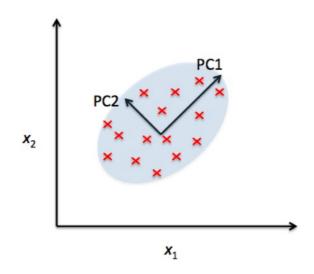


Workshop:
Machine Learning and Prediction Modelling

# Multidimensional Data and Dimensionality Reduction



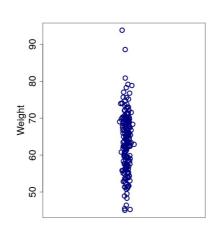
Yannick Rothacher

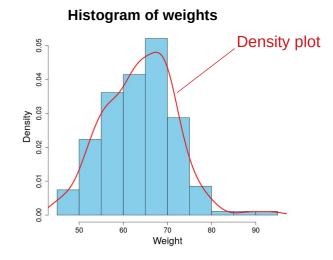
SPF, HS2025



▶ This is the most simple data set possible

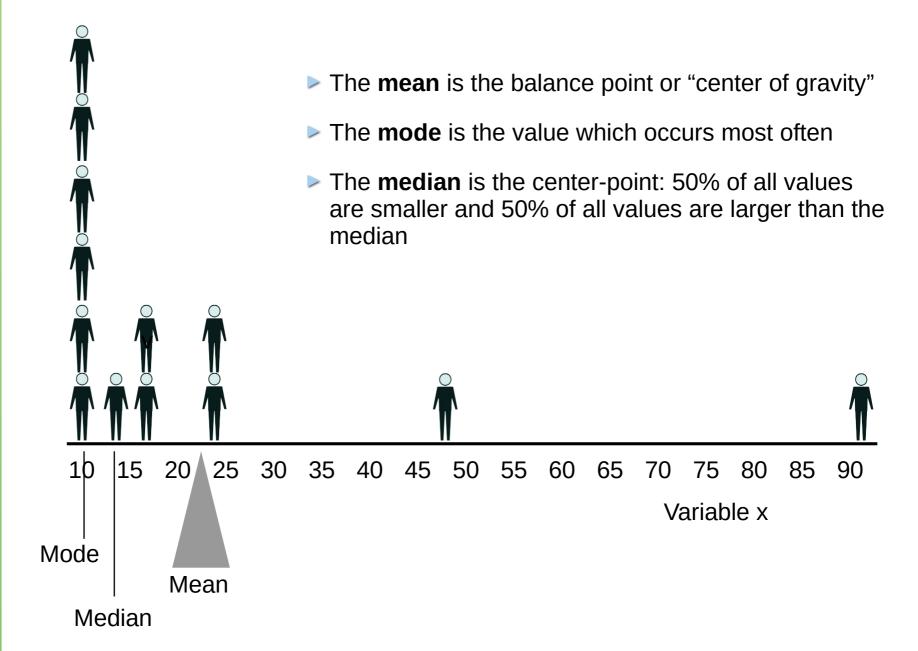
Participant	Weight (kg)
S1	64
S2	80
S3	55
S4	84







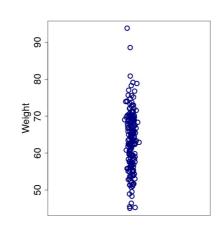
### Recap: Descriptive statistics

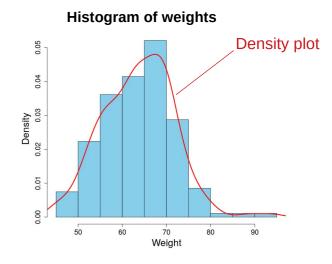




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Participant	Weight (kg)
S1	64
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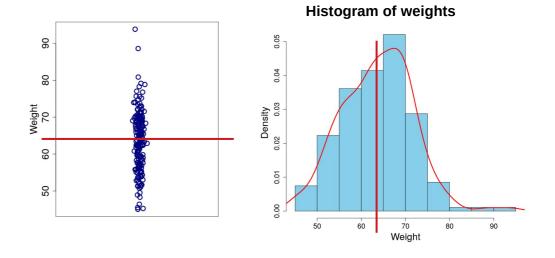






Descriptive statistics

Participant	Weight (kg)
S1	64
S2	80
S3	55
S4	84



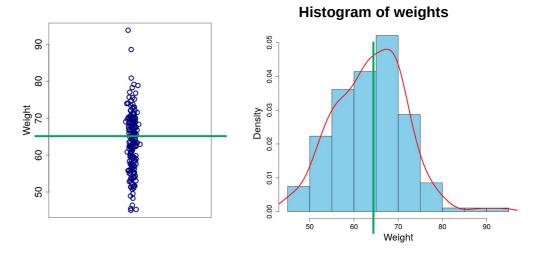
Mean: "Average of a distribution"





Descriptive statistics

Participant	Weight (kg)
S1	64
S2	80
S3	55
S4	84



**Mean**: "Average of a distribution"

$$\bar{x} = \frac{\sum x_i}{n}$$
 63.7 kg

Median: "Midpoint of a distribution"

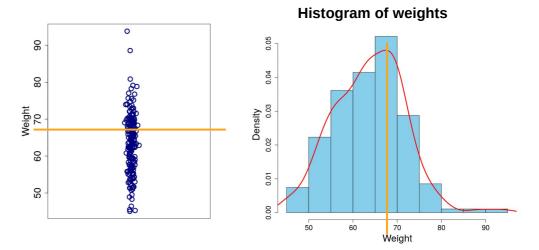
63.9 kg





Descriptive statistics

Participant	Weight (kg)
S1	64
S2	80
S3	55
S4	84



Mean: "Average of a distribution"

$$\bar{x} = \frac{\sum_{i} x_{i}}{n}$$
 63.7 kg

Median: "Midpoint of a distribution"

63.9 kg

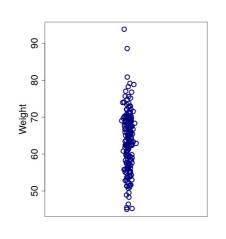
Mode: "The peak(s) of a distribution"

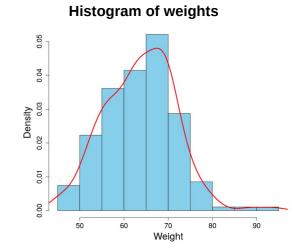
67.4 kg



Descriptive statistics

Participant	Weight (kg)
S1	64
S2	80
S3	55
S4	84





**Mean**: "Average of a distribution"

$$\bar{x} = \frac{\sum x_i}{n}$$
 63.7 kg

Median: "Midpoint of a distribution"

63.9 kg

Mode: "The peak(s) of a distribution"

67.4 kg

Standard deviation: "Spread of distribution"

7.9 kg

$$s = \sqrt{\frac{\sum (x_i - \overline{x})^2}{n - 1}}$$
 (sample standard deviation)

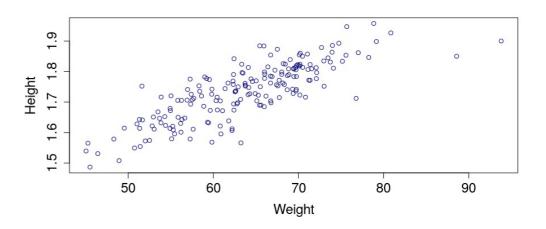


## Multivariate data set

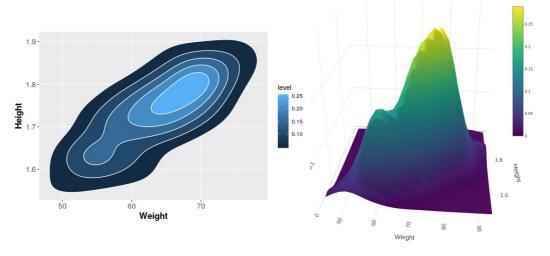
+

Two variables

Participant	Weight (kg)	Height (m)
S1	64	1.71
S2	80	1.82
S3	55	1.65
S4	84	1.84

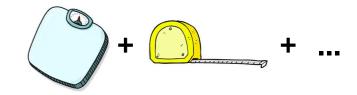


We can still draw a density plot!





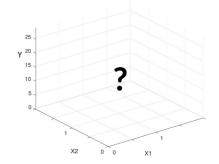
#### Multivariate data set



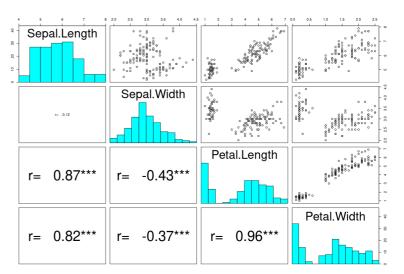
More than two variables

Participant	Weight (kg)	Height (m)	
S1	64	1.71	
S2	80	1.82	
S3	55	1.65	
S4	84	1.84	

Data cannot be visualized anymore...

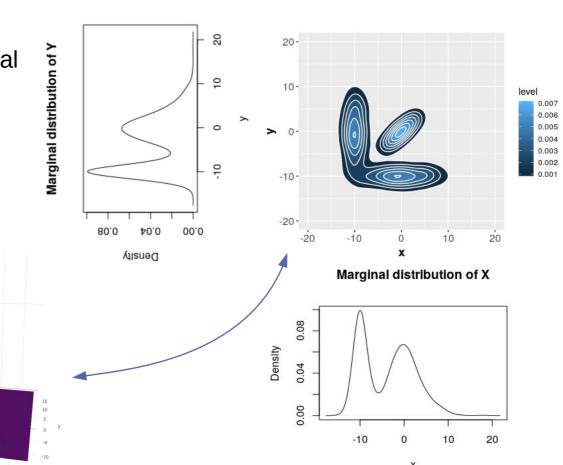


Pairs-plot can help to get an overview of data:





- Does a multivariate distribution have a mode?
  - Yes, the mode is the most probable realization of the multidimensional distribution
- The multidimensional mode is **not necessarily composed** of the marginal distributions' modes!





- Does a multivariate distribution have a mode?
  - Yes, the mode is the most probable realization of the multidimensional distribution
- The multidimensional Mode mode is **not necessarily** 20 20-Marginal distribution of Y composed of the marginal distributions' modes! 10 10level 0.006 0.005 0.004 0.003 0.002 -10-Mode -20-40.0 00.0 -20 10 X Density Marginal distribution of X Density

Component-wise mode

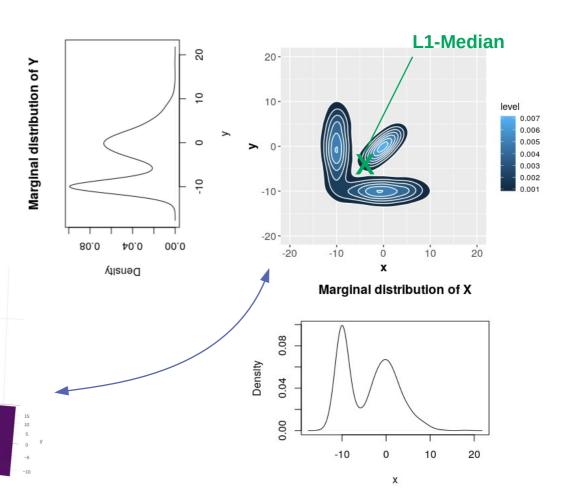
-10

10

20

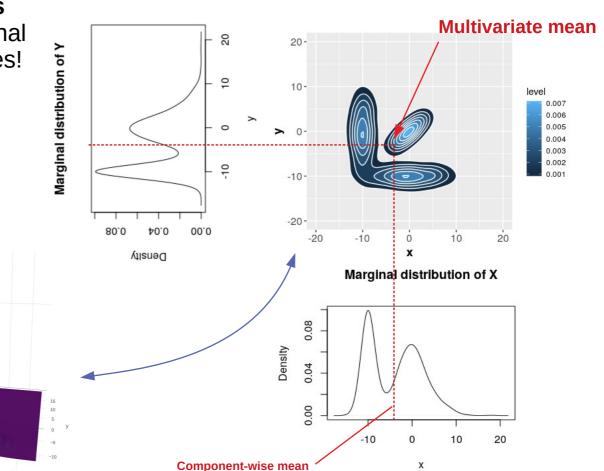


- Does a multivariate distribution have a median?
  - A universally accepted definition of a multivariate median does not exist!
- The "L1-median" is the point with a minimal sum of absolute distances to all other points.
- The L1-median does not have to be composed of the component-wise medians!



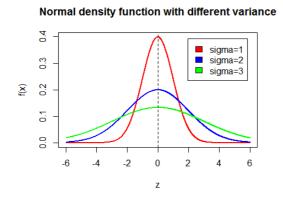


- Does a multivariate distribution have a mean?
  - The multivariate mean is the "balance point" of the distribution
- The multivariate mean is composed of the marginal distributions' mean values!



#### Variance of multivariate distribution

Variance is a measure of the amount of "spread" in a univariate distribution



$$s = \sqrt{\frac{\sum (x_i - \bar{x})^2}{n - 1}}$$
 (sample standard deviation)

The variance is the squared standard deviation:

$$Var = s^2$$

In the case of multivariate distributions the variance is represented in a variance-covariance matrix

$$\mathbf{S}_{pxp} = \begin{pmatrix} \mathbf{X}_{1} & \mathbf{X}_{2} & \mathbf{X}_{p} \\ \mathbf{S}_{1}^{2} & S_{12} & \cdots & S_{1p} \\ \mathbf{S}_{21} & S_{2}^{2} & \cdots & S_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{X}_{p} & S_{p1} & S_{p2} & \cdots & S_{p}^{2} \end{pmatrix}$$

The diagonal elements hold the variances of variables

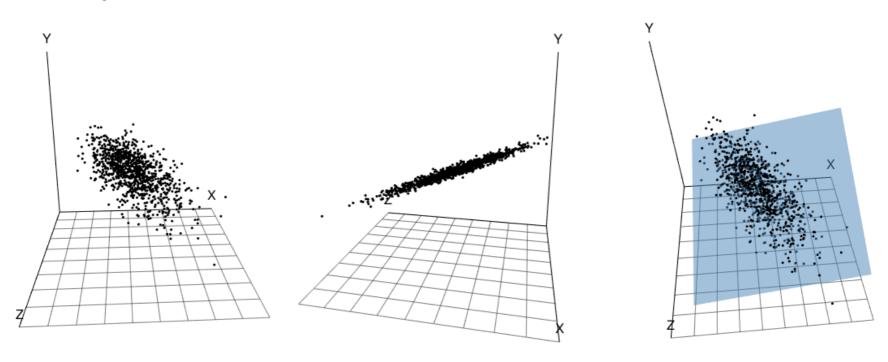
The off-diagonal elements hold the covariance between the respective variables

$$Cov(x,y) = \frac{\sum (x_i - \bar{x})^2 (y_i - \bar{y})^2}{n-1}$$
 (sample covariance)



## Dimensionality reduction

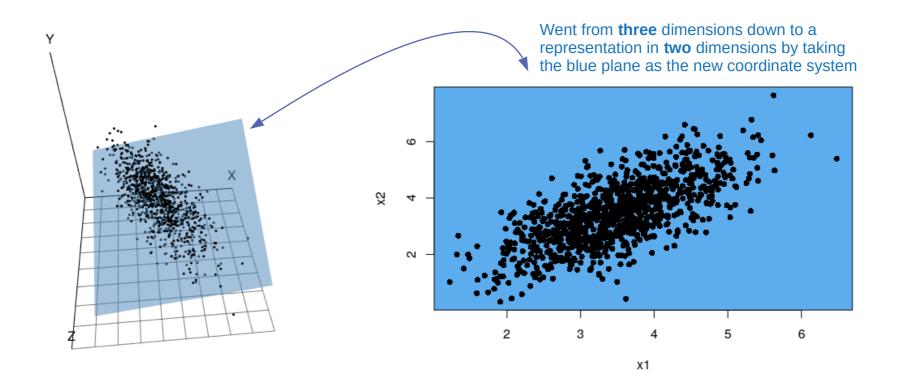
- Is the high number of dimensions really necessary to represent the data?
- Example: The data below is almost lying on a plane
  - The data could also be represented in a two-dimensional coordinate system, without losing a lot of information





## Dimensionality reduction

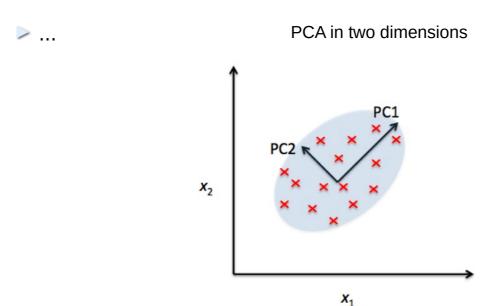
- Is the high number of dimensions really necessary to represent the data?
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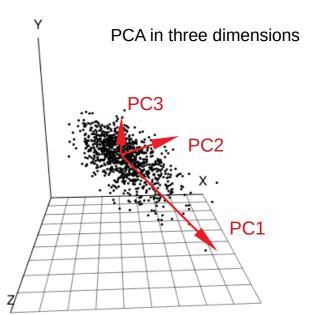




## Principal Component Analysis

- PCA (Principal Component Analysis) is a common method used for dimensionality reduction
- ▶ The idea behind PCA is a rotation of the coordinate system (new axes are principal-component 1, principal-compoinent2, ...)
  - Principal-component 1 is the direction in which the data shows the highest variance
  - Principal-component 2 lies orthogonal to PC1 and is the direction of the second-highest variance

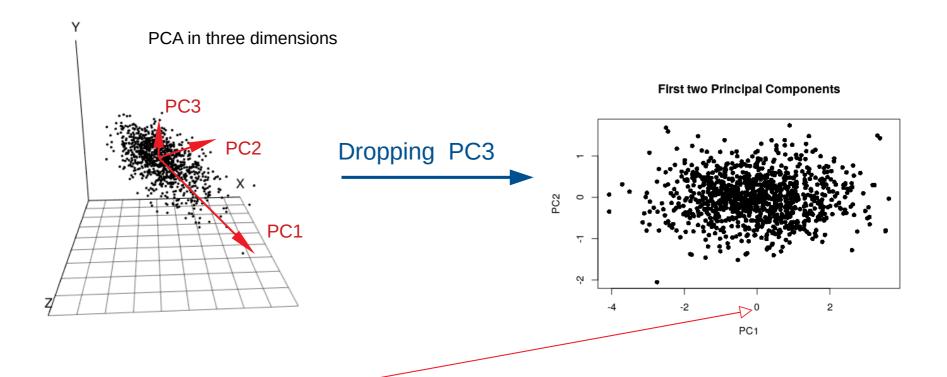






## Principal Component Analysis

▶ In practice, the goal is often to visualize the high-dimensional data in a 2D-plot, by dropping all principal components except the first two:



- ➤ The data is centered (subtracting the mean value from each variable, respectively) when PCA is applied
  - Often the data is also scaled (see later slides)



#### How do we find the PCs?

- After the data has been centered, PCA is just a rotation of the coordinate system (no information is lost)
  - PCA is always possible, one just has to find the right rotation matrix
- It can be shown with linear algebra, that finding the rotation matrix is easier than expected
- The rotation matrix is formed by the eigen-vectors of the variance-covariance-matrix of the centered data
  - In the case of scaled data, the principal components are the eigen-vectors of the correlation matrix

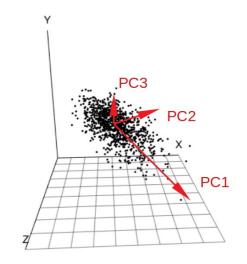


#### PCA in R

PCA can be performed in R using the prcomp() function

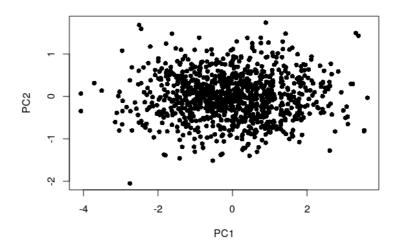
```
> pc <- prcomp(x = myData) # perform PCA
> summary(pc)
Importance of components:
```

PC1 PC2 PC3
Standard deviation 1.270 0.5430 0.08999
Proportion of Variance 0.842 0.1538 0.00423
Cumulative Proportion 0.842 0.9958 1.00000



Plot the first two PCs:

> plot(PC2~PC1, data=pc\$x)





#### PCA in R

▶ PCA can be performed in R using the **prcomp()** function

```
> pc <- prcomp(x = myData) # perform PCA
> pc
Standard deviations (1, .., p=3):
[1] 1.27035807 0.54295340 0.08998992
```

```
Rotation (n x k) = (3 x 3):

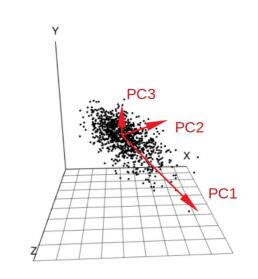
PC1 PC2 PC3

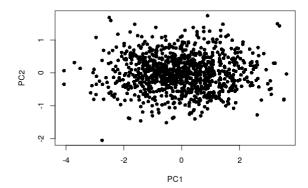
x 0.4918003 0.8525767 0.1767639

y -0.2408321 0.3282877 -0.9133603

z 0.8367391 -0.4066205 -0.3667798
```

shows the contribution of the individual variables to the PCs (also referred to as **variable loadings**)





The first principal component is composed in the following way:

```
PC1 = 0.4918003*x - 0.2408321*y + 0.8367391*z
```

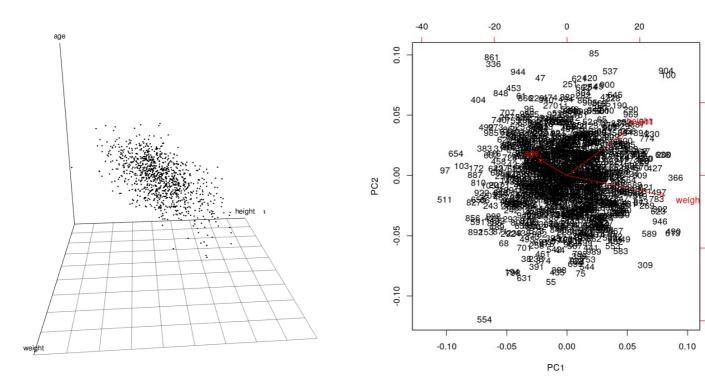
## Biplot – projection of variables

In Biplots the original variables are projected into the PC-coordinate system

20

-20

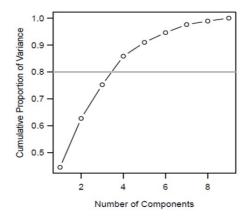
Gives an impression of the contribution of each variable to the PCs



> In R: > pc <- prcomp(x = myData) # perform PCA
> biplot(pc)

## How many PCs are needed?

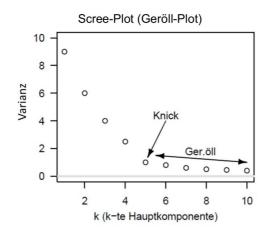
- Take a look at the explained variance by the PCs
- Rule of thumb is that ~80% of total variance should be explained by the first k PCs



proportion of variance explained by the first k principle components:

$$P_k = \frac{\sum_{j=1}^k \text{var}(Y_j)}{V_{total}} \in [0,1]$$

- → Following this rule, we need the **first four PCs** to explain data
- Other method: Check for a "bend" in the Scree-Plot (Geröll-Plot)



#### # In R:

- > screeplot(pca\_object, npcs=10, type="l")
  - → In this case, we need the **first four PCs**, afterwards not much more information is won



#### When to scale the data?

- Often the data is scaled before performing the PCA
- Scaling means that the values of each variable are divided by the variable's standard deviation
  - Now, every variable has unit-variance (variance = 1)
  - Scaling does affect the results of the PCA! (e.g. variable measured in ms vs variable in hours, see covariance-table below)
- In R scaling can be set in the options of the prcomp() command
  - Default is not to scale: prcomp(mydata, scale.=FALSE)
- Scaling is generally advisable when the variables have different scales (e.g. cm, m, kg, ...)
- Scaling is not advised when the variables have the same scale and are comparable with respect to their variability

Example: "Assault" will have a large contribution to PC1 because of its large variance (see covariance matrix)



## Other dimensionality reduction methods

- There are many different methods which can be used for dimensionality reduction, e.g.
  - > PCA
  - Factor analysis
  - t-Distributed Stochastic Neighbor Embedding (t-SNE)
  - Multidimensional Scaling (MDS)
  - Independent Component Analysis (ICA)
  - Feature selection (e.g. by LASSO regression)
  - >