



# Principal Component Analysis

Machine Learning  
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**Acknowledgement**

Slides courtesy of Manuel Blum

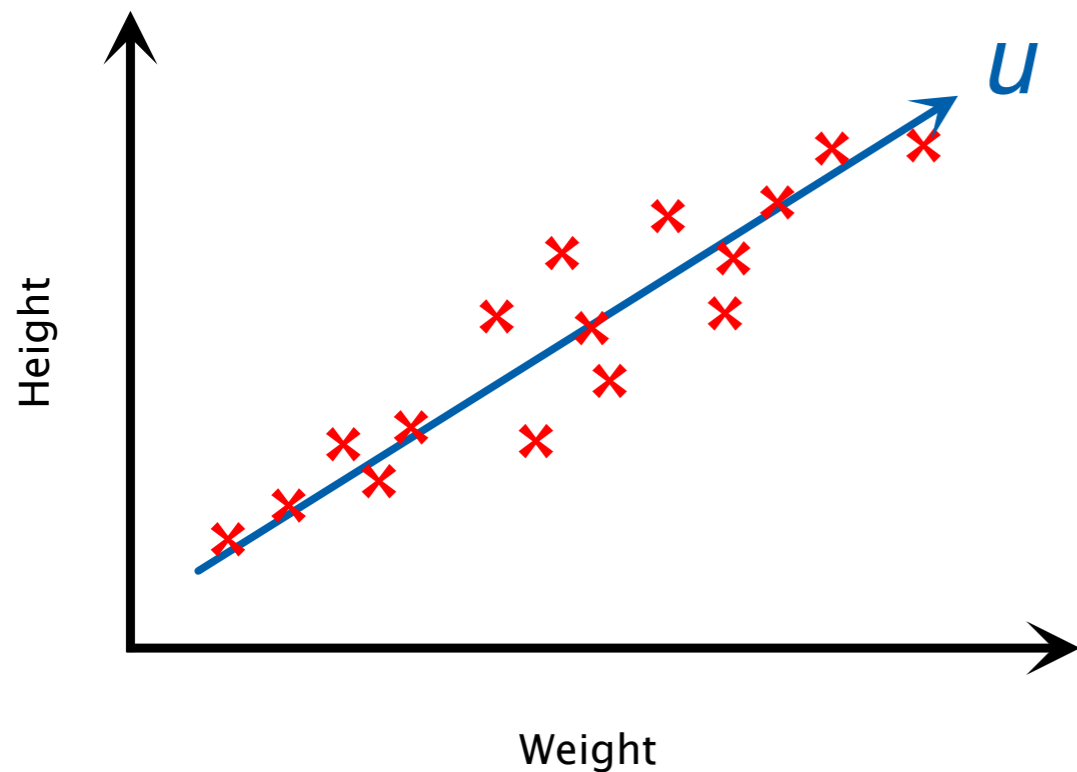
# Motivation

**dimensionality reduction** transforms a  $n$ -dimensional dataset to a  $k$ -dimensional dataset with  $k < n$

- **dataset compression**
  - less memory storage consumption
  - machine learning algorithms run faster on low-dimensional data
- **data visualization**
  - high-dimensional data can be transformed to 2D or 3D for plotting

# Principal Component Analysis

- most commonly used dimensionality reduction method
- projects the data on  $k$  orthogonal bases vectors  $u$  that minimize the projection error



Example:

- original **2D dataset** containing features *weight* and *height*



- projection on vector  $u$

# PCA Algorithm

input:  $x^{(1)}, x^{(2)}, \dots, x^{(m)}$

preprocessing:

- **mean normalization**

1. compute mean of each feature  $j$

$$\mu_j = \frac{1}{m} \sum_{i=1}^m x_j^{(i)}$$

2. subtract the mean from data

$$x_j^{(i)} \leftarrow x_j^{(i)} - \mu_j$$

- **feature scaling**

$$x_j^{(i)} \leftarrow a_j x_j^{(i)}$$

# PCA Algorithm

compute **covariance matrix**  $\Sigma = \frac{1}{m} \sum_{i=1}^m x^{(i)} x^{(i)T}$

diagonalize covariance matrix (using SVD)

$$S = U^{-1} \Sigma U$$

$U$  is the matrix of **Eigenvectors**

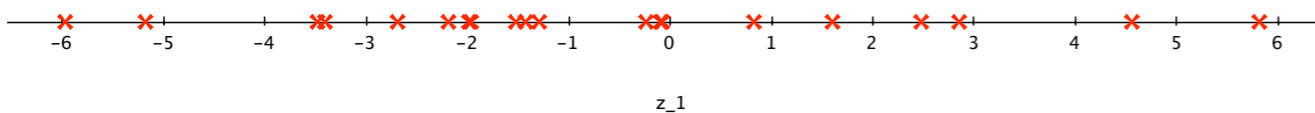
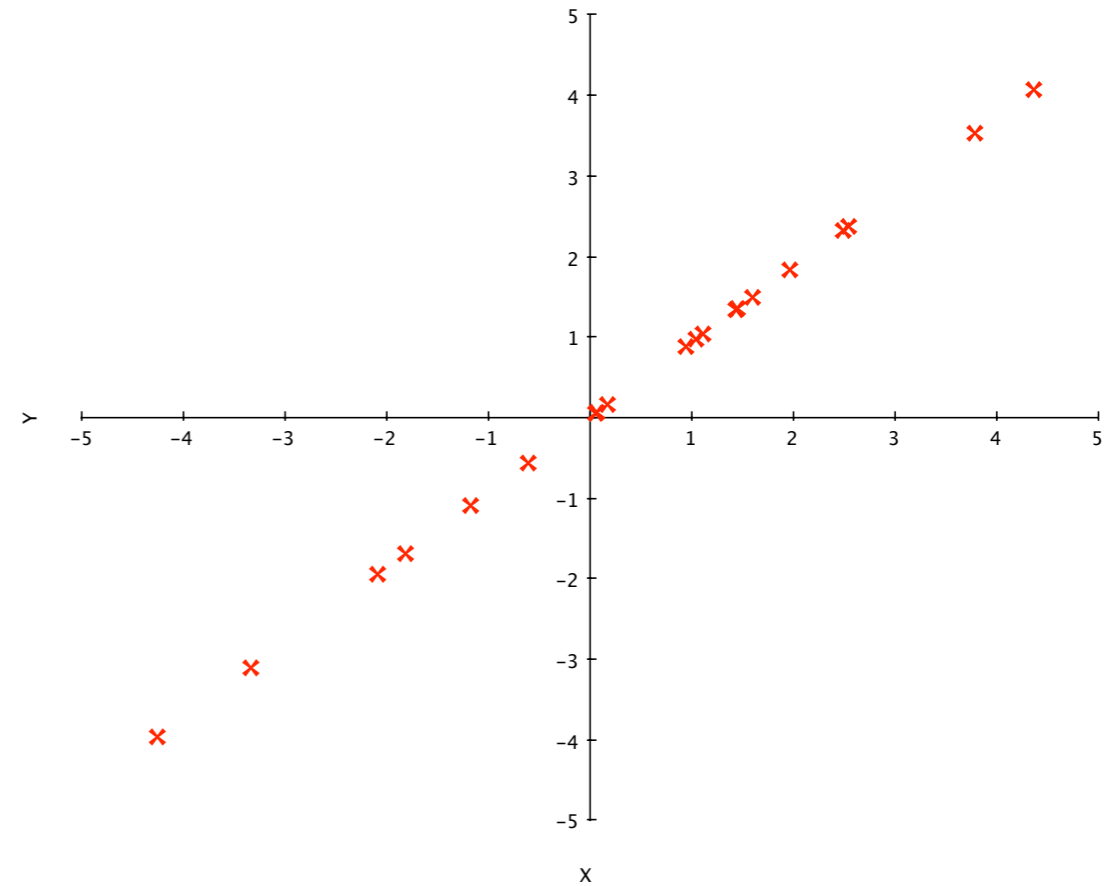
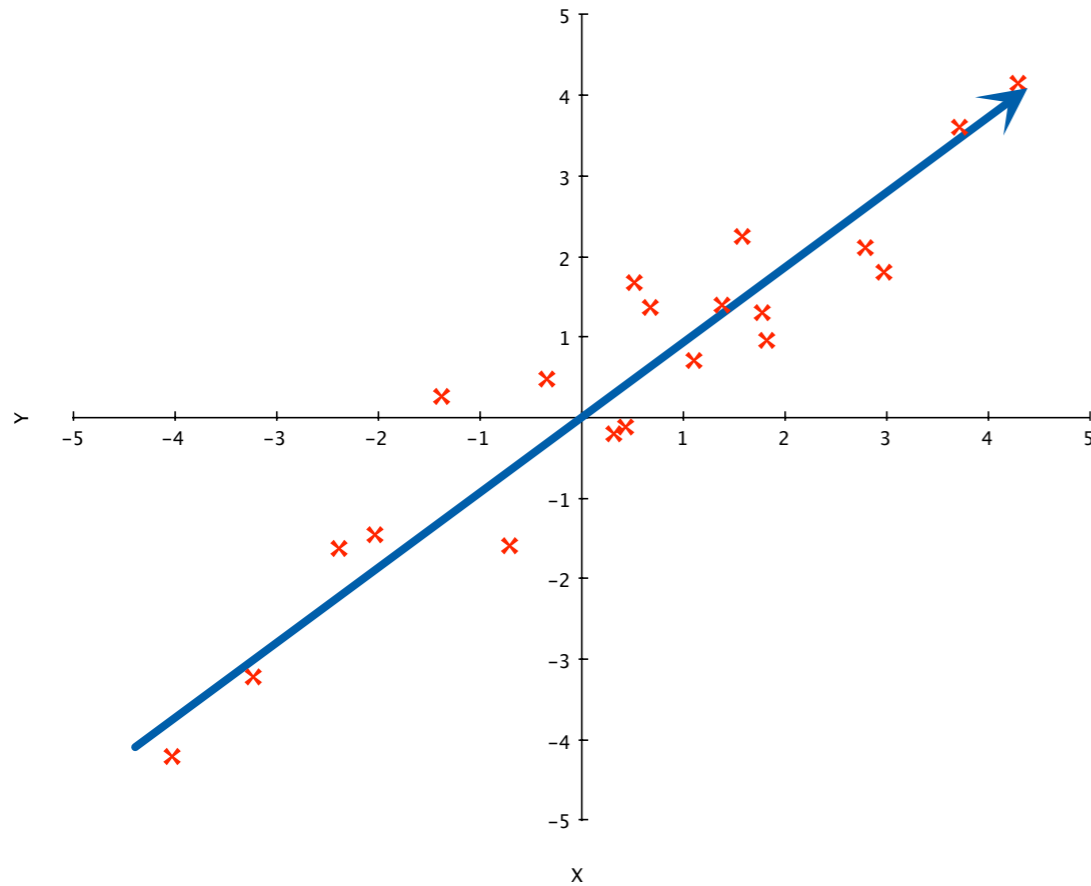
$S$  a diagonal matrix containing the **Eigenvalues**

dimensionality reduction from  $n$  to  $k$  dimensions:

project the data onto the Eigenvectors corresponding to the  $k$  largest Eigenvalues

$$z^{(i)} = U_{reduce}^T x^{(i)}$$

# Reconstruction



$$x_{approx} = U_{reduce} * z^{(i)}$$

$$z^{(i)} = U_{reduce}^T x^{(i)}$$

the reconstruction of compressed data points is an approximation of the original data

# Choosing $k$

average squared projection error:

$$\frac{1}{m} \sum_{i=1}^m \left\| x^{(i)} - x_{approx}^{(i)} \right\|^2$$

total variation in the data:

$$\frac{1}{m} \sum_{i=1}^m \left\| x^{(i)} \right\|^2$$

to retain 99% of the variance, choose  $k$  to be the smallest value, such that

$$\frac{\frac{1}{m} \sum_{i=1}^m \left\| x^{(i)} - x_{approx}^{(i)} \right\|^2}{\frac{1}{m} \sum_{i=1}^m \left\| x^{(i)} \right\|^2} = 1 - \frac{\sum_{i=1}^k S_{ii}}{\sum_{i=1}^n S_{ii}} \leq 0.01$$

$$\frac{\sum_{i=1}^k S_{ii}}{\sum_{i=1}^n S_{ii}} \geq 0.99$$

# Example using Real-world Data



<http://archive.ics.uci.edu/ml/>

- offers 223 datasets
- datasets can be used for the evaluation of ML methods
- results can be compared to those of other researchers



# Iris Data Set

*Download:* [Data Folder](#), [Data Set Description](#)

**Abstract:** Famous database; from Fisher, 1936



<b>Data Set Characteristics:</b>	<b>Multivariate</b>	<b>Number of Instances:</b>	<b>150</b>	<b>Area:</b>	<b>Life</b>
<b>Attribute Characteristics:</b>	<b>Real</b>	<b>Number of Attributes:</b>	<b>4</b>	<b>Date Donated</b>	<b>1988-07-01</b>
<b>Associated Tasks:</b>	<b>Classification</b>	<b>Missing Values?</b>	<b>No</b>	<b>Number of Web Hits:</b>	<b>348488</b>

## Source:

Creator: R.A. Fisher

Donor: Michael Marshall ([MARSHALL%PLU '@' io.arc.nasa.gov](mailto:MARSHALL%PLU '@' io.arc.nasa.gov))

## Data Set Information:

This is perhaps the best known database to be found in the pattern recognition literature. Fisher's paper is a classic in the field and is referenced frequently to this day. (See Duda & Hart, for example.) The data set contains 3 classes of 50 instances each, where each class refers to a type of iris plant. One class is linearly separable from the other 2; the latter are NOT linearly separable from each other.

Predicted attribute: class of iris plant.

This is an exceedingly simple domain.

This data differs from the data presented in Fishers article (identified by Steve Chadwick, [spchadwick '@' espeedaz.net](mailto:spchadwick '@' espeedaz.net) ). The 35th sample should be: 4.9,3.1,1.5,0.2,"Iris-setosa" where the error is in the fourth feature. The 38th sample: 4.9,3.6,1.4,0.1,"Iris-setosa" where the errors are in the second and third features.

## Attribute Information:

1. sepal length in cm
2. sepal width in cm
3. petal length in cm
4. petal width in cm
5. class: Iris Setosa, Iris Versicolour, Iris Virginica

# PCA on the Iris dataset



given: data matrix  $X$

preprocessing:

- mean normalization
- feature scaling

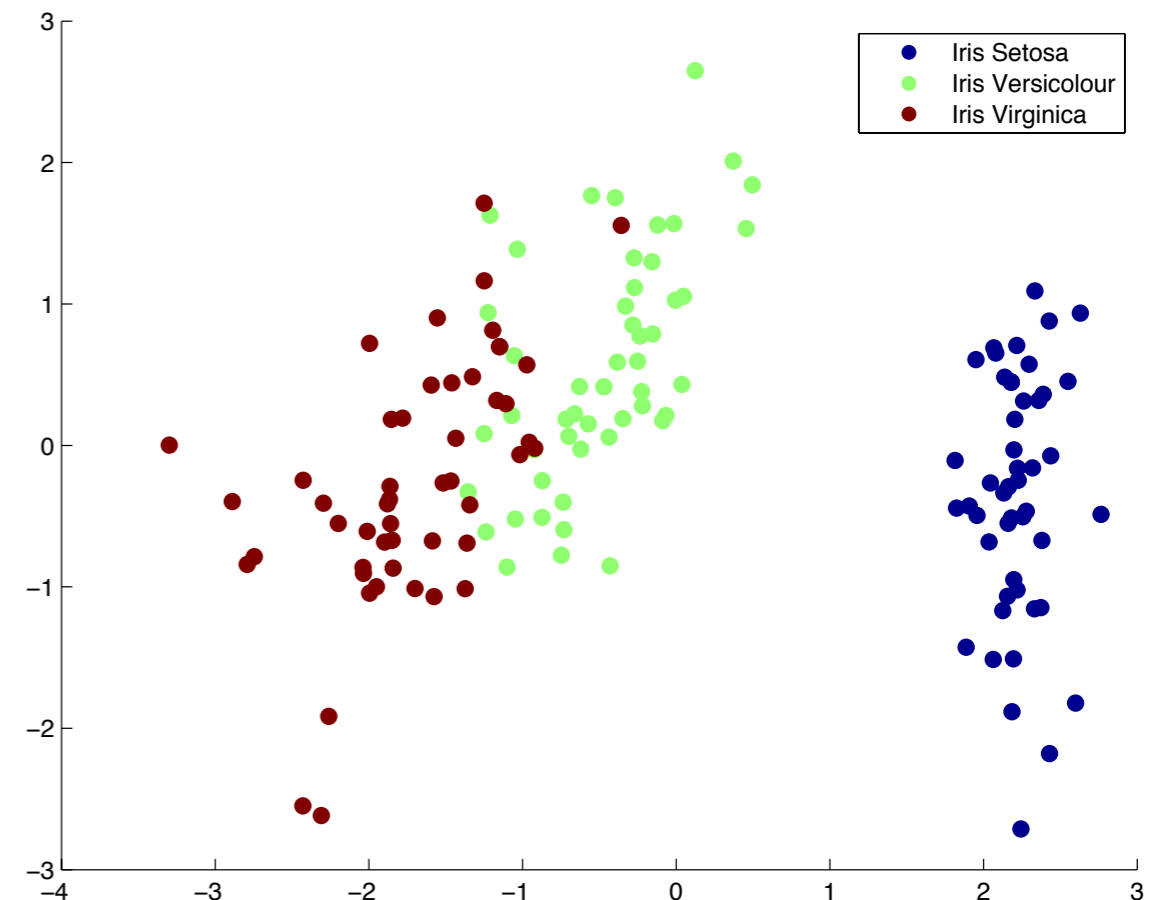
compute covariance matrix:

$$\Sigma = \frac{1}{m} \sum_{i=1}^m x^{(i)} x^{(i)T}$$

compute eigenvectors and eigenvalues:

$$U = \begin{pmatrix} -0.5224 & -0.3723 & 0.7210 & 0.2620 \\ 0.2634 & -0.9256 & -0.2420 & -0.1241 \\ -0.5813 & -0.0211 & -0.1409 & -0.8012 \\ -0.5656 & -0.0654 & -0.6338 & 0.5235 \end{pmatrix}$$

$$S = \begin{pmatrix} 2.8914 & 0 & 0 & 0 \\ 0 & 0.9151 & 0 & 0 \\ 0 & 0 & 0.1464 & 0 \\ 0 & 0 & 0 & 0.0205 \end{pmatrix}$$



reduce  $U$  to  $k$  components

$$z^{(i)} = U_{reduce}^T x^{(i)}$$

# Final Remarks

- PCA can only realize linear transformations
- there exist nonlinear extensions (Kernel PCA)
- PCA-transformed data is uncorrelated
- PCA assumes that most of the information is contained in the direction with the highest variance
- PCA is often used to reduce the noise in a signal
- PCA is an unsupervised method - when used as a preprocessing step for supervised learning the performance can drop significantly

