

# Lecture 14: Feature extraction

## Introduction to Machine Learning

Sophie Robert

L3 MIASHS — Semestre 2

2023-2024

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- Principle
- Algorithm
- Advantages and drawbacks

## 3 t-SNE

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- Example
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# Introduction

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## Question

Do you remember what is the definition of feature extraction/projection ?

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## Question

Do you remember what is the definition of feature extraction/projection ?

- Find a lower dimensional space to project the data in

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## Question

Do you remember what is the definition of feature extraction/projection ?

- Find a lower dimensional space to project the data in
- While keeping as much information as possible

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Many methods exist in the literature:

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Many methods exist in the literature:

- **Linear transformations:** PCA, ...

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Many methods exist in the literature:

- **Linear transformations:** PCA, ...
- **Non-linear transformations:** t-distributed stochastic neighbourhood embedding (**t-SNE**), Uniform Manifold Approximation and Projection (**UMAP**) ...

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# Principal Component Analysis

# Principal Component Analysis

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## Principal Component Analysis

**Principal Component analysis** is a feature projection method that consists in finding a **new coordinate system as a linear combination of the input features** to project the data: this system is orthogonal and a linear combination of the features that **maximizes the variance**.

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## Principal Component Analysis

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The goal of PCA is to:

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## Principal Component Analysis

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The goal of PCA is to:

- Find new axis more relevant to represent the data

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## Principal Component Analysis

**Principal Component analysis** is a feature projection method that consists in finding a **new coordinate system as a linear combination of the input features** to project the data: this system is orthogonal and a linear combination of the features that **maximizes the variance**.

The goal of PCA is to:

- Find new axis more relevant to represent the data
- Give us the importance of each axis

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## Principal Component Analysis

**Principal Component analysis** is a feature projection method that consists in finding a **new coordinate system as a linear combination of the input features** to project the data: this system is orthogonal and a linear combination of the features that **maximizes the variance**.

The goal of PCA is to:

- Find new axis more relevant to represent the data
- Give us the importance of each axis
- Remove unimportant axis and reduce dimension

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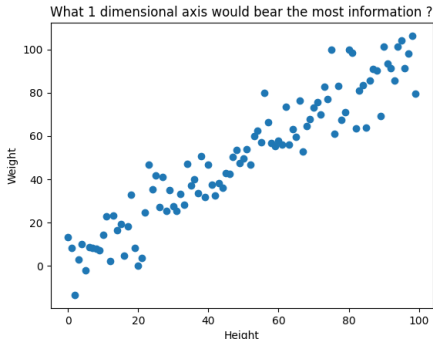
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## Question

What would be an adequate axis to project the data on to reduce to a single axis ?

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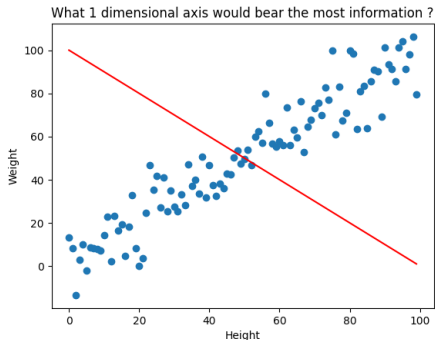
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Possibility:  $y = -x$



## Question

What would a projection on this new axis look like ?

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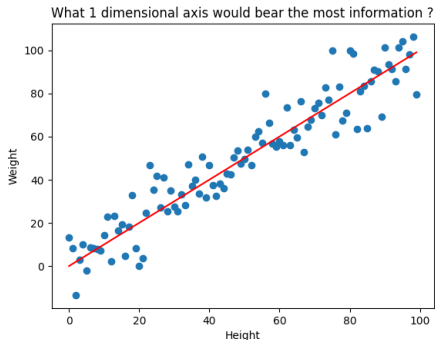
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Possibility:  $y = x$



### Question

What would a projection on this new axis look like ?

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We are looking for new axis that **maximize the variance** and are **uncorrelated**.

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We are looking for new axis that **maximize the variance** and are **uncorrelated**.

It can be shown that these **principal components** are eigenvectors of the data's:

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We are looking for new axis that **maximize the variance** and are **uncorrelated**.

It can be shown that these **principal components** are eigenvectors of the data's:

- Covariance matrix (**Centered PCA**)

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We are looking for new axis that **maximize the variance** and are **uncorrelated**.

It can be shown that these **principal components** are eigenvectors of the data's:

- Covariance matrix (**Centered PCA**)
- Correlation matrix (**Normed PCA**)

# Example dataset

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The **Iris dataset**  $D$ :

sepal length (cm)	sepal width (cm)	petal length (cm)
5.1	3.5	1.4
4.9	3.0	1.4
4.7	3.2	1.3
4.6	3.1	1.5
5.0	3.6	1.4
5.4	3.9	1.7
4.6	3.4	1.4
5.0	3.4	1.5
4.4	2.9	1.4
4.9	3.1	1.5

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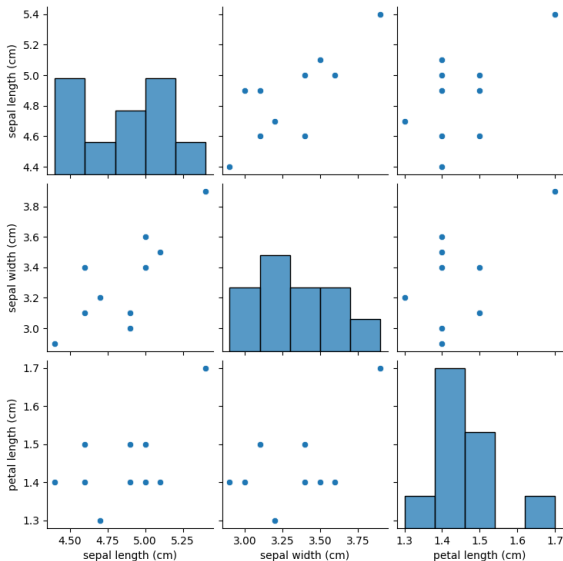
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# Standardize matrix

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For **centered PCA**, center matrix.

For **normed PCA**, standardize matrix (remove mean and divide by standard error):  $Z$ .

# Compute the correlation matrix

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Compute correlation matrix  $C$ .

With  $Z$  the standardized matrix and  $n$  the number of individuals within the dataset,

$$C = \frac{1}{n} Z^t Z$$

	Sepal length	width	Petal length
Sepal length	1.00	0.79	0.60
width	0.79	1.00	0.52
Petal length	0.60	0.52	1.00

# Find eigenvalues and eigenvectors

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Eigen values (sorted) of the correlation matrix are:  
[2.27, 0.51, 0.20]

Eigen vectors matrix  $P$  (sorted by eigen values) is:

Feature	Component 0	Component 1	Component 2
S. length	0.61	-0.26	0.75
Width	0.59	-0.48	-0.65
P. length	0.53	0.84	-0.14

# Project matrix into new space

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Multiply the standardized matrix  $Z$  by the eigenvectors to have the matrix  $Z^*$  in the new projected space  $Z^* = ZP$ : **this is the projection of the individuals in the new feature space.**

ID	Component 0	Component 1	Component 2
0	0.66	-0.95	0.30
1	-0.80	0.07	0.87
2	-1.35	-0.90	0.02
3	-0.74	1.00	-0.31
4	0.64	-1.02	-0.20
5	3.68	0.57	-0.20
6	-0.65	-0.31	-0.83
7	0.75	0.13	0.11
8	-2.11	0.70	-0.26
9	-0.08	0.72	0.51

# Select number of axis

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## Axis importance

The value of each eigenvalue is the **importance of the axis** and is used to select the number of axis to keep as a percentage of the explained variance.

# Select number of axis

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### Axis importance

The value of each eigenvalue is the **importance of the axis** and is used to select the number of axis to keep as a percentage of the explained variance.

Usually, compute the cumulated sum of  $\frac{\lambda_i}{\sum_{i=0}^n \lambda_i}$  for the i-th eigenvalue  $\lambda_i$ : the selected value is the **total explained variance of the dataset**.

# Select number of axis

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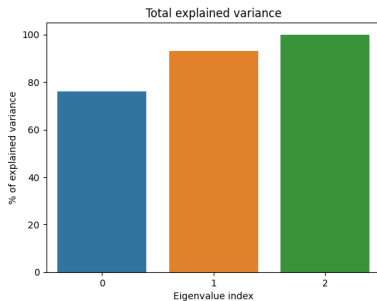
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Here the cumulated sum is:  $[0.76, 0.93, 1]$



We select **two** axis.

# Final projection

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We have our final projection for each individual within the reduced space:

ID	Component 0	Component 1
0	0.66	-0.95
1	-0.80	0.07
2	-1.35	-0.90
3	-0.74	1.00
4	0.64	-1.02
5	3.68	0.57
6	-0.65	-0.31
7	0.75	0.13
8	-2.11	0.70
9	-0.08	0.72

# Interpret results

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A possible interpretation is the plot of the **correlation circle**.

# Interpret results

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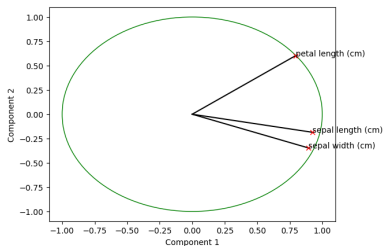
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A possible interpretation is the plot of the **correlation circle**.

## Correlation circle

The correlation circle consists in computing **the correlation of the original features with the new components** and deducing from it the contribution of each variable to the axis.



# Advantages and drawbacks

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## Advantages

- Simple to implement

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## Advantages

- Simple to implement
- Simple to evaluate data loss because of transformation

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## Advantages

- Simple to implement
- Simple to evaluate data loss because of transformation

## Drawbacks

- Features are transformed and lose interpretability of some results

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## Advantages

- Simple to implement
- Simple to evaluate data loss because of transformation

## Drawbacks

- Features are transformed and lose interpretability of some results
- Sensitive to outliers

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# t-distributed Stochastic Neighbor Embedding

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## t-SNE

**t-distributed Stochastic neighbor embedding** (t-SNE) is a nonlinear feature reduction method that consists in associating for each pair of individual **the probability of being close** to each other in the new space by relying on a probability distribution in the original space.

# t-distributed Stochastic Neighbor Embedding

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## t-SNE

**t-distributed Stochastic neighbor embedding** (t-SNE) is a nonlinear feature reduction method that consists in associating for each pair of individual **the probability of being close** to each other in the new space by relying on a probability distribution in the original space.

Given a matrix  $X$  with  $k$  individuals and  $n$  features, find a matrix  $Q$  with  $k$  individuals and  $d$  features, with  $d \ll n$

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We need two probability distributions:

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We need two probability distributions:

- $p_{ij}$ : measures the similarity between the individuals within the initial space.

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We need two probability distributions:

- $p_{ij}$ : measures the similarity between the individuals within the initial space.
- $q_{ij}$ : measures the similarity between the individuals within the new space.

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We need two probability distributions:

- $p_{ij}$ : measures the similarity between the individuals within the initial space.
- $q_{ij}$ : measures the similarity between the individuals within the new space.

The divergence between these probability distributions **needs to be minimized**.

# Kullback-Leibler divergence

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## Kullback-Leibler divergence

The **Kullback-Leibler divergence** is a statistical similarity that measures how one probability distribution  $P$  is different from a distribution  $Q$ .

$$D_{KL}(P||Q) = \sum_{x \in \mathcal{X}} P(x) \log\left(\frac{P(x)}{Q(x)}\right)$$

In the case of t-SNE we want to minimize:

$$KL(P||Q) = \sum_{i \neq j} p_{ij} \log\left(\frac{p_{ij}}{q_{ij}}\right)$$

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We define the conditional probability of  $j$  being the neighbor of  $i$  as a Gaussian around  $x_i$  ( $\sigma_i$  measures the density around  $x_i$  and should be estimated):

$$p_{j|i} = \frac{\exp(-\|\mathbf{x}_i - \mathbf{x}_j\|^2 / 2\sigma_i^2)}{\sum_{k \neq i} \exp(-\|\mathbf{x}_i - \mathbf{x}_k\|^2 / 2\sigma_i^2)}$$

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and set:

$$p_{ij} = \frac{p_{j|i} + p_{i|j}}{2N}$$

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and set:

$$p_{ij} = \frac{p_{j|i} + p_{i|j}}{2N}$$

In the new dimensional space, we set a heavy-tailed Student law:

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and set:

$$p_{ij} = \frac{p_{j|i} + p_{i|j}}{2N}$$

In the new dimensional space, we set a heavy-tailed Student law:

$$q_{ij} = \frac{(1 + \|\mathbf{q}_i - \mathbf{q}_j\|^2)^{-1}}{\sum_k \sum_{l \neq k} (1 + \|\mathbf{q}_k - \mathbf{q}_l\|^2)^{-1}}$$

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and set:

$$p_{ij} = \frac{p_{j|i} + p_{i|j}}{2N}$$

In the new dimensional space, we set a heavy-tailed Student law:

$$q_{ij} = \frac{(1 + \|\mathbf{q}_i - \mathbf{q}_j\|^2)^{-1}}{\sum_k \sum_{l \neq k} (1 + \|\mathbf{q}_k - \mathbf{q}_l\|^2)^{-1}}$$

and use gradient descent to **minimize the KL divergence**.

# Example: Iris dataset

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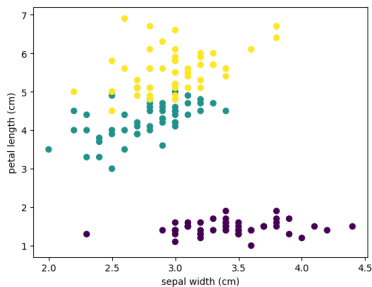
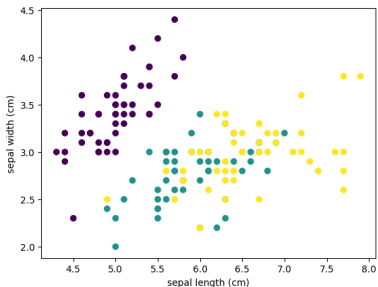
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Data repartition without any transformation:



# Example: Iris dataset

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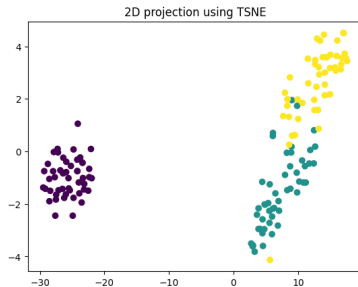
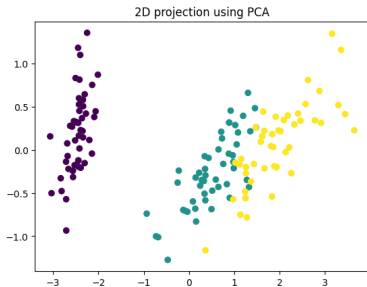
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## Projection in 2-D using t-SNE and PCA:



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## Advantages:

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# Advantages and drawbacks

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- Hard to interpret
- Does not create new variables that can be understood

# Questions

## Lecture 14: Feature extraction

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