples/samples/records: [rows]
- Features/attributes/dimensions/independent variables/ covariates/predictors/regressors: [columns]

The Goal

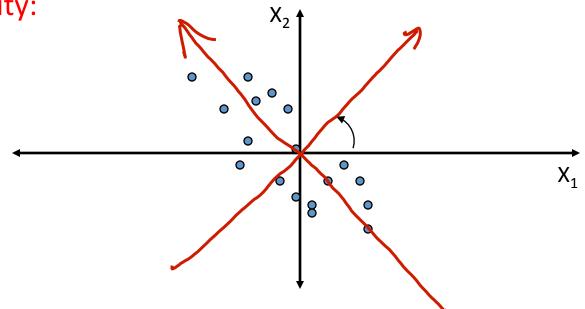
We wish to explain/summarize the underlying variance-covariance structure of a large set of variables through a few linear combinations of these variables.

PCA is introduced by Pearson (1901) and Hotelling (1933)

Trick: Rotate Coordinate Axes

Suppose we have a sample population measured on p random variables $X_1,...,X_p$.

Our goal is to develop a new set of K(K < p) axes (linear combinations of the original p axes) in the directions of greatest variability:



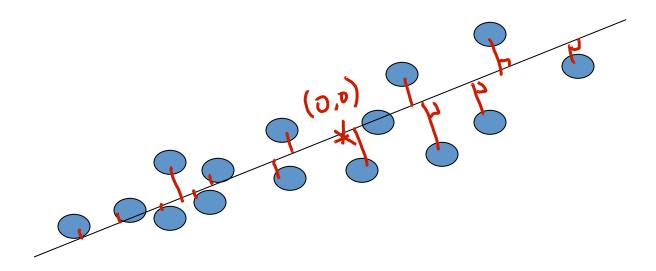
This could be accomplished by rotating the axes (if data is centered).

- Given n points in a p dimensional space,
- for large p, how to project on to a lower-dimensional (K<p) space while preserving broad trends in the data and allowing it to be visualized?

FROM NOW we assume Data matrix is centered:

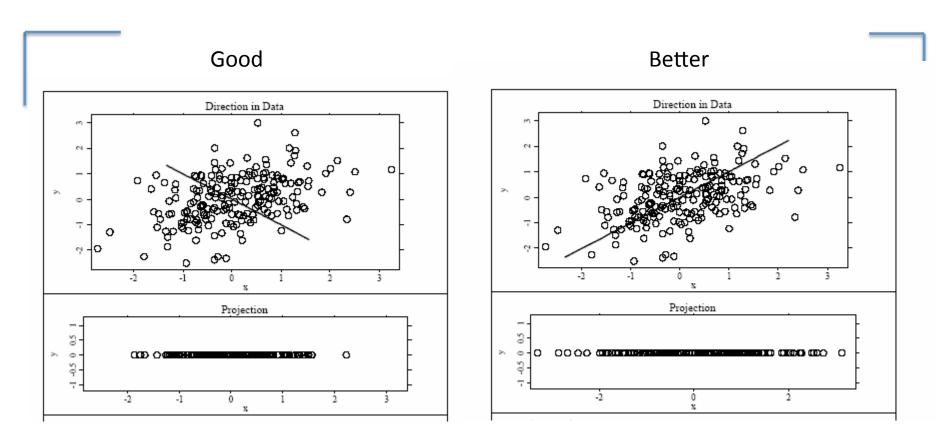
(we subtract the mean along each dimension, and center the original axis system at the centroid of all data points, for simplicity)

• Given n points in a p dimensional space, for large p, how to project on to a 1 dimensional space?

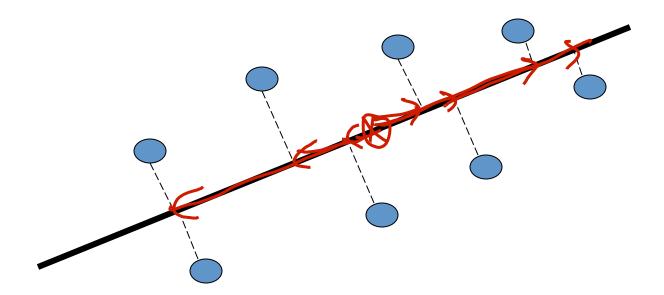


 Choose a line that fits the data so the points are spread out well along the line

Let us see it on a figure



 Formally, to find a line that → Maximizing the sum of squares of data samples' projections on that line



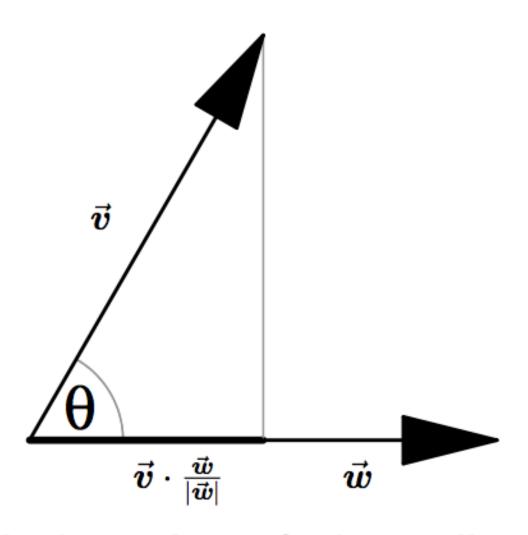


Figure 1: The dot product is fundamentally a projection.

 Formally, to find a line (direction) that → Maximizing the sum of squares of data samples' projections on that line





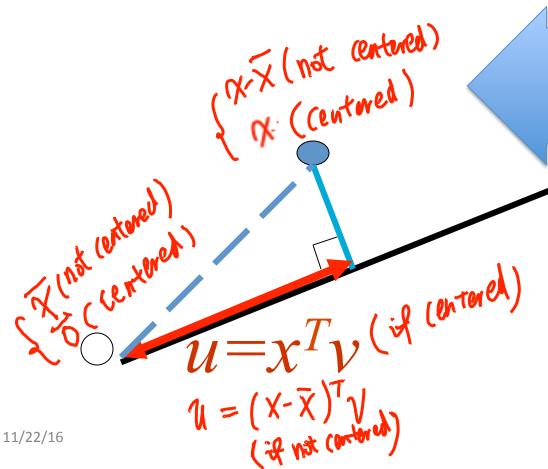
 $\frac{\partial}{\partial t} = \frac{\partial}{\partial t} = \frac{\partial}$

x: p*1 vector

v: p*1 vector

u: 1*1 scalar

 Formally, to find a line (direction) that → Maximizing the sum of squares of data samples' projections on that line



Center X_{n*p} vs. not-centered X_{n*p}

subject to $v^Tv = 1$

x: p*1 vector

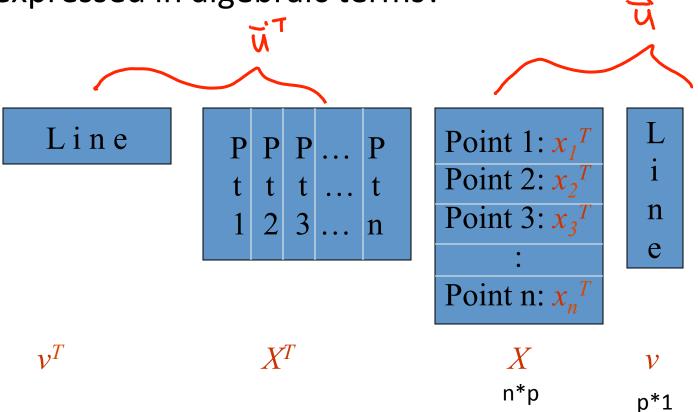
v: p*1 vector

u: 1*1 scalar

arg max
$$\sum_{u_{i}} (u_{i})^{2} \qquad \text{Max} \left\{ \sum_{i=1}^{h} \mathcal{N}_{i}^{2} \right\} =$$

$$= \left[\mathcal{N}_{1} \mathcal{N}_{2}, \mathcal{N}_{3}, \dots, \mathcal{N}_{h} \right] \left[\begin{array}{c} u_{i} \\ u_{2} \\ \vdots \\ u_{h} \end{array} \right] = \left[\begin{array}{c} \mathcal{N}_{1} \\ \mathcal{N}_{2} \\ \vdots \\ \mathcal{N}_{h} \end{array} \right]$$
Here $\mathcal{N}_{i} = \left[\begin{array}{c} u_{1} \\ u_{2} \\ \vdots \\ u_{h} \end{array} \right] = \left[\begin{array}{c} \chi_{1}^{T} \vee \\ \chi_{2}^{T} \vee \\ \vdots \\ \chi_{h}^{T} \vee \end{array} \right] = \left[\begin{array}{c} \chi_{1}^{T} \vee \\ \chi_{2}^{T} \vee \\ \vdots \\ \chi_{h}^{T} \vee \end{array} \right] = \left[\begin{array}{c} \chi_{1}^{T} \vee \\ \chi_{2}^{T} \vee \\ \vdots \\ \chi_{h}^{T} \vee \end{array} \right] = \left[\begin{array}{c} \chi_{1}^{T} \vee \\ \chi_{2}^{T} \vee \\ \vdots \\ \chi_{h}^{T} \vee \end{array} \right] = \left[\begin{array}{c} \chi_{1}^{T} \vee \\ \chi_{2}^{T} \vee \\ \vdots \\ \chi_{h}^{T} \vee \end{array} \right]$

 How is the sum of squares of projection lengths expressed in algebraic terms?



 How is the sum of squares of projection lengths expressed in algebraic terms?

$$max(v^TX^TX^V)$$
, subject to $v^Tv = 1$

Rewriting this:

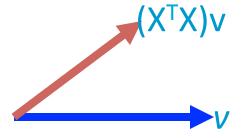
max(
$$v^TX^T Xv$$
), subject to $v^Tv = 1$

$$x^{T}X^{T}Xv = \lambda = \lambda v^{T}v = x^{T}(\lambda v)$$

 $<=> v^{T}(X^{T}Xv - \lambda v) = 0$

- Show that the maximum value of V^TX^TXV is obtained for those vectors /directions satisfying $X^TXV = \lambda V$
- So, find the largest λ and associated u such that the matrix X^TX when applied to u, yields a new vector which is in the same direction as u, only scaled by a factor λ .

 (X^TX)v points in some other direction (different from v) in general



 \rightarrow If u is an eigenvector and λ is corresponding eigenvalue

$$X^TX u = \lambda u$$

So, find the largest λ and associated u such that the matrix X^TX when applied to u, yields a new vector which is in the same direction as u, only scaled by a factor λ .

Algebraic Interpretation – beyond 1D

- How many eigenvectors are there?
- For Real Symmetric Matrices
 - except in degenerate cases when eigenvalues repeat, there are peigenvectors

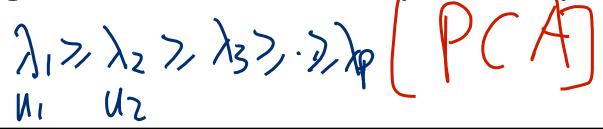
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u_{1,}...,u_{p} are the eigenvectors \lambda_{1}...\lambda_{p} are the eigenvalues, large to small, ordered by its value
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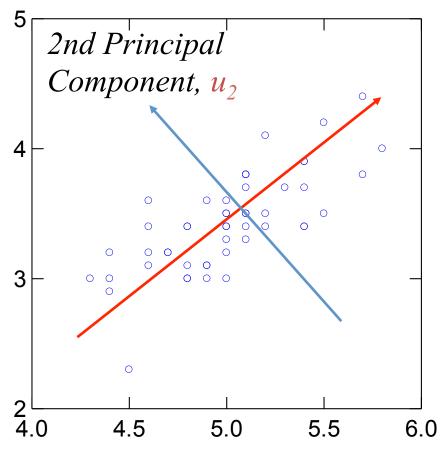
- all eigenvectors are mutually orthogonal and therefore form a new basis space
 - Eigenvectors for distinct eigenvalues are mutually orthogonal
 - Eigenvectors corresponding to the same eigenvalue have the property that any linear combination is also an eigenvector with the same eigenvalue; one can then find as many orthogonal eigenvectors as the number of repeats of the eigenvalue

Algebraic Interpretation – beyond 1D

- For matrices of the form (symmetric) X^TX
 - All eigenvalues are non-negative
 - See Handout-1 "linear algebra review" / Page 18,19,20
 - $-\lambda_1...\lambda_n$ are the eigenvalues, ordering from large to small,
 - i.e. Ordered by the PC's importance

PCA Eigenvectors -> Principal Components



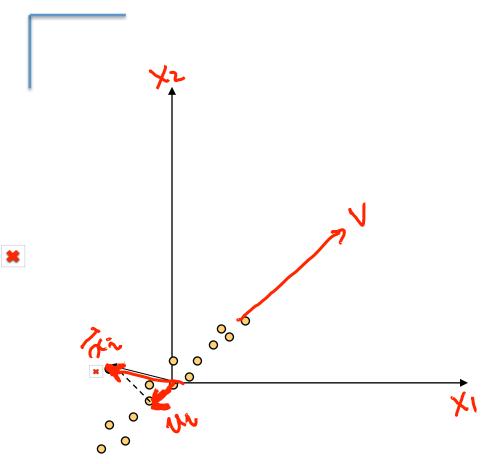


1st Principal Component, *u*₁

PCA 1-D: How the sum of squares of projection lengths relates to VARIANCE?

 In a new coordinate system with v as axis, u is the position of sample x on this axis

PCA 1-D: How the sum of squares of projection fis lengths relates to VARIANCE?



convert **x_i** onto **V**, coordinate

Consider the variation along direction v among all of the orange points {**x_1**, **x_2**,..., **x_n**}:

The variance of positions {u_1, u_2,..., u_n}

PCA 1-D: How the sum of squares of projection lengths relates to VARIANCE?

$$Var(u) = \sum_{u_i} (u_i - \mu)^2 P(u = u_i) = \sum_{u_i} (u_i)^2$$

Assuming centered data matrix

argmax Variance (U)
= argmax Variance

Formally, to find a line (direction v) that

- → Maximizing the sum of squares of data samples' projections on that v line
- → Maximizing the variance of data samples' projected representations on the vaxis

Today

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- PCA for dimension reduction
- □ Eigenface → PCA for face recognition

Applications

Uses:

- Data Visualization
- Data Reduction
- Data Classification
- Trend Analysis
- Factor Analysis
- Noise Reduction

• Examples:

- How many unique "sub-sets" are in the sample?
- How are they similar / different?
- What are the underlying factors that influence the samples?
- How to best present what is "interesting"?
- Which "sub-set" does this new sample rightfully belong?
- **—**

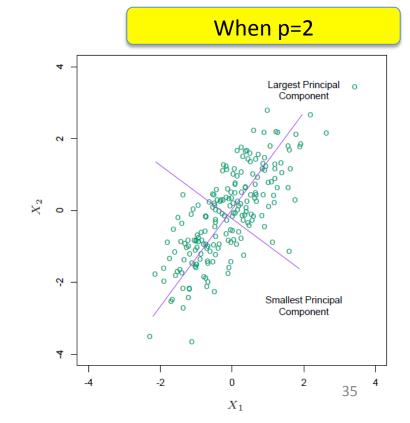
Interpretation of PCA

From p original variables: $x_1, x_2, ..., x_p$:

Produce k new variables: $u_1, u_2, ..., u_k$:

$$u_1 = a_{11}x_1 + a_{12}x_2 + \dots + a_{1p}x_p$$

 $u_2 = a_{21}x_1 + a_{22}x_2 + \dots + a_{2p}x_p$
...



Interpretation of PCA

u_k 's are Principal Components

such that:

 u_k 's are uncorrelated (orthogonal)

 u_1 explains as much as possible of original variance in data set u_2 explains as much as possible of remaining variance

etc.

 k^{th} PC retains the k^{th} greatest fraction of the variation in the samples

Interpretation of PCA

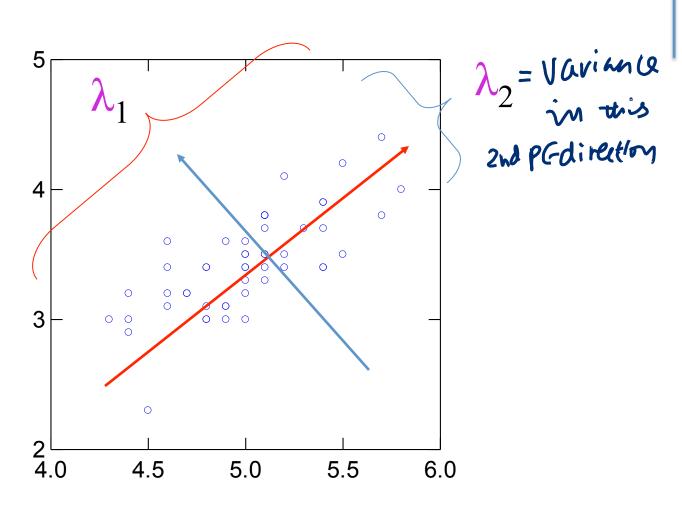
 The new variables (PCs) have a variance equal to their corresponding eigenvalue, since

$$Var(u_i) = u_i^T X^T X u_i = u_i^T \lambda_i u_i = \lambda_i u_i^T u_i = \lambda_i$$
for all $i=1...p$

• Small $\lambda_i \Leftrightarrow$ small variance \Leftrightarrow data change little in the direction of component u_i

PCA is useful for finding new, more informative, uncorrelated features; it reduces dimensionality by rejecting low variance features

PCA Eigenvalues

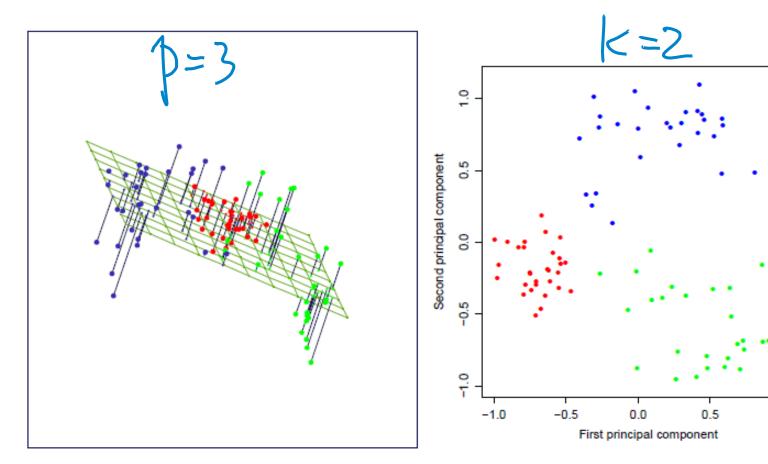


PCA Summary until now

 Rotates multivariate dataset into a new configuration which is easier to interpret

 PCA is useful for finding new, more informative, uncorrelated features; it reduces dimensionality by rejecting low variance features

PCA for dimension reduction e.g. p=3 → (pick top 2 PCs)



corresponds to choosing a "2D linear plane"

1.0

Use PCA to reduce higher dimension

- Suppose each data point is p-dimensional
 - The eigenvectors of data covariance matrix define a new coordinate system
 - We can compress (i.e. perform projection) the data points by only using the top few eigenvectors
 - corresponds to choosing a "linear subspace"
 - represent points on a line, plane, or "hyper-plane"
 - these eigenvectors are known as the *principal components*

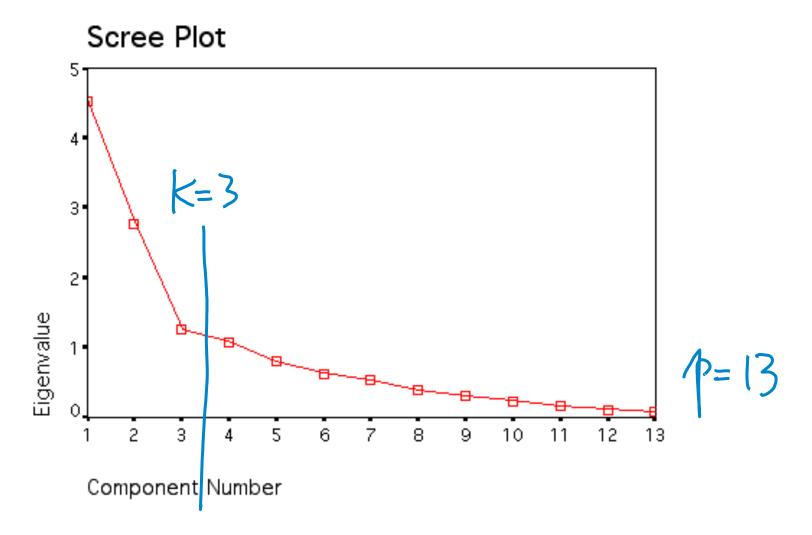
How many components to keep?

 I. Variance: Enough PCs to have a cumulative variance explained by the PCs that is >50-70%

 II. Scree plot: represents the ability of PCs to explain the variation in data, e.g. keep PCs with eigenvalues >1

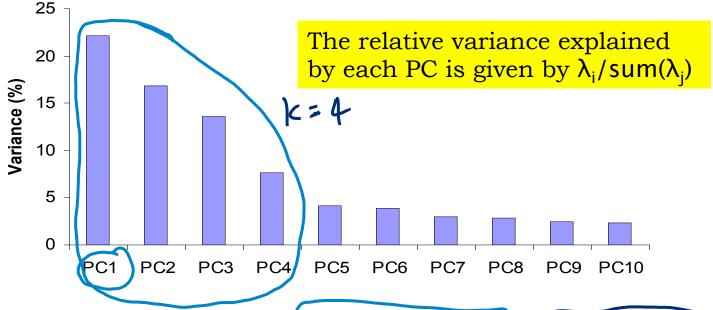
$$Var(N_k) = \lambda k$$

Dimensionality Reduction e.g. check eigenvalue (I)



Dimensionality Reduction(2) e.g. check percentage of kept variance





You do lose some information, but if the eigenvalues are small, you don't lose much

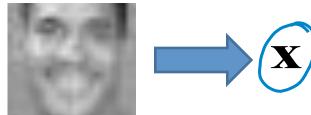
- p dimensions in original data
- Calculate p eigenvectors and eigenvalues
- choose only the first k eigenvectors, based on their eigenvalues
- final projected data set has only k dimensions

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- Eigenface > PCA for face recognition

Example 1: Application to image, e.g. a task of face recognition

1. Treat pixels as a vector



2. Recognize face by 1-nearest neighbor





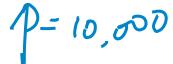
$$\mathbf{y}_1 ... \mathbf{y}_n$$

A face-image database of totally n different people

$$k = \underset{k}{\operatorname{argmin}} \left\| \mathbf{y}_{k}^{T} - \mathbf{x} \right\|$$

Example 1: the space of all face images

- When viewed as vectors of pixel values, face images are extremely high-dimensional
 - 100x100 image ≠ 10,000 dimensions
 - Slow and lots of storage

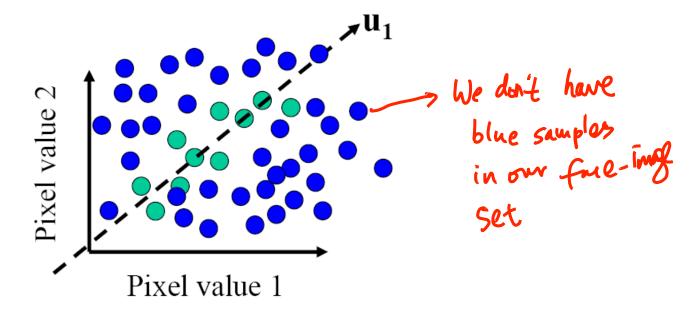


- But very few 10,000-dimensional vectors are valid face images
- We want to effectively model the subspace of face images



Example 1:The space of all face images

 Eigenface idea: construct a low-dimensional linear subspace that best explains the variation in the set of face images



- A face image
- A (non-face) image

Example 1: Application to Faces, e.g. Eigenfaces (PCA on face images)

- 1. Compute covariance matrix of face images
- 2. Compute the principal components ("eigenfaces")
 - K eigenvectors with largest eigenvalues
- 3. Represent all face images in the dataset as linear combinations of eigenfaces
 - Perform nearest neighbors on these projected low-d coefficients

Example 1: Application to Faces

Training images



Example 1: Eigenfaces example

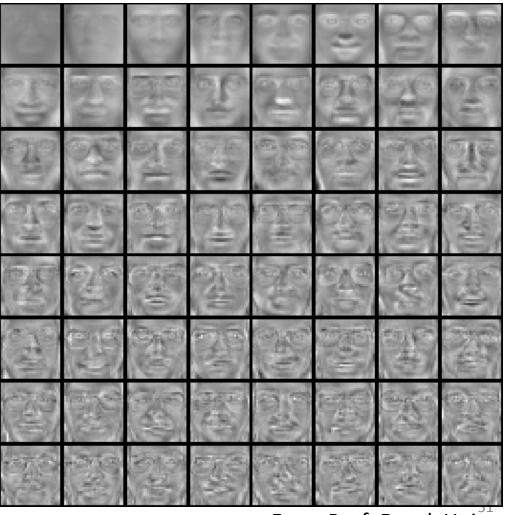
Top eigenvectors: $u_1, ... u_k$

k=64

Mean: µ

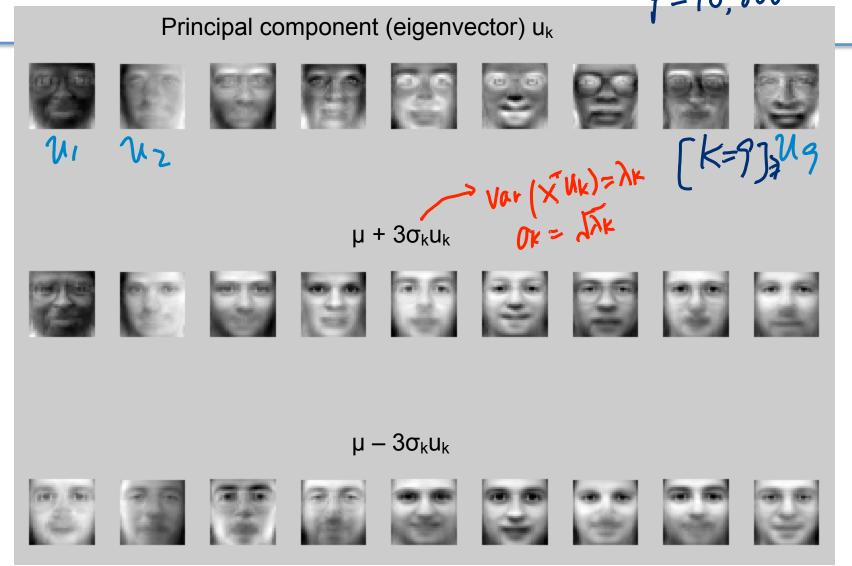


$$\mathbf{X} = \mu = \frac{1}{N} \sum_{k=1}^{N} x_k$$



From Prof. Derek Hoiem

Example 1: Visualization of eigenfaces



Example 1: Representation and reconstruction of original x

• Face **x** in "face space," coordinates:



$$\mathbf{x} \rightarrow [\mathbf{u}_1^{\mathrm{T}}(\mathbf{x} - \mu), \dots, \mathbf{u}_k^{\mathrm{T}}(\mathbf{x} - \mu)]$$

$$g = w_1, \ldots, w_k$$

New representation

Remarkably few eigenvector terms are needed to give a fair likeness of most people's faces.

→ subtract the mean along each dimension, in order to center the original axis system at the centroid of all data points

Representation and reconstruction

Face x in "face space" coordinates:



$$\mathbf{x} \to [\mathbf{u}_1^{\mathrm{T}}(\mathbf{x} - \mu), \dots, \mathbf{u}_k^{\mathrm{T}}(\mathbf{x} - \mu)]$$

$$w_1,\ldots,w_k$$



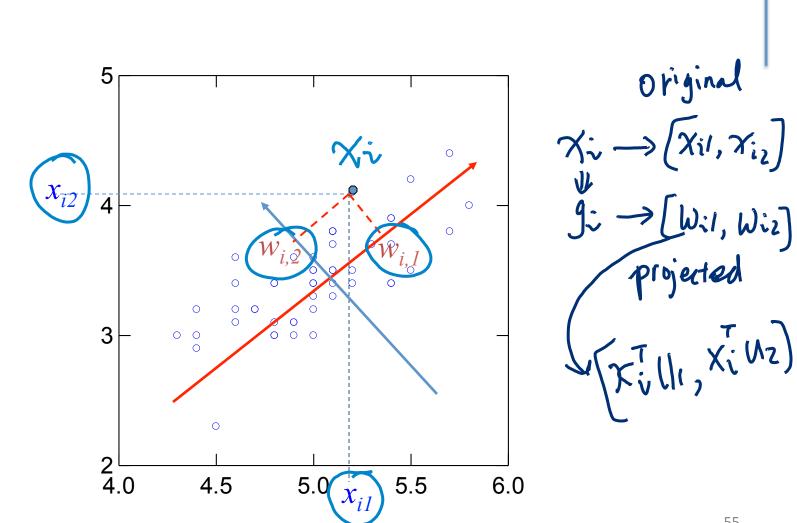




+ $w_1u_1+w_2u_2+w_3u_3+w_4u_4+...+$

A human face may be considered to be a linear combination of these standardized eigen faces

New representation in the lower-dim PC space



Key Property of Eigenspace Representation

Given

- ullet 2 images $\hat{\mathcal{X}}_1,\hat{\mathcal{X}}_2$ that are used to construct the Eigenspace
- $\hat{\mathcal{g}}_1$ is the eigenspace projection of image $\hat{\mathcal{X}}_1$
- \hat{g}_2 is the eigenspace projection of image \hat{x}_2

Then,

$$\|\hat{g}_2 - \hat{g}_1\| \approx \|\hat{x}_2 - \hat{x}_1\|$$

That is, distance in Eigenspace is approximately equal to the distance between two original images.

Classify / Recognition with eigenfaces

Step 1: Process labeled training images

• Find mean μ and covariance matrix

- Find k principal components (i.e. eigenvectors of Σ) \rightarrow $\mathbf{u}_1,...\mathbf{u}_k$
- Project each training image \mathbf{x}_i onto subspace spanned by the top principal components: $(\mathbf{w}_{i1},...,\mathbf{w}_{ik}) = (\mathbf{u}_1^T(\mathbf{x}_i \boldsymbol{\mu}),...,\mathbf{u}_k^T(\mathbf{x}_i \boldsymbol{\mu}))$

Classify / Recognition with eigenfaces

Step 2: Nearest neighbor based face classification

Given a novel image x

- Project onto k PC's subspace: $(\mathbf{w}_1,...,\mathbf{w}_k) = (\mathbf{u}_1^T(\mathbf{x} - \boldsymbol{\mu}), ..., \mathbf{u}_k^T(\mathbf{x} - \boldsymbol{\mu}))$
- Optional: check reconstruction error $\mathbf{x} \hat{\mathbf{x}}$ to determine whether the image is really a face
- Classify as closest training face(s) in the lower k-dimensional subspace

Is this a face or not?

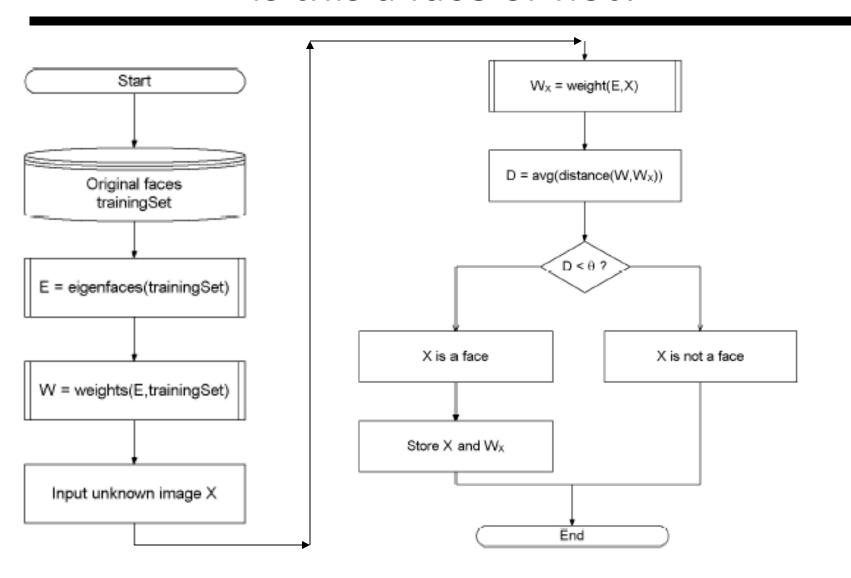


Figure 1: High-level functioning principle of the eigenface-based facial recognition algorithm 59

Example 2: e.g. Handwritten Digits

- 16 x 16 gray scale
- Total 658 such 3's
- 130 is shown \$=756
- Image $x_i : R^{256}$
- Compute principal components

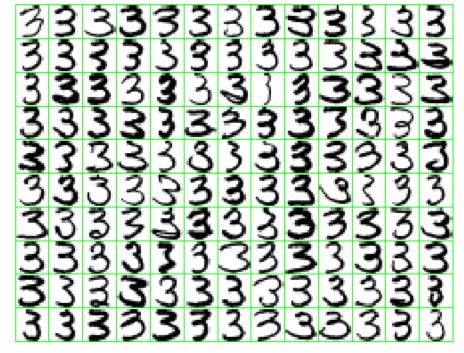
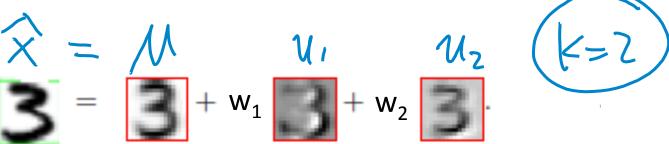


FIGURE 14.22. A sample of 130 handwritten 3's shows a variety of writing styles.



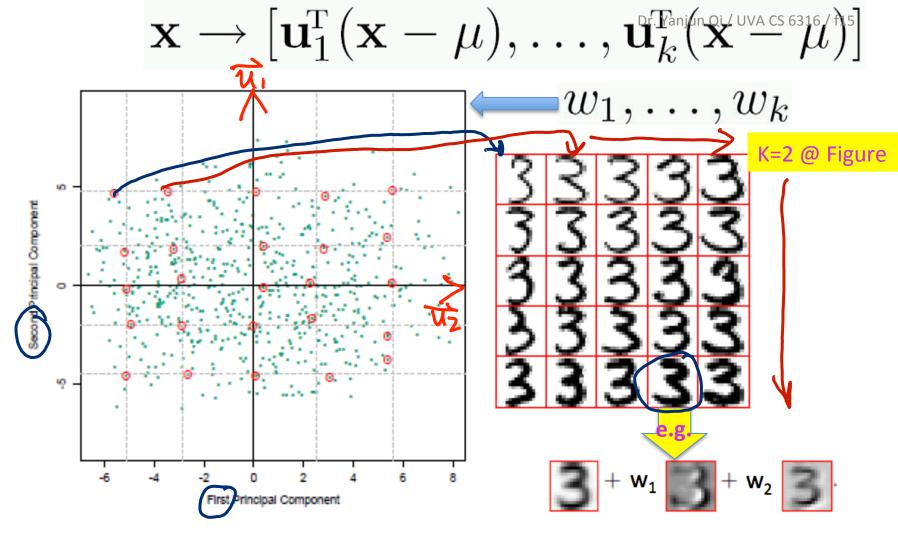


FIGURE 14.23. (Left panel:) the first two principal components of the handwritten threes. The circled points are the closest projected images to the vertices of a grid, defined by the marginal quantiles of the principal components. (Right panel:) The images corresponding to the circled points. These show the nature of the first two principal components.

Principal Components projection of the digits (time 0.02s) The new reduced representation is easier to visualize and interpret e.g. From scikit learn 11

PCA summary

General dimensionality reduction technique

- Preserves most of variance with a much more compact representation
 - Lower storage requirements (eigenvectors + a few numbers per face/sample)
 - Faster matching (since matching within lower-dim)

PCA & Gaussian Distributions.

- PCA is similar to learning a Gaussian distribution for the data.
- μ is the mean of the distribution.
- Then the estimate of the covariance.

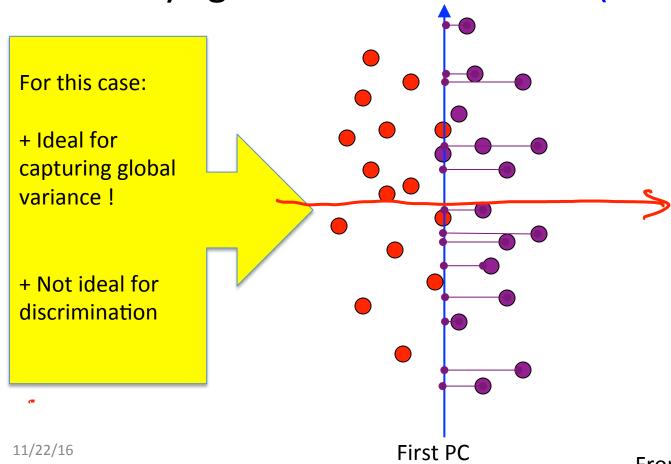
 Dimension reduction occurs by ignoring the directions in which the covariance is small.

(1) Limitations of PCA

- PCA is not effective for some datasets.
- For example, if the data is a set of strings
- (1,0,0,0,...), (0,1,0,0...),...,(0,0,0,...,1) then the eigenvalues do not fall off as PCA requires,

(2) PCA Limitations

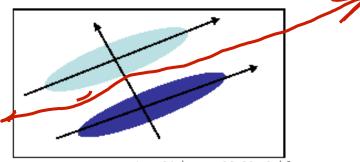
 The direction of maximum variance is not always good for classification (Example 1)



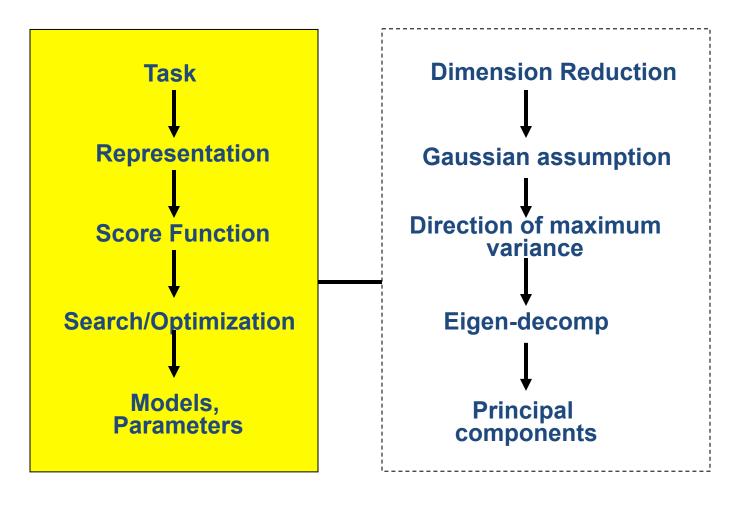
From Prof. Derek Hoiem

PCA and Discrimination

- PCA may not find the best directions for discriminating between two classes. (Example 2)
- Example: suppose the two classes have 2D Gaussian densities as ellipsoids.
- 1st eigenvector is best for representing the probabilities / overall data trend
- 2nd eigenvector is best for discrimination.



Principal Component Analysis



References

- ☐ Hastie, Trevor, et al. *The elements of statistical learning*. Vol. 2. No. 1. New York: Springer, 2009.
 - Dr. S. Narasimhan's PCA lectures
 - ☐ Prof. Derek Hoiem's eigenface lecture



 Subtract the mean from each of the data dimensions.

 Subtracting the mean makes variance and covariance calculation easier by simplifying their equations. The variance and co-variance values are not affected by the mean value.

DAT	ГА:	(p=	:2)
		\ -	— <i>,</i>

<u>x1</u>	x2

2.5 2.4

0.5 0.7

2.2 2.9

1.9 2.2

3.1 3.0

2.3 2.7

2 |1.6

1 |1.1

1.5 1.6

1.1|0.9

ZERO MEAN DATA:

x1	x2

.69 .49

-1.31 | -1.21

.39 | .99

.09 | .29

1.29 | 1.09

.49 | .79

.19 | -.31

-.81 | -.81

-.31 | -.31

-.71 | -1.01

From Dr. S. Narasimhan

Calculate the covariance matrix

 since the non-diagonal elements in this covariance matrix are positive, we should expect that the x1 and x2 variable increase together.

 Calculate the eigenvectors and eigenvalues of the covariance matrix

eigenvalues =

1.28402771 .0490833989

eigenvectors =

-.677873399 -.735178656

-.735178656 .677873399

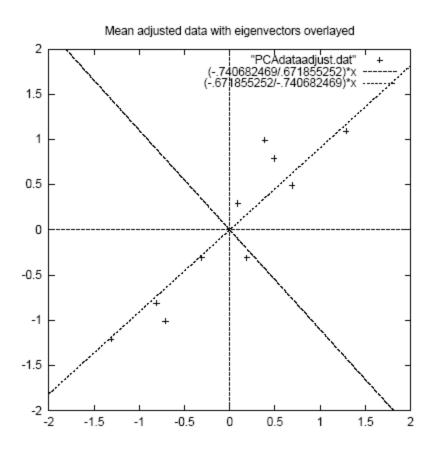


Figure 3.2: A plot of the normalised data (mean subtracted) with the eigenvectors of the covariance matrix overlayed on top.

- eigenvectors are plotted as diagonal dotted lines on the plot.
- Note they are perpendicular to each other.
- •Note one of the eigenvectors goes through the middle of the points, like drawing a line of best fit.
- •The second eigenvector gives us the other, less important, pattern in the data, that all the points follow the main line, but are off to the side of the main line by some amount.

Reduce dimensionality and form feature vector
 the eigenvector with the highest eigenvalue is the principle component of the data set.

In our example, the eigenvector with the largest eigenvalue was the one that pointed down the middle of the data.

Once eigenvectors are found from the covariance matrix, the next step is to order them by eigenvalue, highest to lowest. This gives you the components in order of significance.

Feature Vector

FeatureVector = $(eig_1 eig_2 eig_3 ... eig_n)$

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

Now, if you like, you can decide to *ignore* the components of lesser significance.

You do lose some information, but if the eigenvalues are small, you don't lose much

Deriving the new data

FinalData = RowFeatureVector x RowZeroMeanData

RowFeatureVector is the matrix with the eigenvectors in the columns *transposed* so that the eigenvectors are now in the rows, with the most significant eigenvector at the top

RowZeroMeanData is the mean-adjusted data *transposed*, ie. the data items are in each column, with each row holding a separate dimension.

FinalData transpose: dimensions along columns

w1	w2
827970186	175115307
1.77758033	.142857227
992197494	.384374989
274210416	.130417207
-1.67580142	209498461
912949103	.175282444
.0991094375	349824698
1.14457216	.0464172582
.438046137	.0177646297
1.22382056	162675287

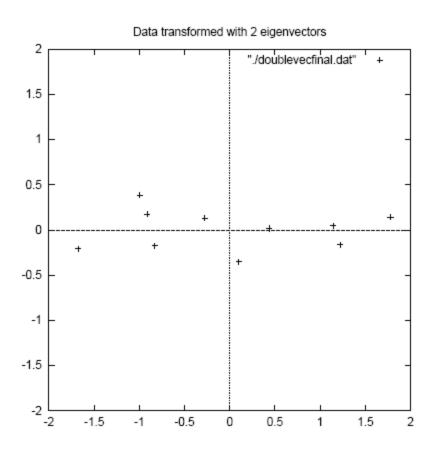


Figure 3.3: The table of data by applying the PCA analysis using both eigenvectors, and a plot of the new data points.

Reconstruction of original Data

 If we reduced the dimensionality, obviously, when reconstructing the data we would lose those dimensions we chose to discard.

 In our example let us assume that we considered only the w1 dimension...

Reconstruction of original Data

w1

-.827970186

1.77758033

-.992197494

-.274210416

-1.67580142

-.912949103

.0991094375

1.14457216

.438046137

1.22382056

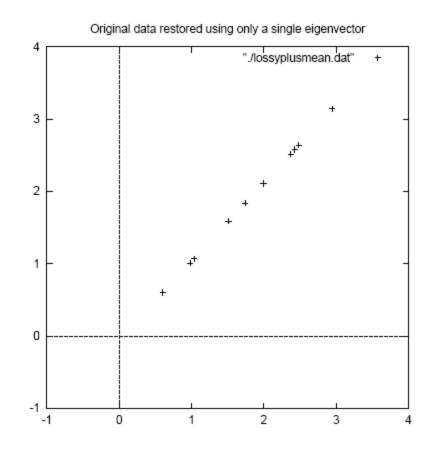


Figure 3.5: The reconstruction from the data that was derived using only a single eigenvector