

# Hypothesis Testing as Bivariate Analyses

j3  
– 01.04.26 –



Plateforme OMICS - MIO

Bio-informatique & Sciences de l'Environnement : Exploration de la Diversité Taxonomique des Ecosystèmes par Metabarcoding



Variability in species richness between the 2 stations, depths



Is there a **real significant** difference or  
just a coincidence ?



Using **statistics** to answer your question !!

# Population VS samples

Population: set of individuals or objects of the same kind (very large or infinite)

- We can't study an entire population: in statistics, we study a limited number of individuals, a part of the population: **a sample**
- We try to **deduce properties** of the population from the sample
- If we want to **study the variability** of a variable of interest in the population, we need a **representative sample** (drawn at random)

In a population, we can measure a characteristic: **a variable** that is the result of a random phenomenon.

- Qualitative
- Quantitative (continuous)

A **probability law** describes the random behavior of a phenomenon that depends on chance.

In a population, we can measure a characteristic: **a variable** that is the result of a random phenomenon.

- Qualitative
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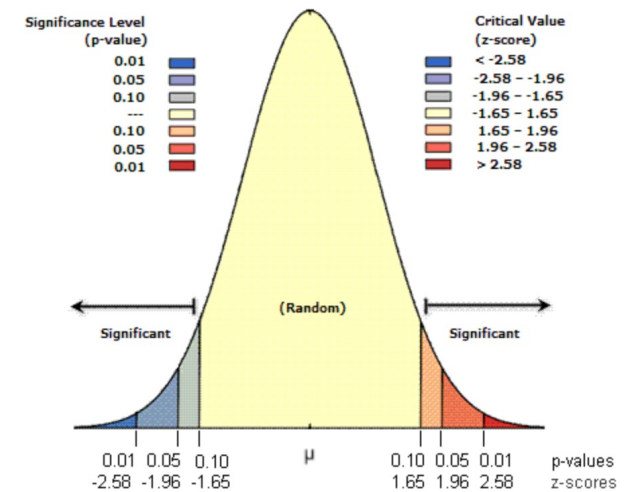
A **probability law** describes the random behavior of a phenomenon that depends on chance.

## THE NORMAL LAW

If we have 1000 samples of a variable following a normal distribution, and plot the number of samples equal to each value, we obtain a "bell" curve / gaussian distribution

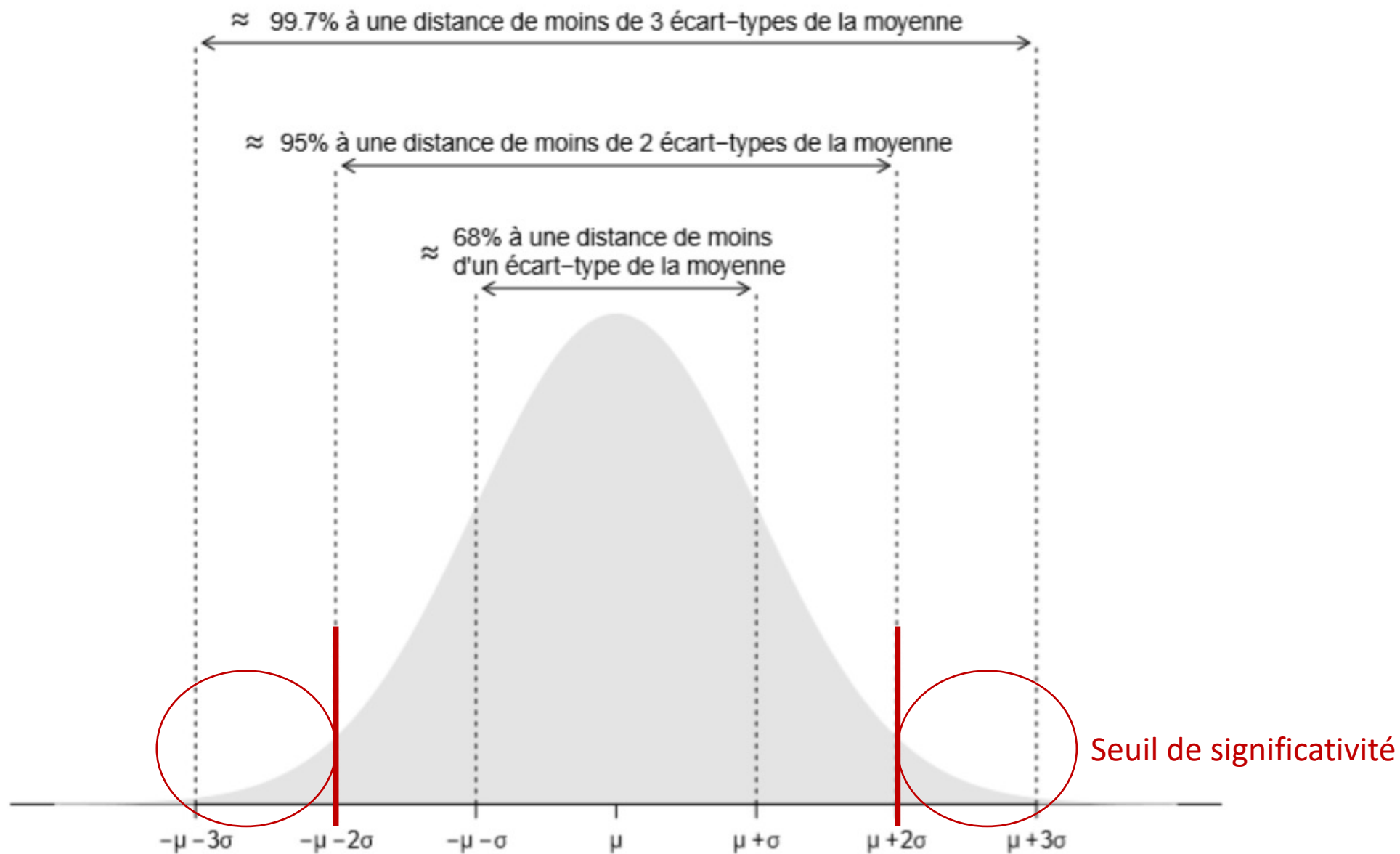
$X \sim N(\mu, \sigma^2)$  with  $\mu$  and  $\sigma^2$  the parameters of the distribution:

- $\mu$ : expectation of  $X$
- $\sigma$  = dispersion around the mean = Standard deviation (SD) of  $X$



- Symetric, unimodal
- Center around the mean

# Répartition des valeurs autour de la moyenne

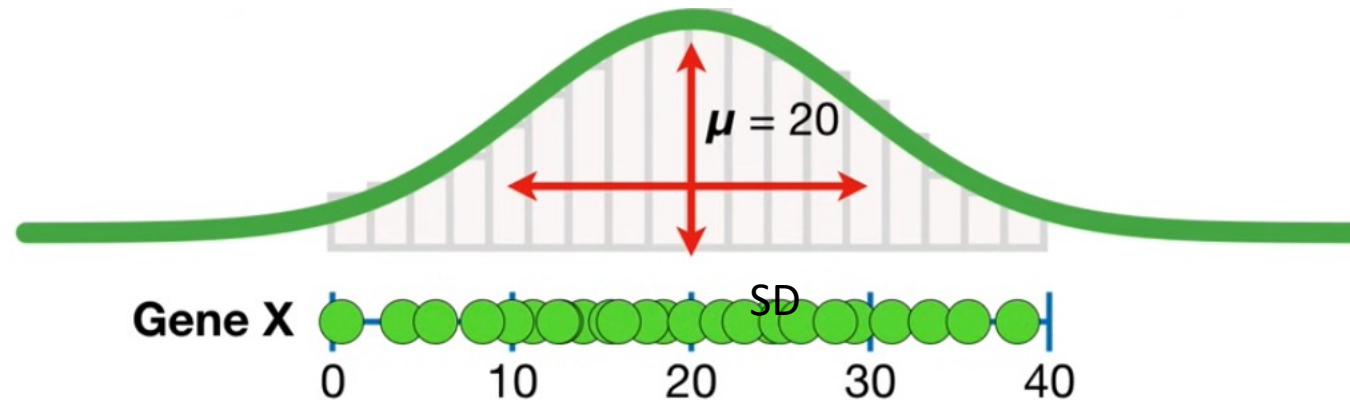


## Remember : Descriptive statistics (Univariate analysis)



Merely describe, show and summarize collected data

- **Central tendency** (mean, median...)
- **Dispersion** (variance, standard deviation)
- **Frequency distribution** (count, relative, cumulative)

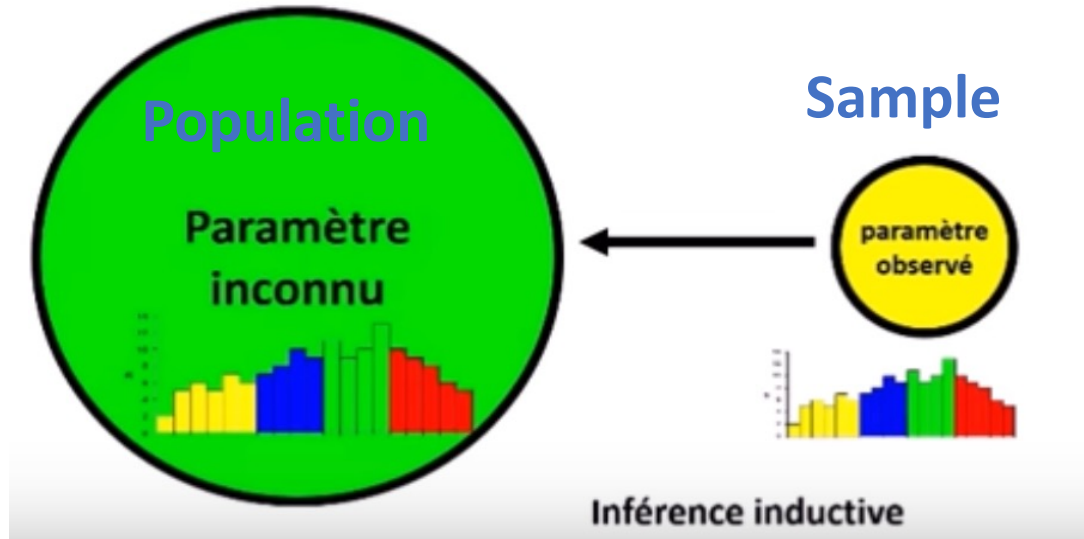
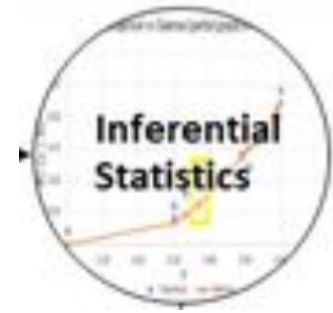


Identify the characteristics of data for each variable(s)

→ Allows you to formulate hypotheses and guide statistical analyzes

# Inferential Statistics

## Predictions - Generalizations



**Make inferences about the population**

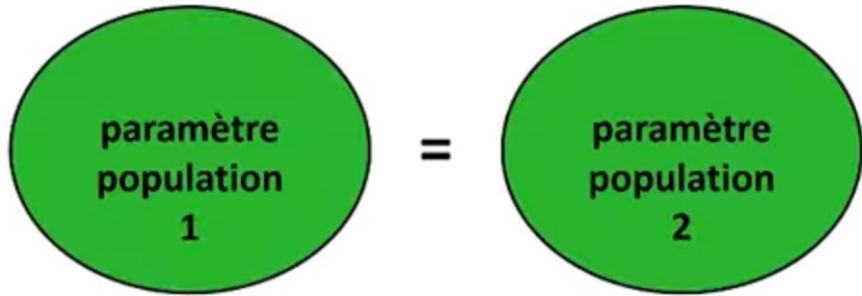
- How can I use my sample to make predictions about the population = **Estimation**
- How do I prove a theory about my data's behaviour (comparison) = **Hypothesis Testing**

# Hypothesis testing approach

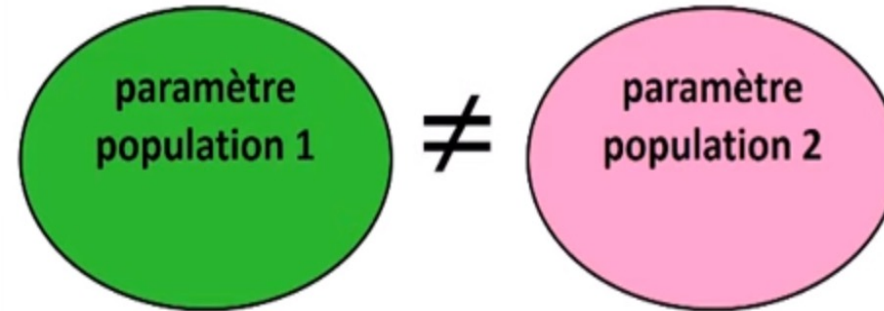
Trying to validate a hypothesis relating to a population parameter from a sample comparisons

Is there a **real** difference or just a coincidence (chance)

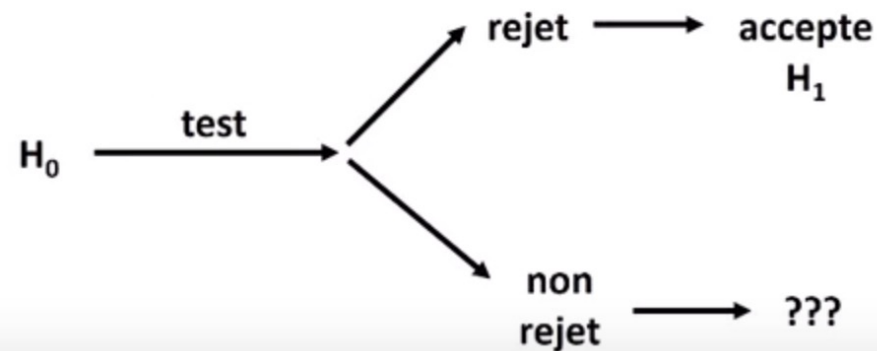
Null Hypothesis  $H_0$



Alternative hypothesis  $H_1$



**We are testing the null hypothesis!**

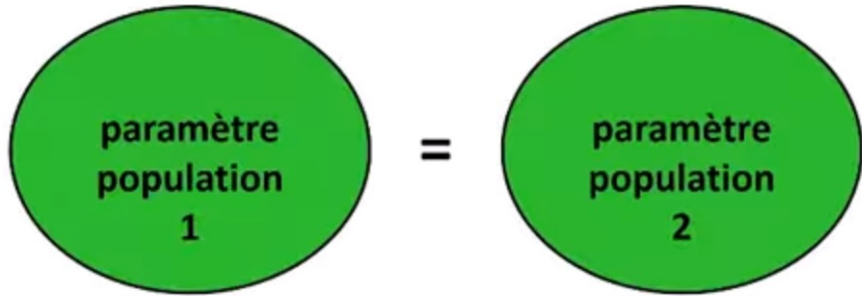


# Hypothesis testing approach

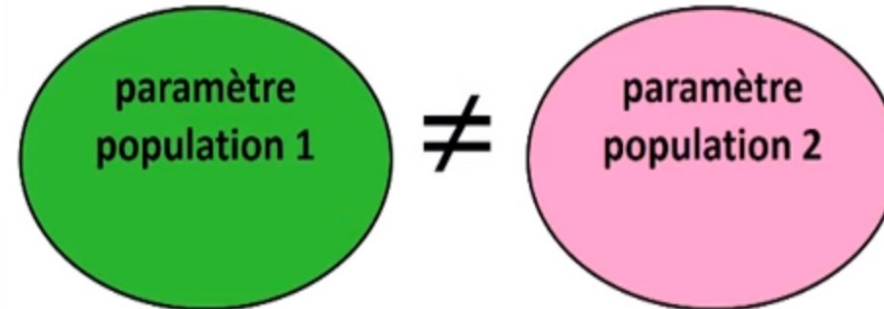
Trying to validate a hypothesis relating to a population parameter from a sample comparisons

Is there a **real** difference or just a coincidence (chance)

Null Hypothesis H0



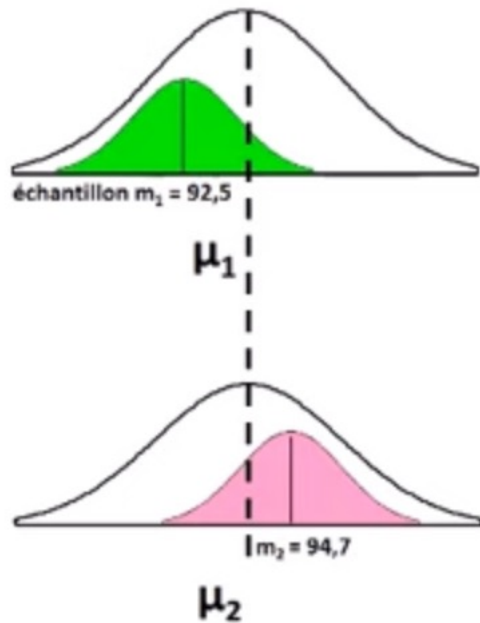
Alternative hypothesis H1



**“Absence of Evidence is not Evidence of Absence”**

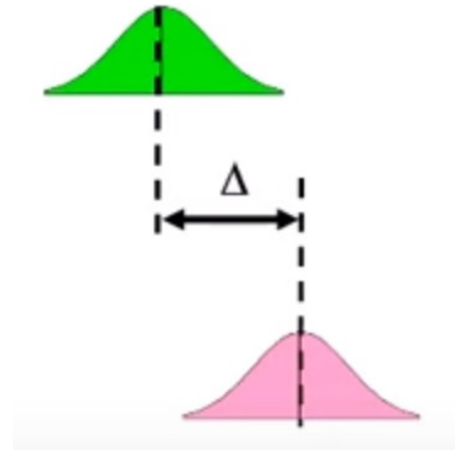
# Hypothesis testing & mean comparison

If  $H_0$  true... no difference



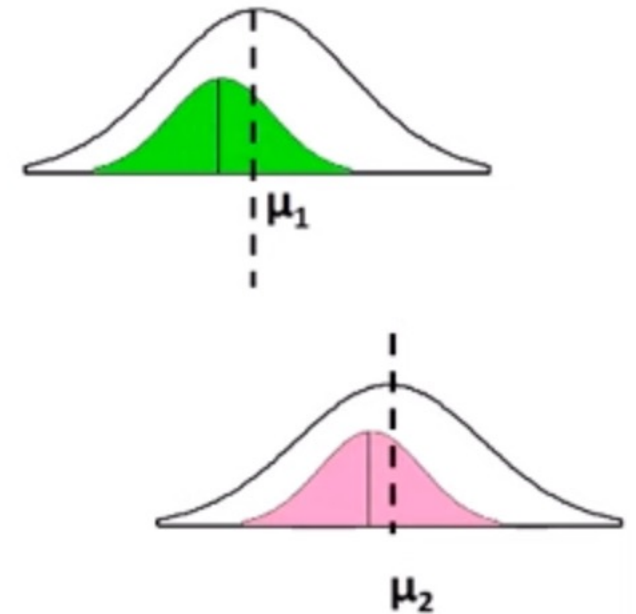
$$H_0 : \mu_1 = \mu_2$$

**SAME distribution**  
→ **Sampling fluctuation**



$$\text{Si } H_0 \text{ vraie : } \Delta = m_1 - m_2 \approx 0$$

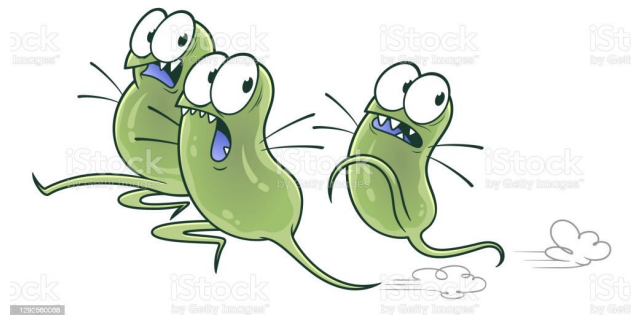
If  $H_0$  rejected,  $H_1$  accepted



$$H_1 : \mu_1 \neq \mu_2$$

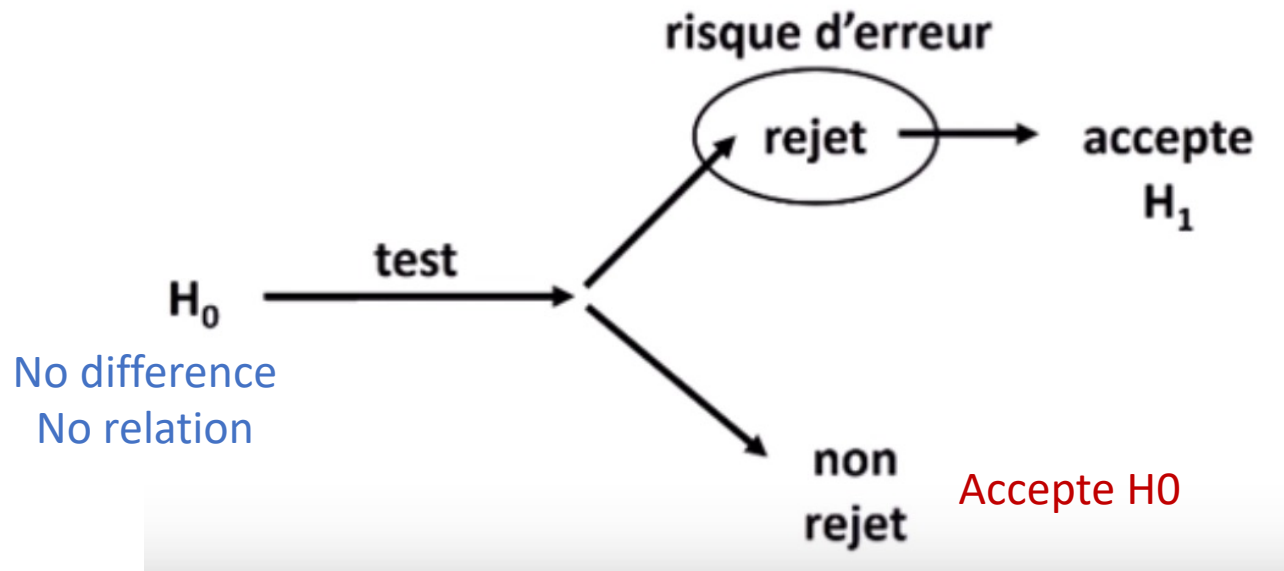
**Two different distributions**

**Inference Issue : Subjected to errors!!**  
**The risk is linked to the result of hypothesis testing**  
**Because of your sampling!**



# The risk of Type I error $\alpha$

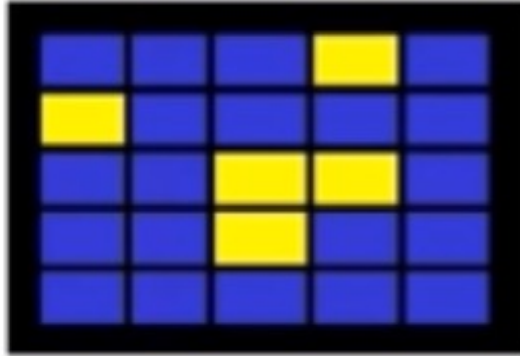
- A probability between 0 and 1, or 0 and 100%
- Is when a difference is affirmed but there is none (=False positive)!!



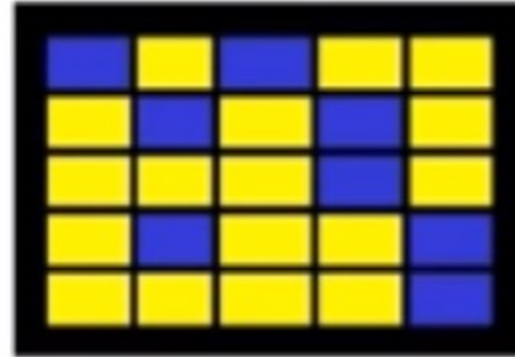
$\alpha$  = Risk to reject  $H_0$  if  $H_0$  is true

## Sampling

25 tiles  
→ 80% blue



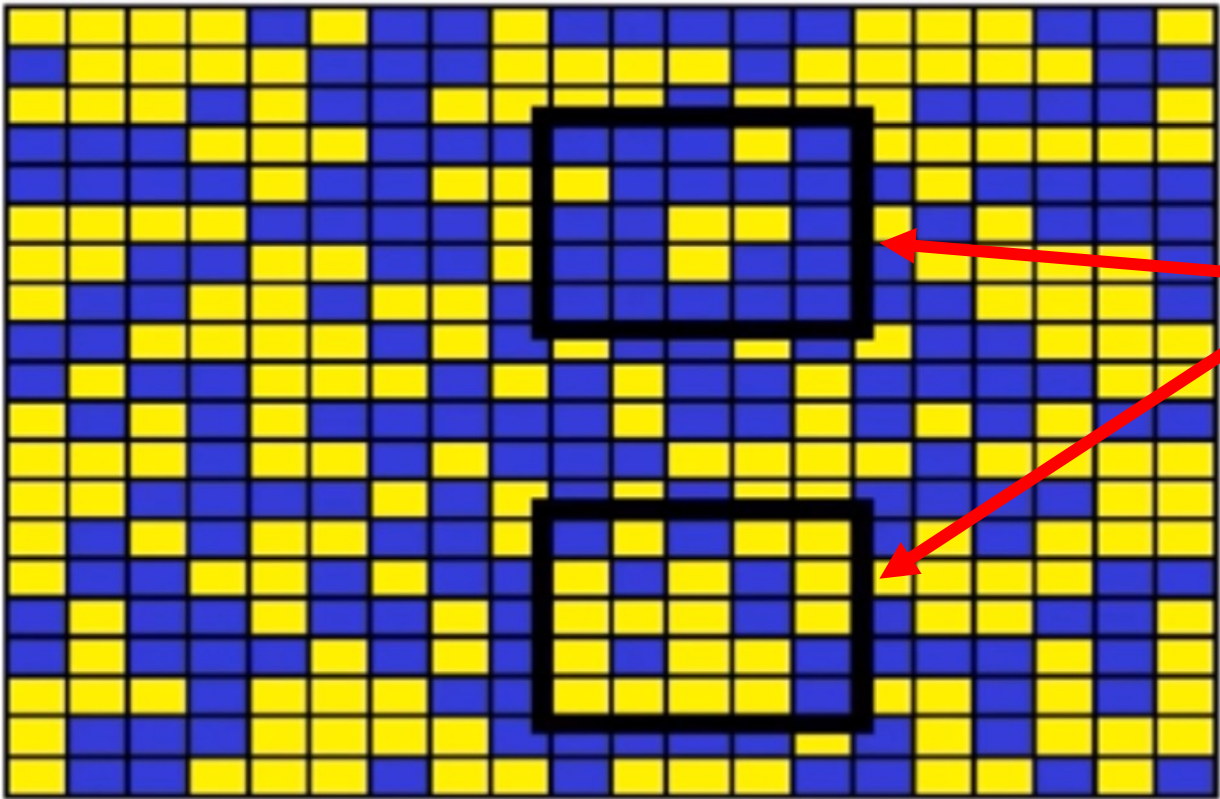
25 tiles  
→ 32% blue



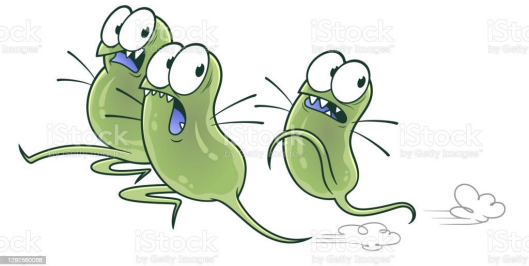
Do the two samples come from the same population? (same distribution)?

- **H<sub>0</sub> is rejected**
- but let's go to the store...see the population

Come from the same population (50% blue, 50 % yellow)!!

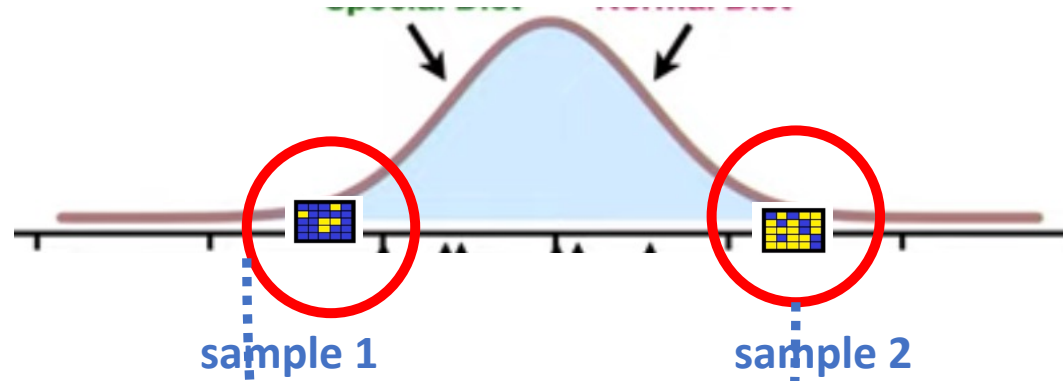


Rare sample type



Conclude on the basis of our samples that they came from two different distributions  
= Type I error

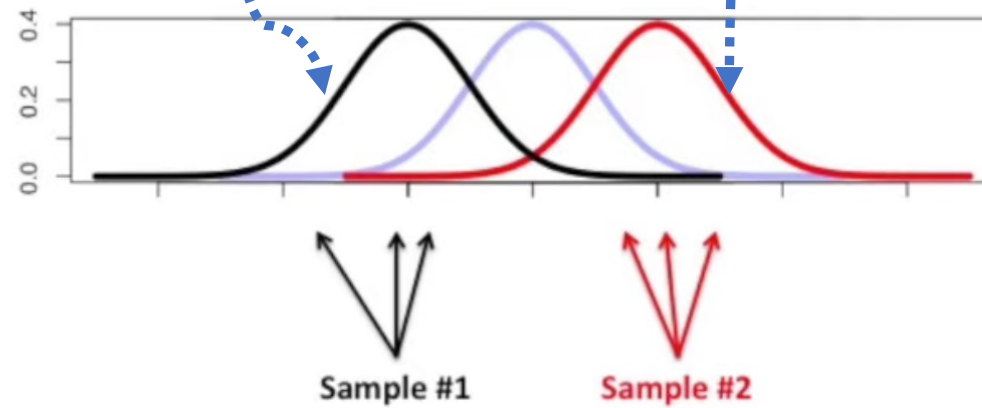
Data come from the same distribution but ...



sample 1

sample 2

The test see...

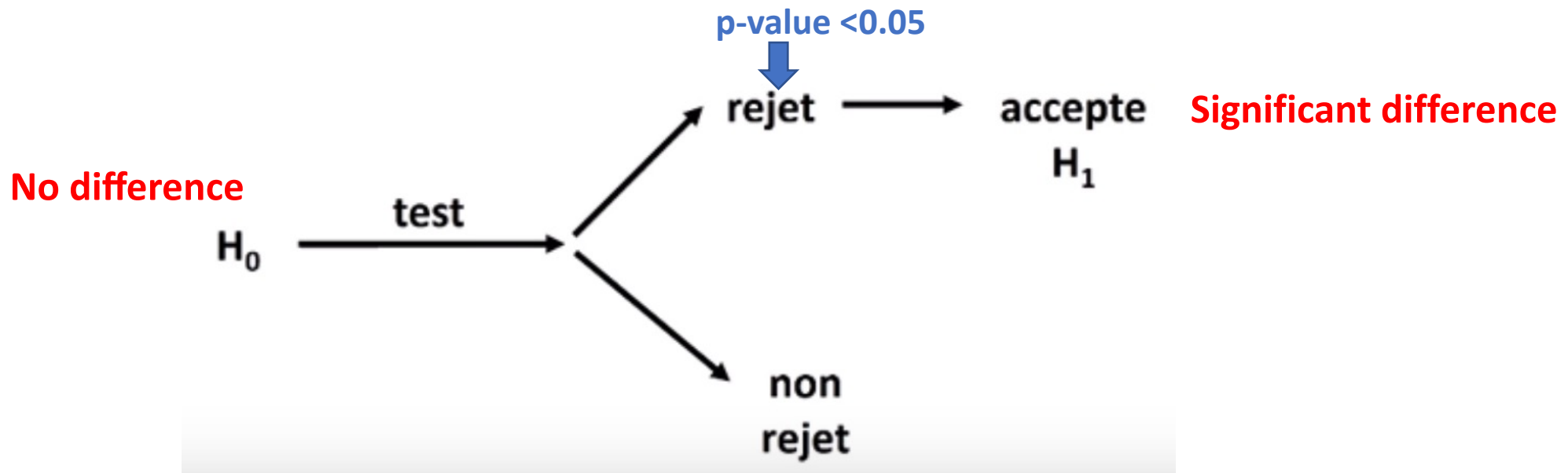


Sample #1

Sample #2

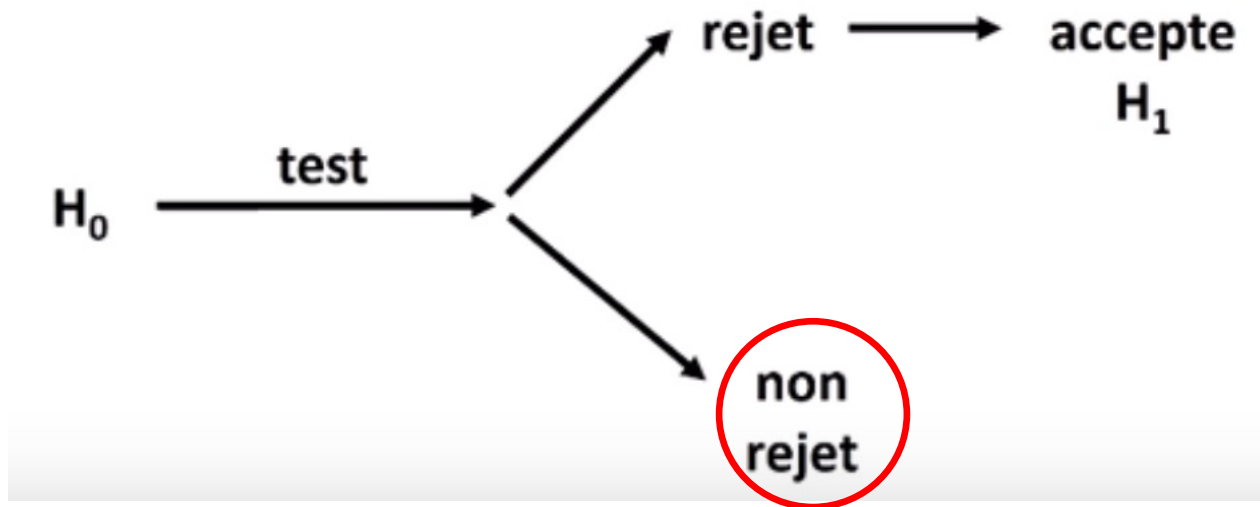
Two different distributions

- $\alpha$  is chosen before the test : **Significance threshold**
- $\alpha$  often set 5% (H0 wrongly rejected)
- In science the "almost no chance" translates to in less than 5% of cases where H0 is true = **p-value < 0.05**



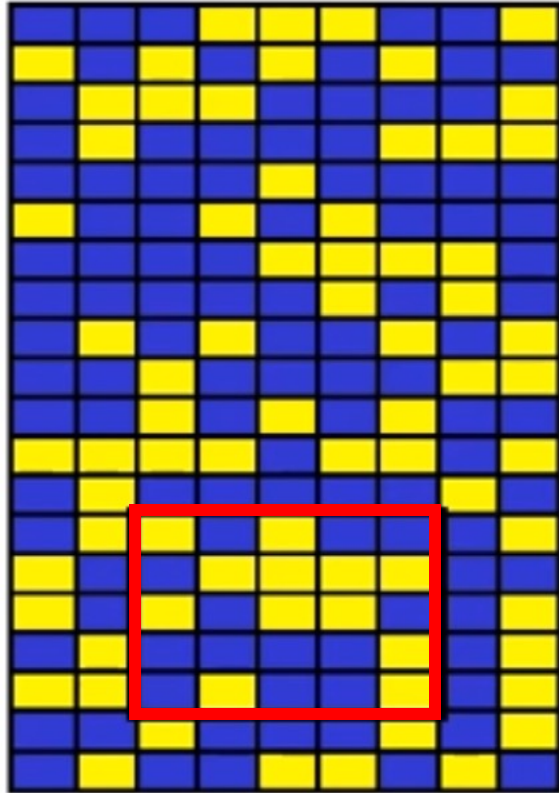
## Risk of Type II Error : $\beta$

Failing to conclude a difference when there is a true one ("False Negative")  
Probability of not rejecting  $H_0$ , if  $H_1$  is true



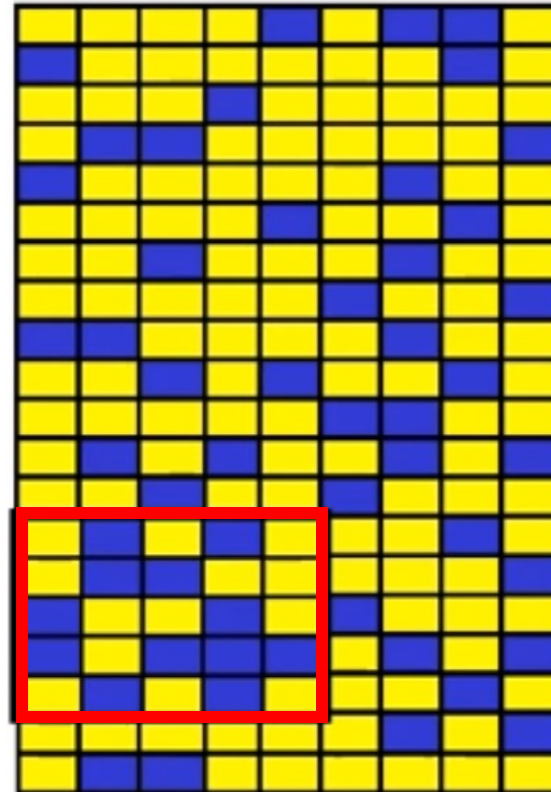
$\beta$  is not calculable

60% of blue



48% blue

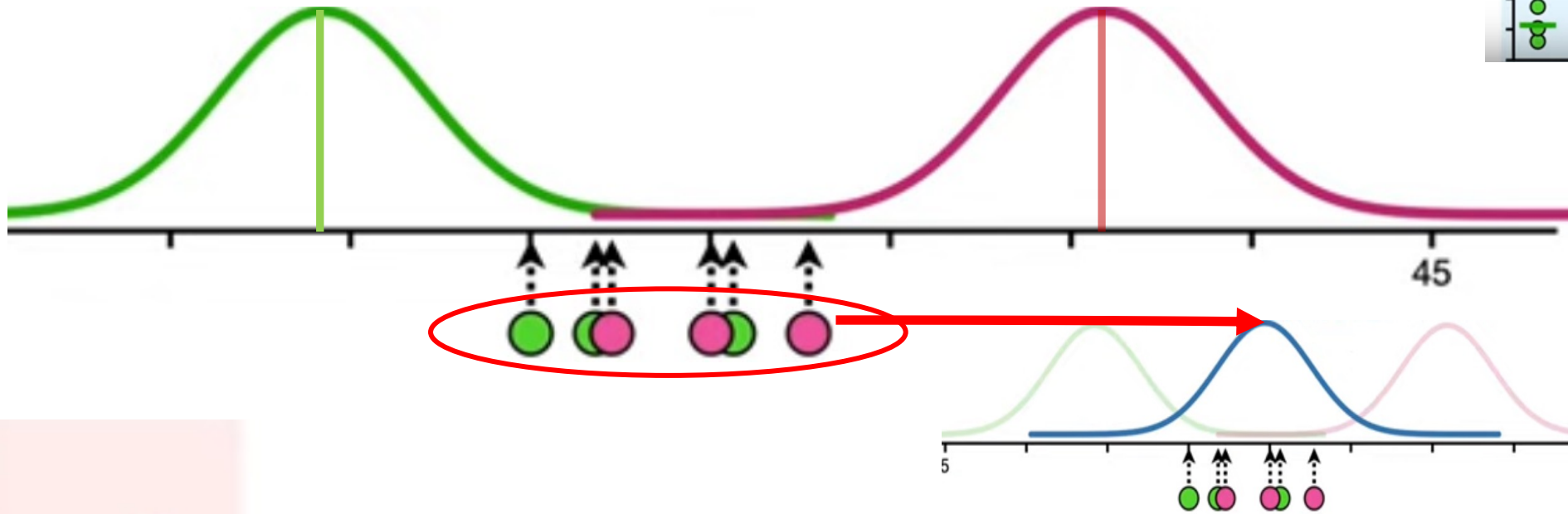
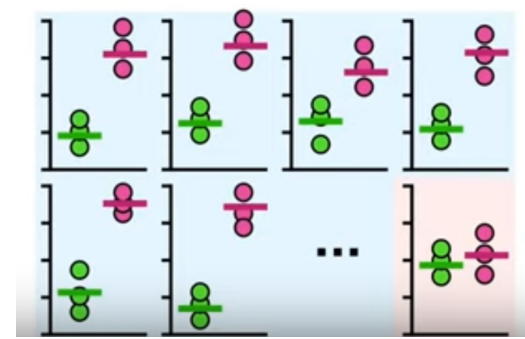
30% of blue



52% blue

- 2 different tiles = 2 different populations, H0 should be rejected But that would not have been the case during the test with our sampling...

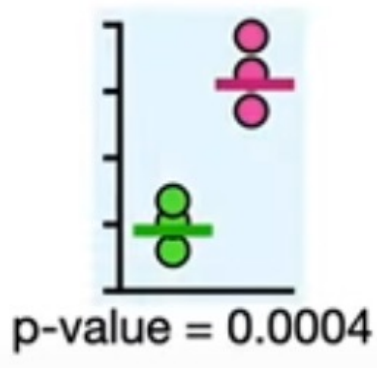
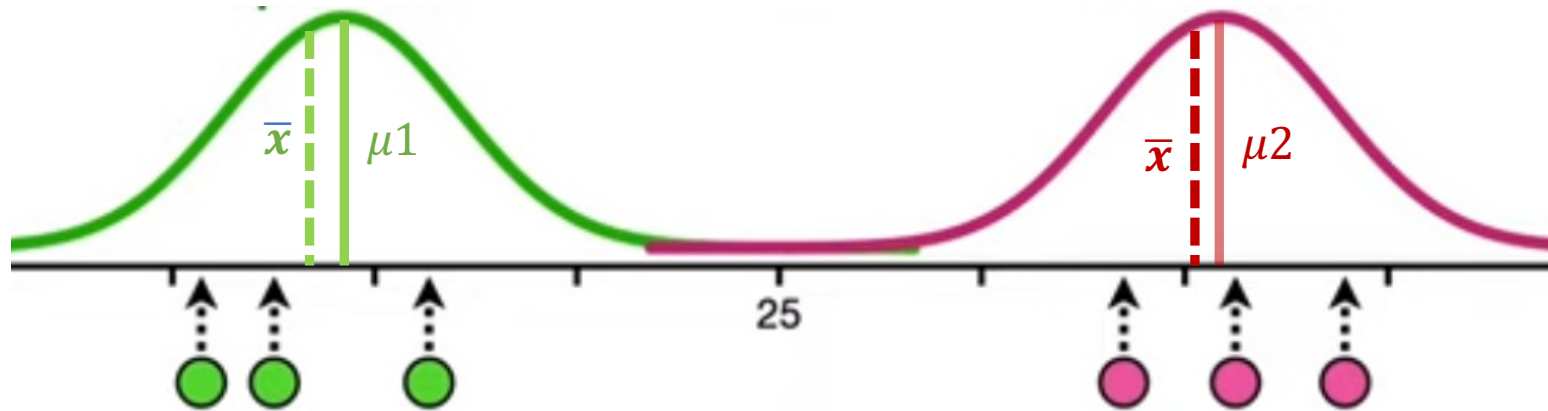
**But sometimes...**



**p=0.23!!!**

**Even if two different distributions (pop)...the test (your data) thinks they come from the SAME distribution!  
Unable to correctly reject H0...**

# Scientifically ... representative sampling of population



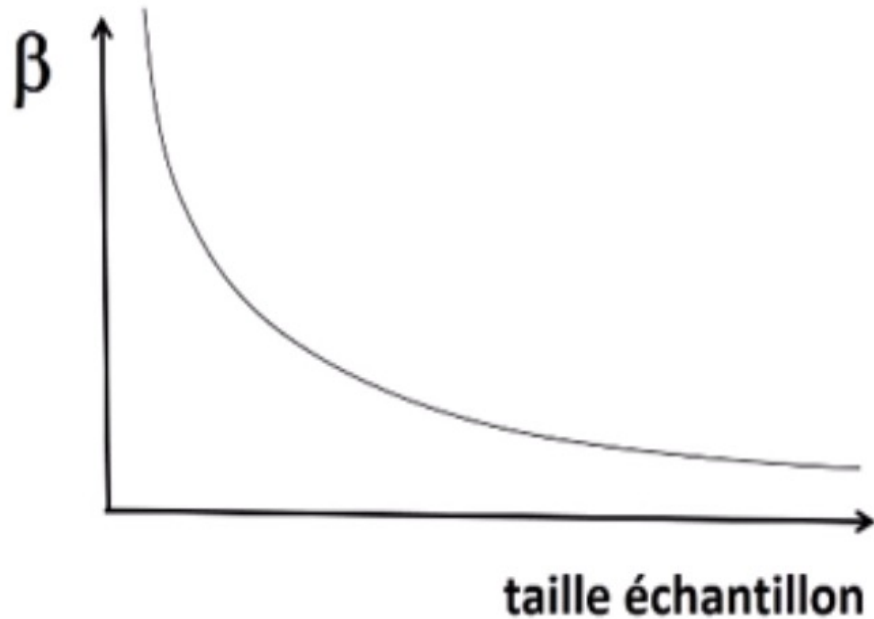
→  $H_0$  correctly rejected

→ = Data do not belong to same distribution

→ **Two different populations**

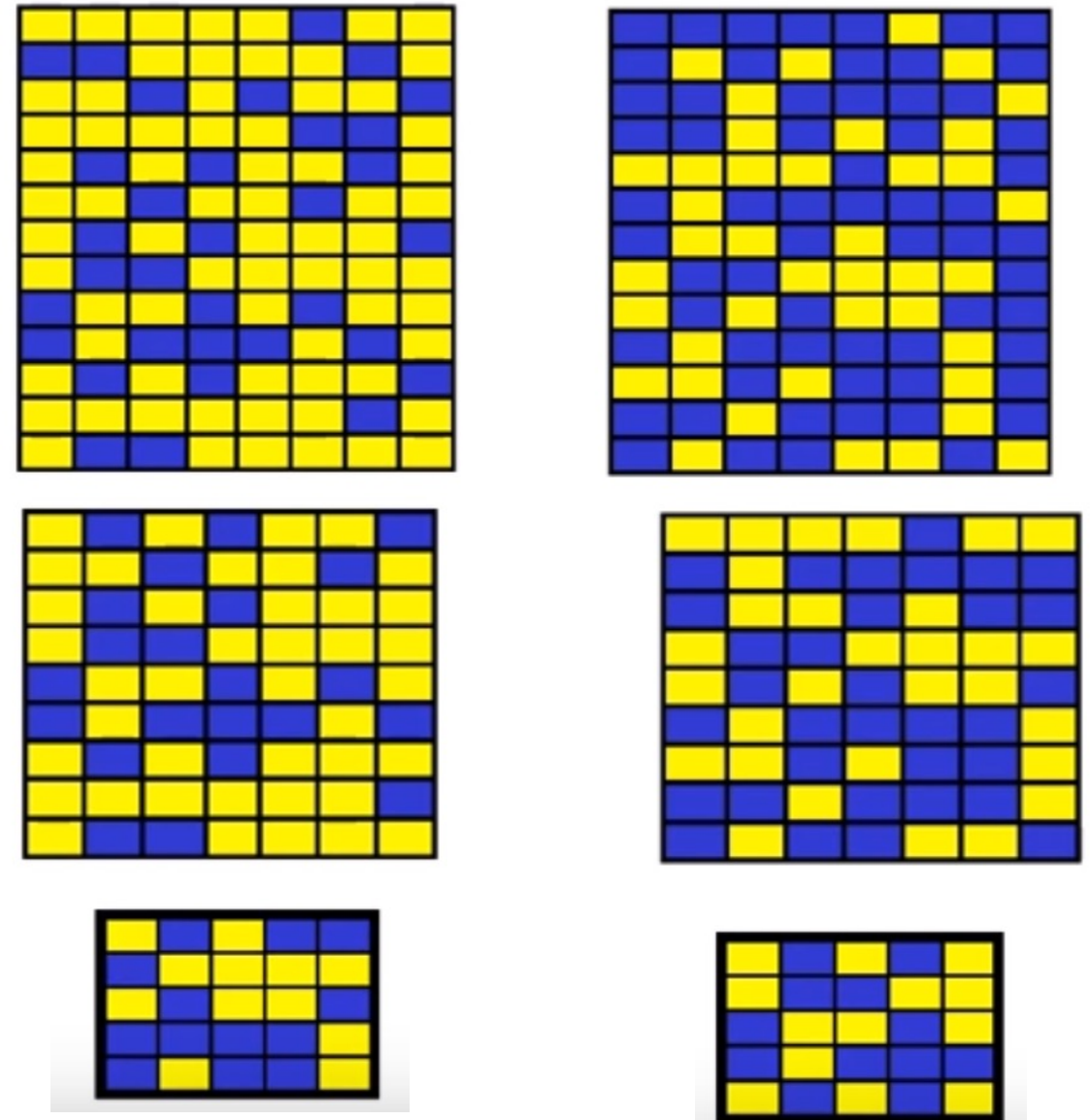
# Fundamental relationship

$$\text{Power} = 1 - \beta$$



**Power:** Probability of correctly reject the  $H_0$  hypothesis  
Ability of a test to detect differences

The more the size increases, the more the differences appear! The power of the test increases!

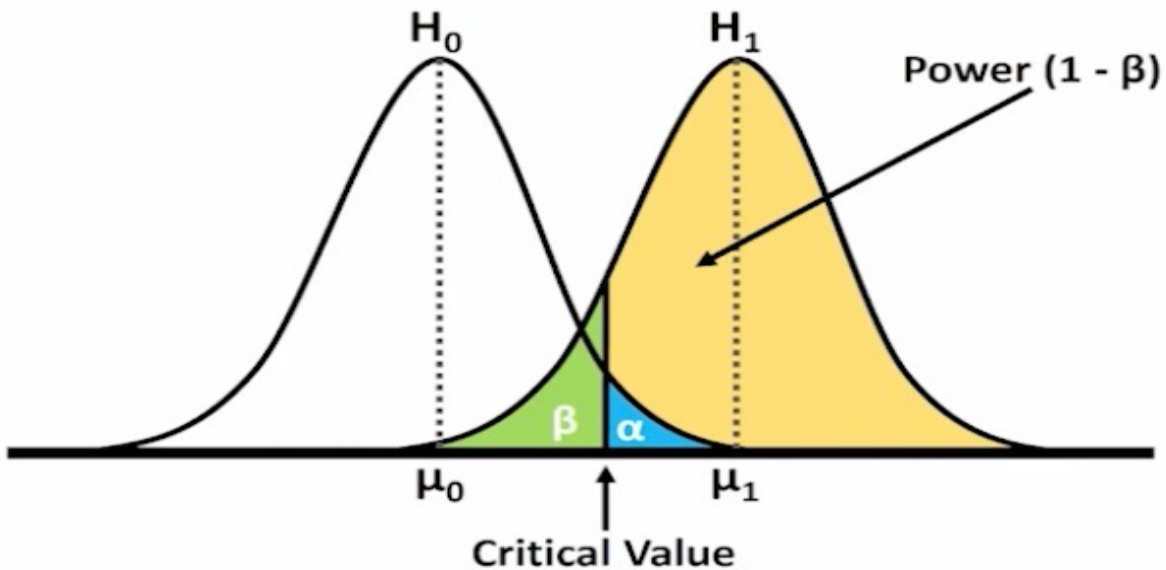


# Summary

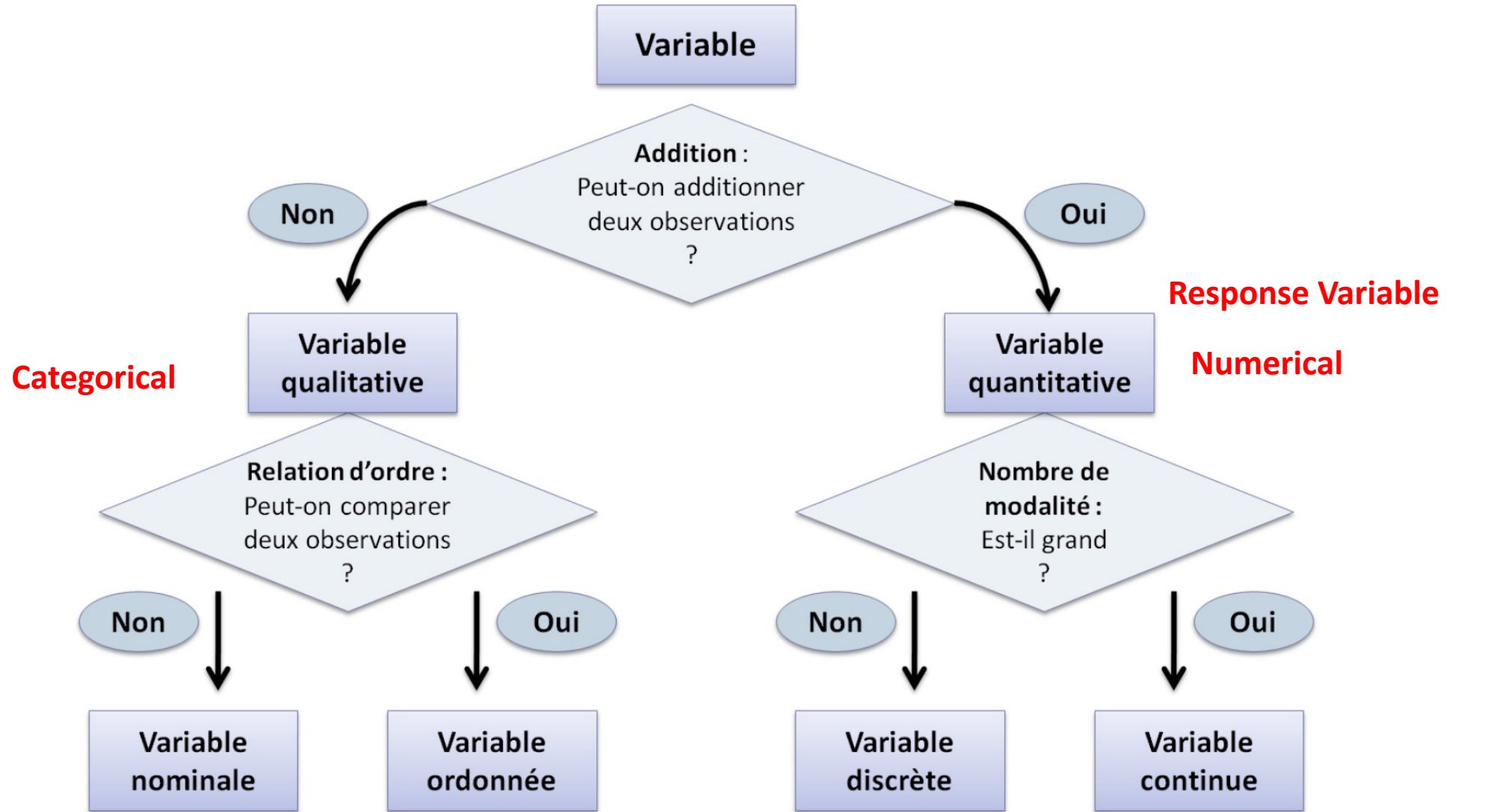
Population

TEST échantillons

	$H_0$ vraie	$H_1$ vraie
accepter $H_0$	OK	erreur type 2 $\beta$ Faux Négatif
rejeter $H_0$	erreur type 1 $\alpha$ Faux positif	OK



# Reminder on variables... important for statistical tests



• Married, single...  
→ No relation order

• Behaviour  
• good, excellent...

Child in family (1,2,3..)  
finite number of real values

Size, weight : infinite

# Bivariate Hypothesis Testing

- Seek to **quantify the association** between a **variable to be explained** (response/Quantitative) and an **explanatory variable** (factor/categorical)
- **Make statistical inferences about the relationship between two variables, One quantitative variable (response) & one qualitative (explicative)!**
  - Can variations in **species richness** (response variable) be explained by the explanatory variable (factor) **Treatment**
    - **Comparison of mean between groups**

- **Parametric or non parametric test??**
- **which test?? significance ? (p-value)**
- **How many groups??**
- **Post hoc test required ??**



Which test for independent samples?

ONE categorical variable (H/F) & ONE continuous variable (numerical)

**Normalité des données?**

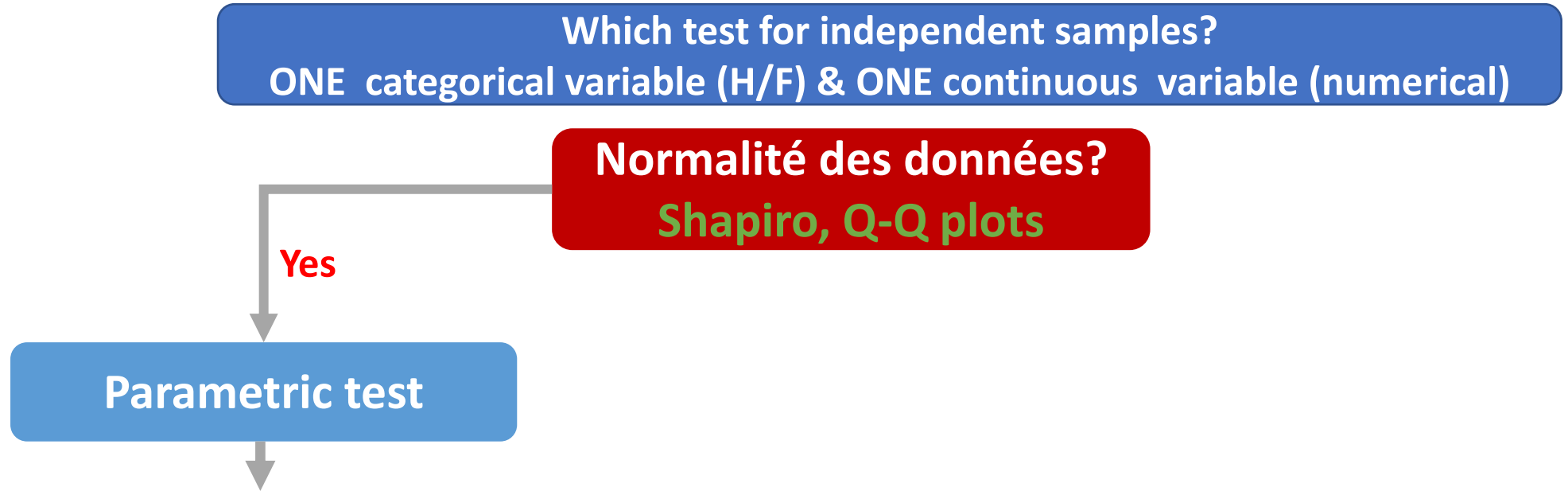
Shapiro, Q-Q plots

Which test for independent samples?  
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Normalité des données?  
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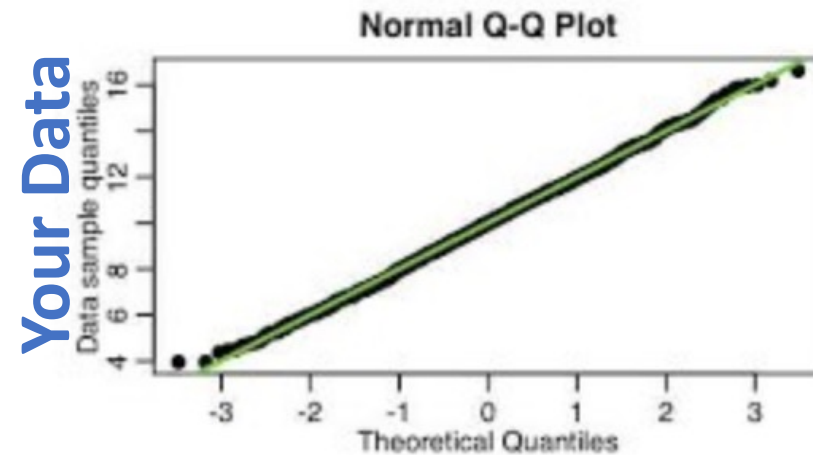
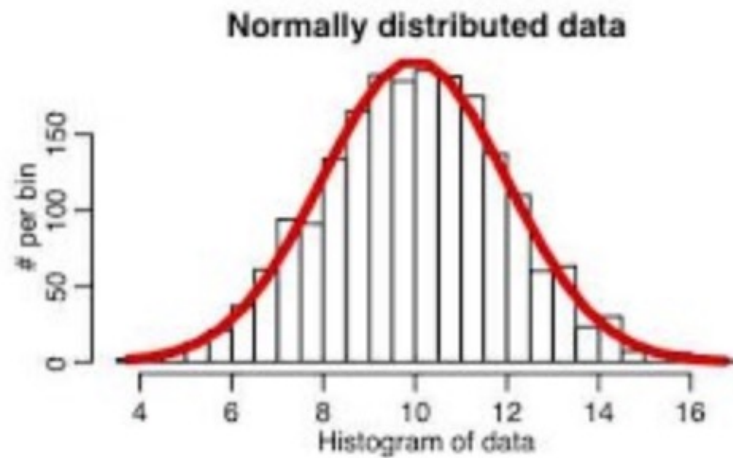
Yes

Parametric test



# Q-Q plot normale: Compare your distribution with a normal distribution

Do my data follow a normal distribution ?

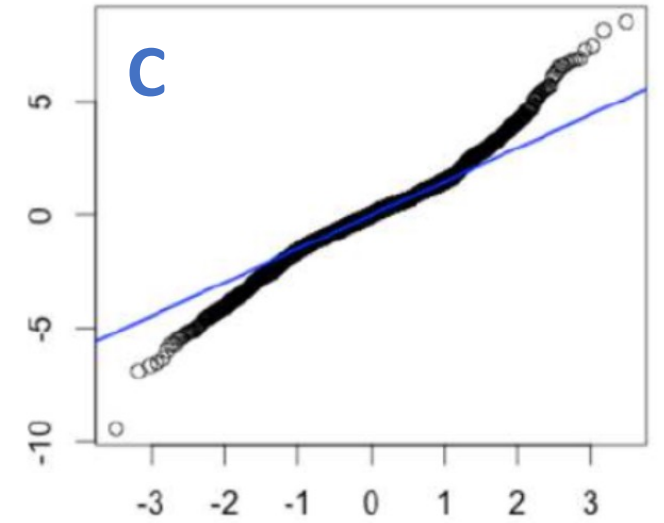
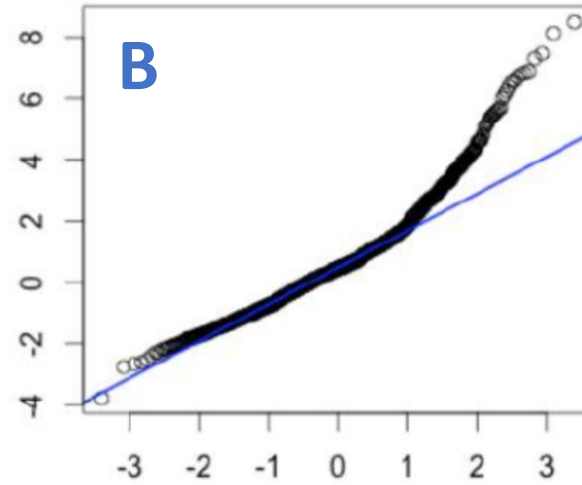
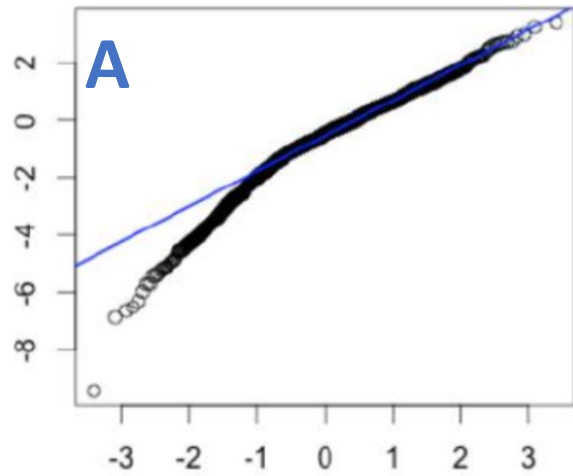


Conclusion?

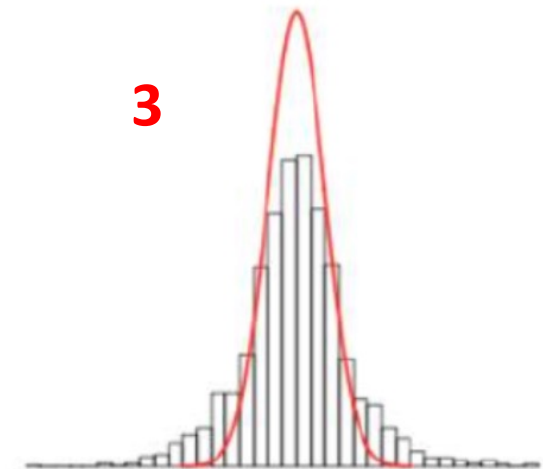
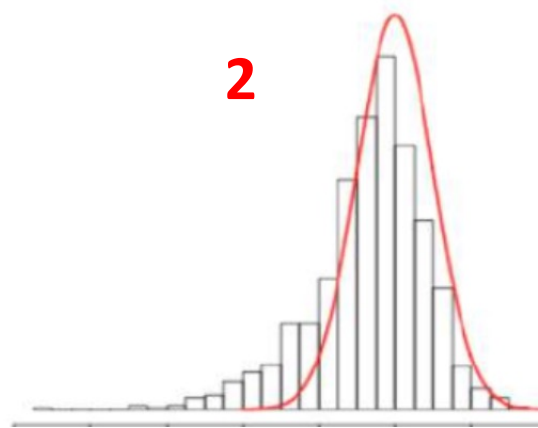
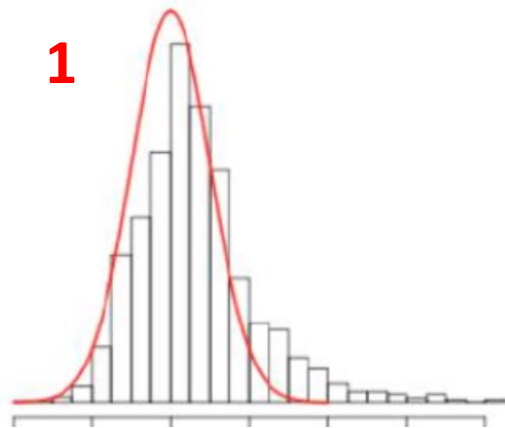
Normal Data ( $\mu=0$ ,  $SD=1$ )

The line drawn by QQ-Plot indicates the position that the points must have to follow a normal distribution

# What are the distributions (bottom) corresponding to these QQ-plots?



??????????



Which test for independent samples?  
ONE categorical variable (H/F) & ONE continuous variable (numerical)

Normalité des données?  
Shapiro, Q-Q plots



Yes

Parametric test



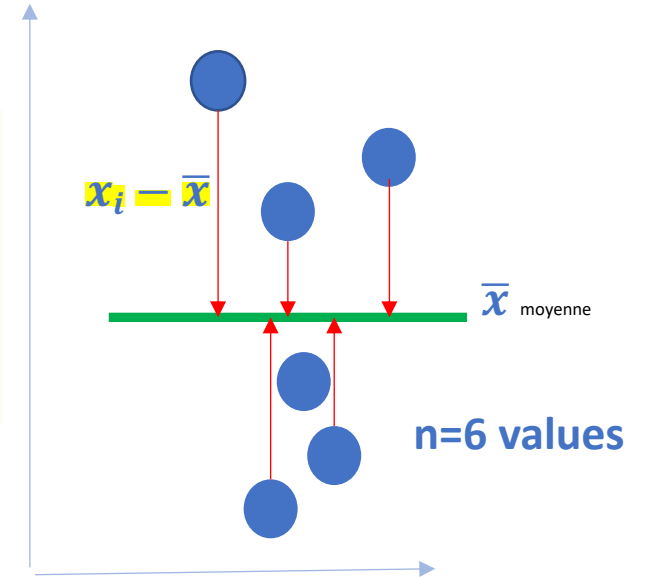
Variance Homogeneity  
Bartlett, levene, F-test

# Variance = $S^2 / \sigma^2$

- Variance measures the degree of dispersion of a data set around the mean
- Arithmetic mean of squared deviations from the mean! ☹️

→ square unit

$$S^2 = \sum_{i=1}^n \frac{(x_i - \bar{x})^2}{n - 1}$$



# Standard Deviation = $S / \sigma$

$S = \sqrt{S^2}$  The advantage of the standard deviation : expressed in the same unit as the data series

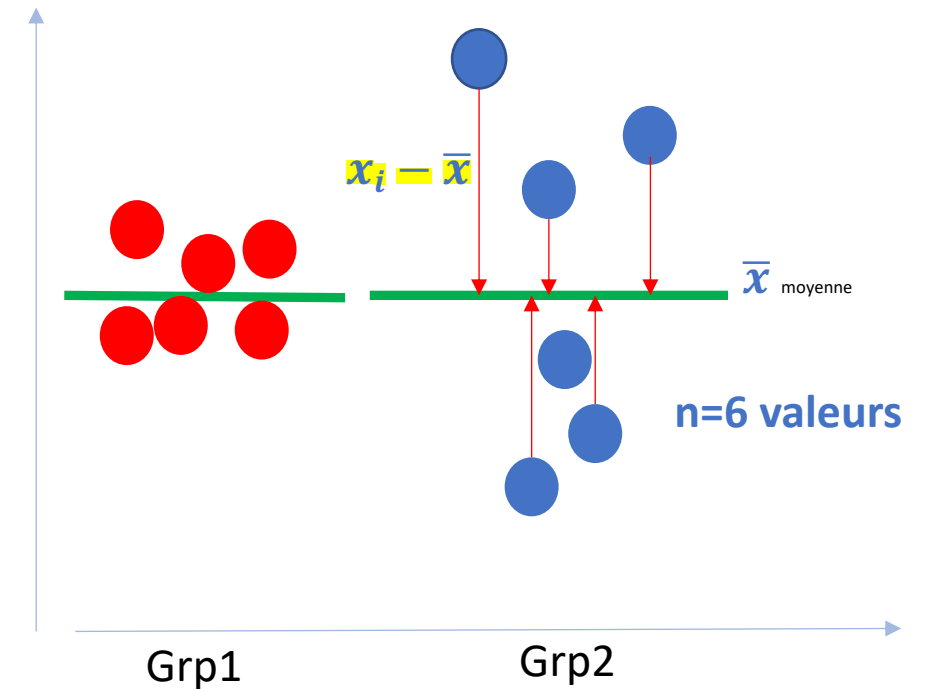
$$S^2 = \sum_{i=1}^n \frac{(x_i - \bar{x})^2}{n-1} = \frac{\text{Sum of Squares (SS)}}{n-1}$$

SS will be greater in the sample...??

Results of test using variance :

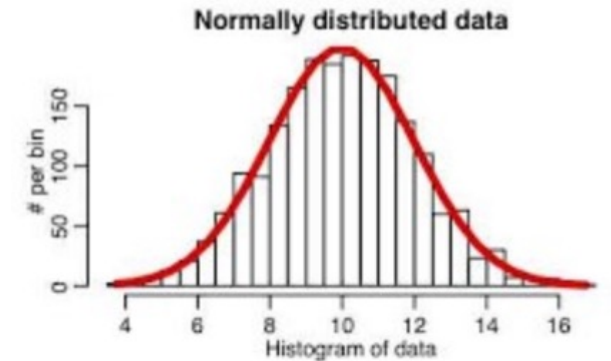
	Df	Sum Sq	Mean Sq	F value	Pr(>F)
groupe	3	13.03	4.343	0.211	0.887
Residuals	14	288.75	20.625		

- **Sum of Squares (= SS, Sum Sq) in your results!**  
→ Numerator of variance!!
- **Mean Square (= Mean Sq= VARIANCE formula!!!)**



# Requirement for parametric test... check-list!

- Check **normality** of data: Shapiro Test & QQ-plots!!
- Shapiro:  $H_0$  is «data follow normal distribution»



- Check **variance Homogeneity**: F-test (2 groups), Bartlett's & Levene's tests

$H_0$ : « No difference »

$$S^2 = 169$$

$$S^2 = 289$$



# Parametric Tests

Follow a known distribution (Normal distribution)



**Position** parameters  
**Dispersion** parameters

Conditions are required (variance homogeneity)

- **T-test (paired or unpaired):** Compare of the means from **2 sample groups** for one variable
- **One way Anova** (variance analysis) : compare the means of **three or more sample groups** for one variable

Which test for independent samples?  
ONE categorical variable (H/F) & ONE continuous variable (numerical)

Normalité des données?  
Shapiro, Q-Q plots

Yes

Parametric test

Variance Homogeneity  
Bartlett, levene, F-test

No

Transformation  
(square root, log)

Yes

Yes

How many groups?

2 Groups

3 Groups  
& more

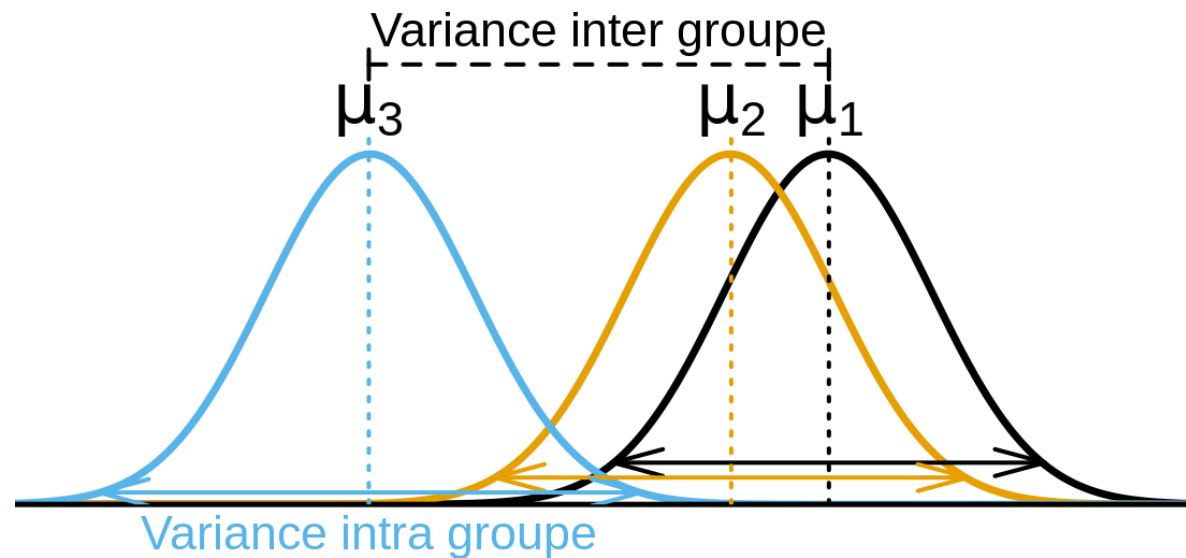
T-test

One way ANOVA

Post hoc test

# ANOVA: ANalysis Of VAriance (One way Anova= Univariate) (3 groups at least)

- Compare the **variance of the group means** to that **within groups** (i.e. intra-group variance) for a **single explanatory variable** (qualitative)



# ANOVA: ANalysis Of VAriance (One way Anova= Univariate)

- Postulate = The **VARIATIONS** observed between the **MEANS** of the different groups (AT LEAST 3) are so small that they are easily explained by chance!!!
- Evaluation : Compare the **variance of the group means** to that **within groups** (i.e. intra-group variance)
- ANOVA → variations through the Variance quantity

$$\boxed{\text{Variance inter-groupes}} + \boxed{\text{Variance intra-groupes}}$$

attribuable au facteur

attribuable à l'expérimentale  
(fluctuation de l'échantillonnage, hasard)

• **Statistic F** = 
$$\frac{\text{Factor effect!} \quad \textit{Inter-group Variance}}{\textit{Intra-group Variance} \quad \text{Chance /fluctuation}}$$

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
groupe	3	13.03	4.343	0.211	0.887
Residuals	14	288.75	20.625		

**Idea :**

if **the factor really has an effect**, the part of the variations that can be attributed to it = **Inter-group variance** will be significantly higher than the part of the variations that cannot be attributed to it = **Intra-group variance!**

**Statistic F** Follows a so-called **Fisher-Snedecor** law:

= **Distribution F** used for test of variances, distribution of variances not being normal

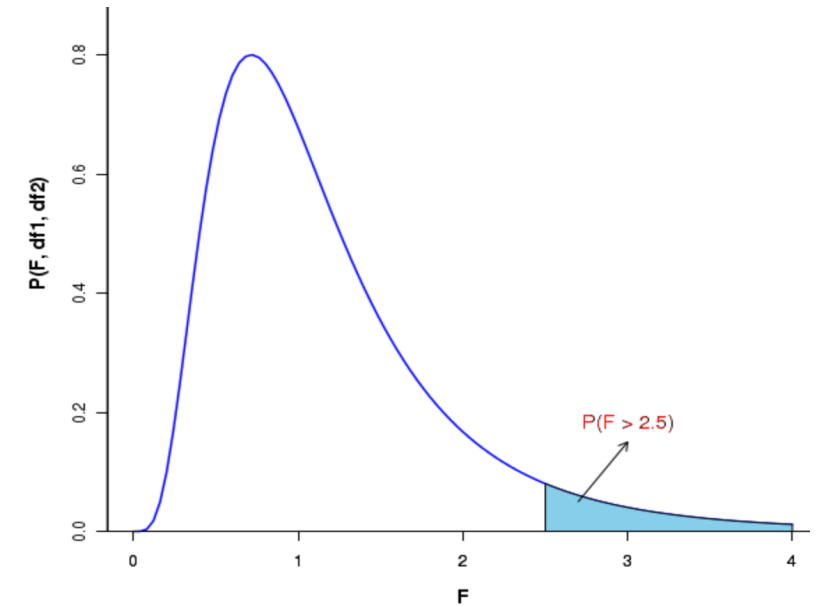
- Relation of an observed value of F with the a priori probability of encountering such a value ( $>$  or  $=$ ) by chance!
- $\rightarrow$  probability given by the law = p-value!

Denominator  $S^2$

Numerator  $S^2$

$S^2$

	Df	Sum Sq	Mean Sq	Sq	F value	Pr(>F)
groupe	3	13.03	4.343		0.211	0.887
Residuals	14	288.75	20.625			



variances	ddl	F	
entre k groupes	$v_k$	$k-1$	$v_k / v_r$
résiduelle	$v_r$	$N - k$	

Degré de liberté

- **Two-ways ANOVA : Influences of TWO qualitative variables on ONE quantitative variable**

**Exple: Influence of soil type and degree of humidity (ordinal variable) on plant yield**

# Non-parametric tests

**No assumptions are made for the distribution of data:  
Distribution-free tests, they are alternative to parametric tests**

- **Wilcoxon Rank test** : samples are paired/unpaired, 2 sample groups
- **Mann-Whitney test**: Independent samples, 2 sample groups
- **Kruskal wallis test** : Independant samples, Three or more groups

→ Based on the average ranks: we classify the values, we replace by a position (1,2 etc),  
Compares the average of the ranks between the groups

Which test for independent samples?  
ONE categorical variable (H/F) & ONE continuous variable (numerical)

Normality of data?  
Shapiro, Q-Q plots

NO

Non parametric test

How many groups?

2 Groups

At least 3 Groups

→ Unpaired Wilcoxon test  
→ Mann-Whitney

Kruskal Wallis

↓  
Post-hoc Test (Dunn)

**Which test for independent samples?**  
ONE categorical variable (H/F) & ONE continuous variable (numerical)

**Normality of data?**  
Shapiro, Q-Q plots

Yes

Parametric test

**Homogeneity of Variance?**  
Bartlett, levene, F-test

NO

Transformation  
(square root, log)

YES

YES

**How many groups?**

2 Groups

At least 3 Groups

T-test

One way ANOVA

Post-hoc Test (Tukey)

NO

Non parametric test

**How many groups?**

2 Groups

At least 3 Groups

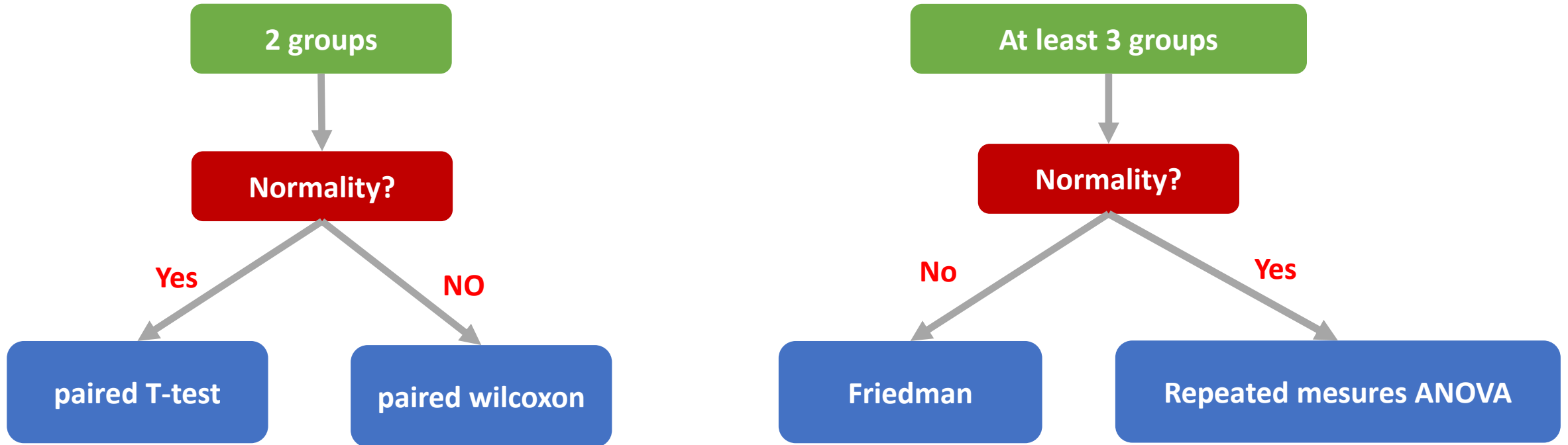
→ Unpaired Wilcoxon test  
→ Mann-Whitney

Kruskal Wallis

Post-hoc Test (Dunn)

# Repeated measurements – paired samples

Exple= time series, Before-After  
Treatment...



# Post-hoc Test

Statistical tests with **at least 3 groups!**

After ANOVA, Kruskal-wallis

→ The result of an ANOVA test is **an Overall p-value**

Exple: You are comparing the effect of 3 soil types (A,B,C) on plant growth

**ANOVA returns a p-value of 0.03**

It does not tell you which pair of groups are significantly different!!!!

→ Post-hoc Test! Multiple comparisons (eg: Gp A vs. Grp. B; GrpB vs. Grp C; Grp C vs. Grp A!)

- Parametric Post-hoc test (ANOVA) → **Tukey Test**
- Non-parametric Post-hoc test (Kruskal wallis) → **Dunn Test**

## Multiple Testing Issue: increasing the risk...

Test is based on **probabilities**, so there is always **a risk** of drawing the **wrong conclusion!**

→ **No hypothesis test is 100% reliable**



Performing hypothesis testing:

- You have two hypotheses :
  - H0: Null hypothesis = the reference hypothesis : No difference
  - H1: Alternative hypothesis: There is a difference

- You encounter: **Type I error :  $\alpha$  = Risk alpha**

$\alpha = 0.05$  Is the **probability** (significance threshold) to incorrectly **reject H0!**  
In other words, an acceptable chance of a false positive!!

# Differential abundance : Multiple testing!!

**ONE TEST :**

$$P_{\text{False Positive}} = P_{\text{error}} = \underline{\alpha} = 0.05$$

Complementary Prob

$$P_{\text{no\_error}} = 1 - \underline{\alpha} = 0.95$$

**TWO TEST without making error :**  $P_{\text{no\_error in two tests}} = (1 - \underline{\alpha}) * (1 - \underline{\alpha}) = (1 - \underline{\alpha})^2$

Complementary Prob

$$P_{\text{at\_least\_ONE\_error in two tests}} = 1 - (1 - \underline{\alpha})^2$$

**Generalization to n TESTS**

$$P_{\text{at\_least\_ONE\_error in n tests}} = 1 - (1 - \underline{\alpha})^n$$

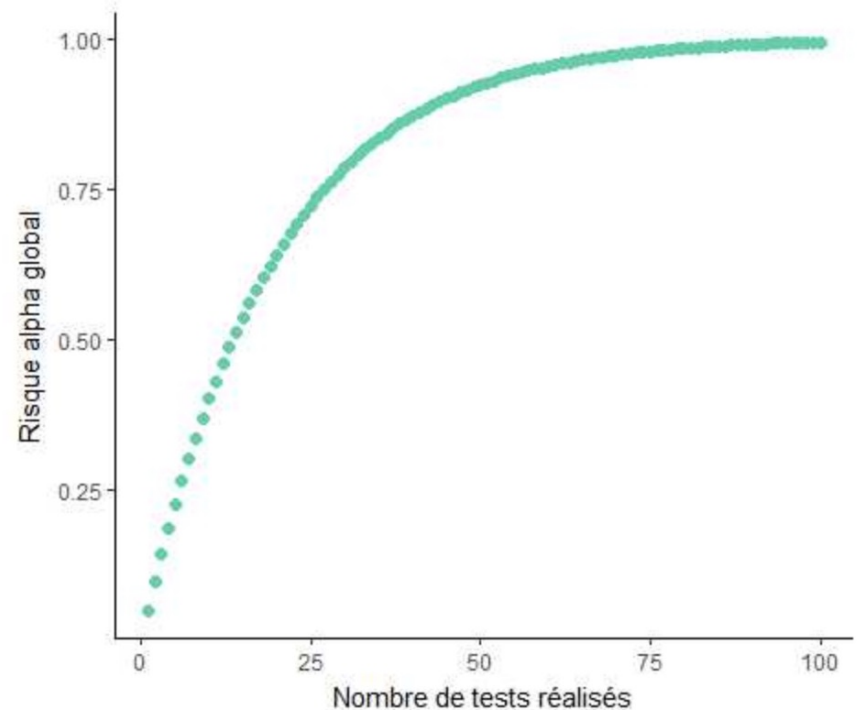
It's called the global  $\underline{\alpha}$  risk

# What does it means...

- You test **ONE** ASVs (n=1) for differential abundance:  $1-(1-\alpha)^n = 1-(1-0.05)^1 = 0.05$
- You test **3** ASVs (n=3):  $1-(1-0.05)^3 = 0.14$
- You test **100** ASVs (n=100):  $1-(1-0.05)^{100} = 0.9941$

The global risk  $\alpha$  reach **0.9941=99.41%!!!!**

**→ 99% to wrongly reject the H0 at least  
One times**



Need to ajusted this phenomen by using p-value **adjusted!**

# FDR : False Discovery Rate : Benjamini-Hochker

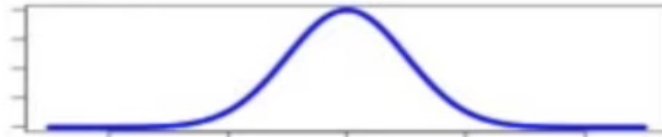
The idea : Discard bad data that looks good!!!

Benjamini-hochker **adjusts p-values**  
to limit the number of **false positives**  
that are reported as **significant** (pvalue < 0.05)

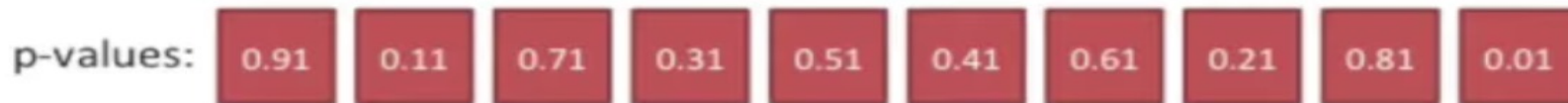
**Adjusts p-values**  
means that it makes them **larger!**

**Using FDR cutoff < 0.05**  
means less than 5% of the significant results will be false positives

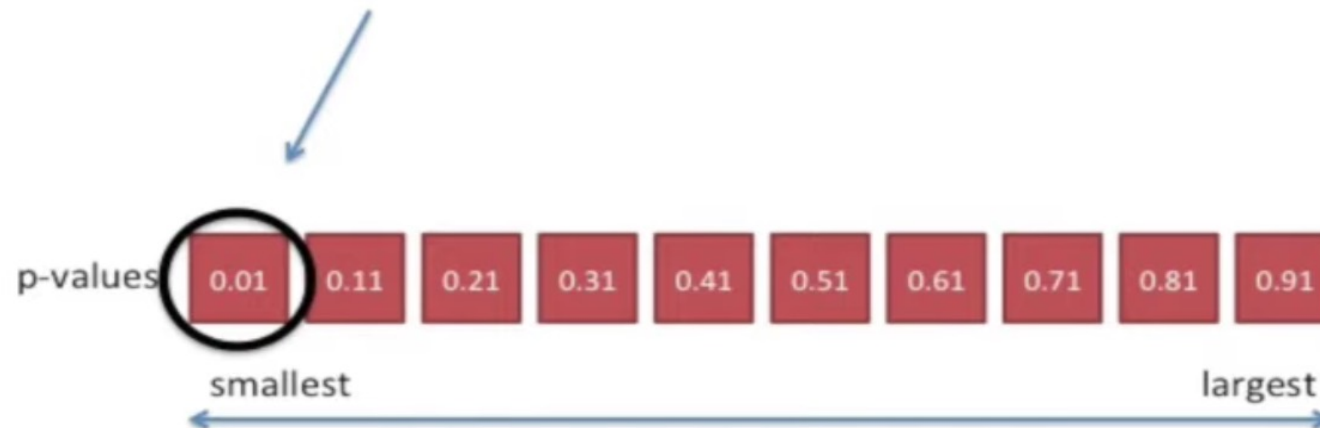
# Mathematical approach FDR-Benjamini-Hochker



10 pairs of samples taken from the same distribution. (i.e. 10 genes that were not effected by the drug).

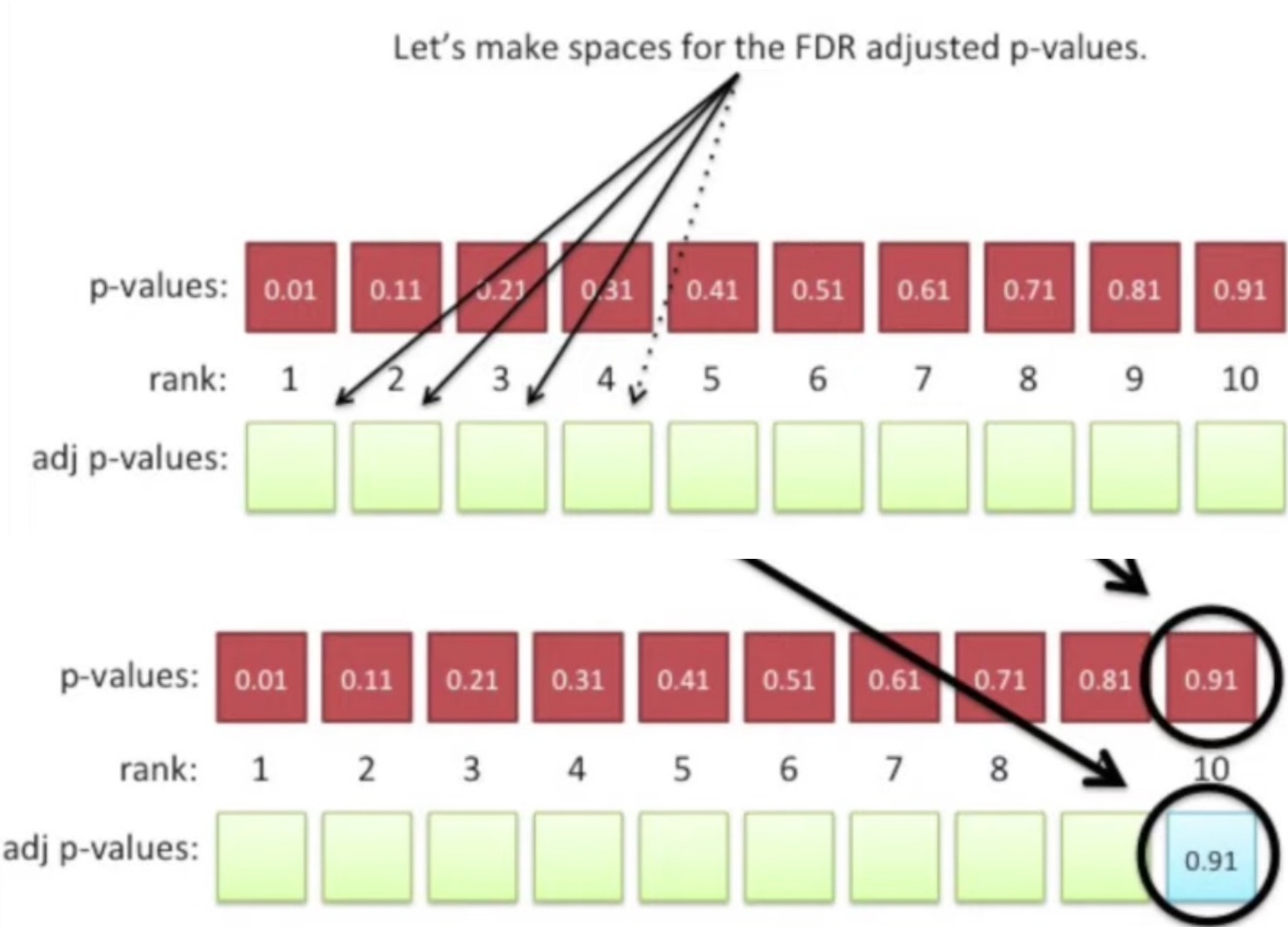


Notice that one of the p-values is a false positive (that is to say, less than 0.05)



**1- Ranking pvalue**

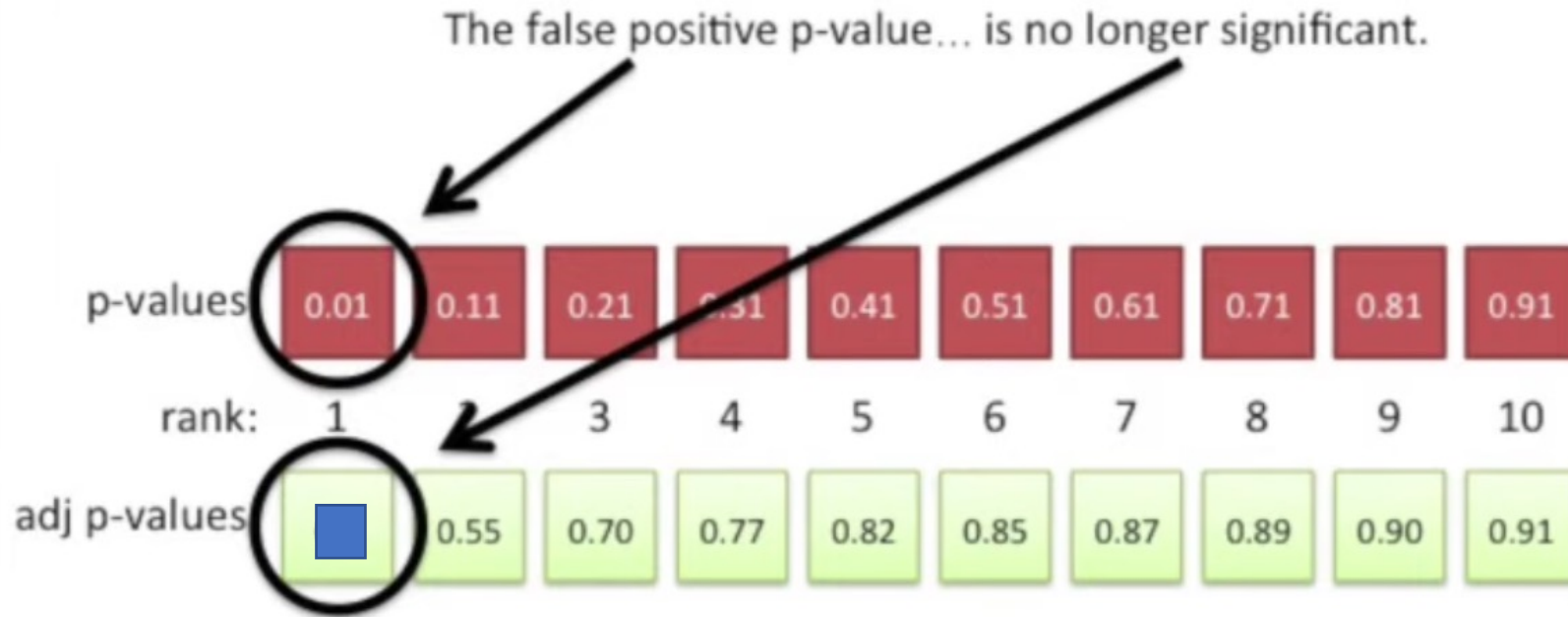
# Prepare space for adjusted p-value



2- Largest adjusted pvalue and larger pvalue are same



# Finally...



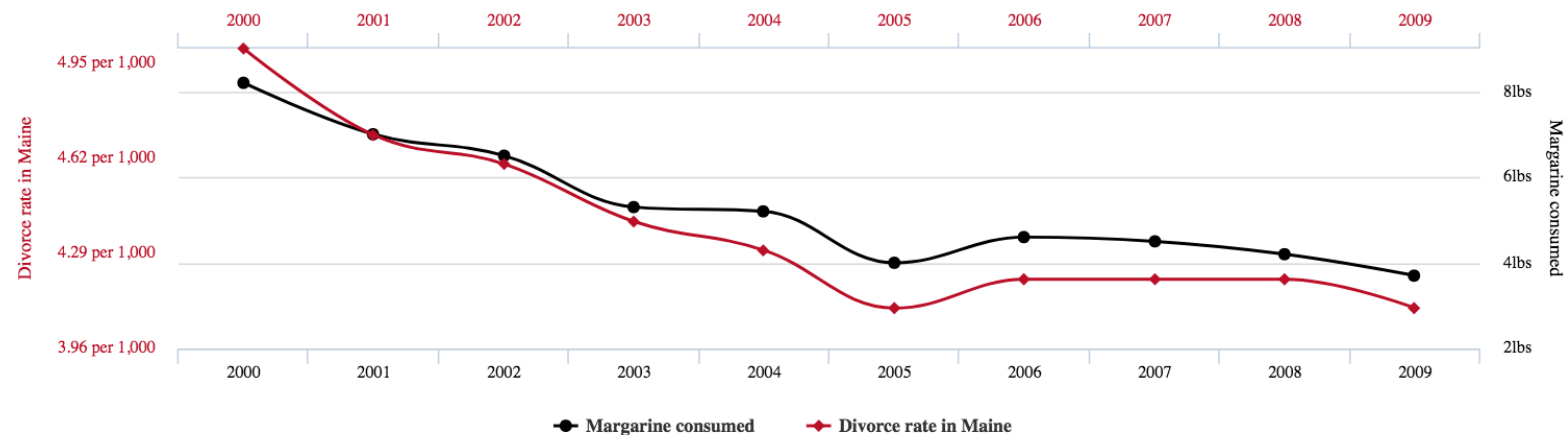
# **Correlation & linear Regression as Bivariate Analyses**

**Objective :** Analyze the **link** that may exist between **two variables** (here: **quantitatives**)  
(Two qualitative variables -> Khi2 test)

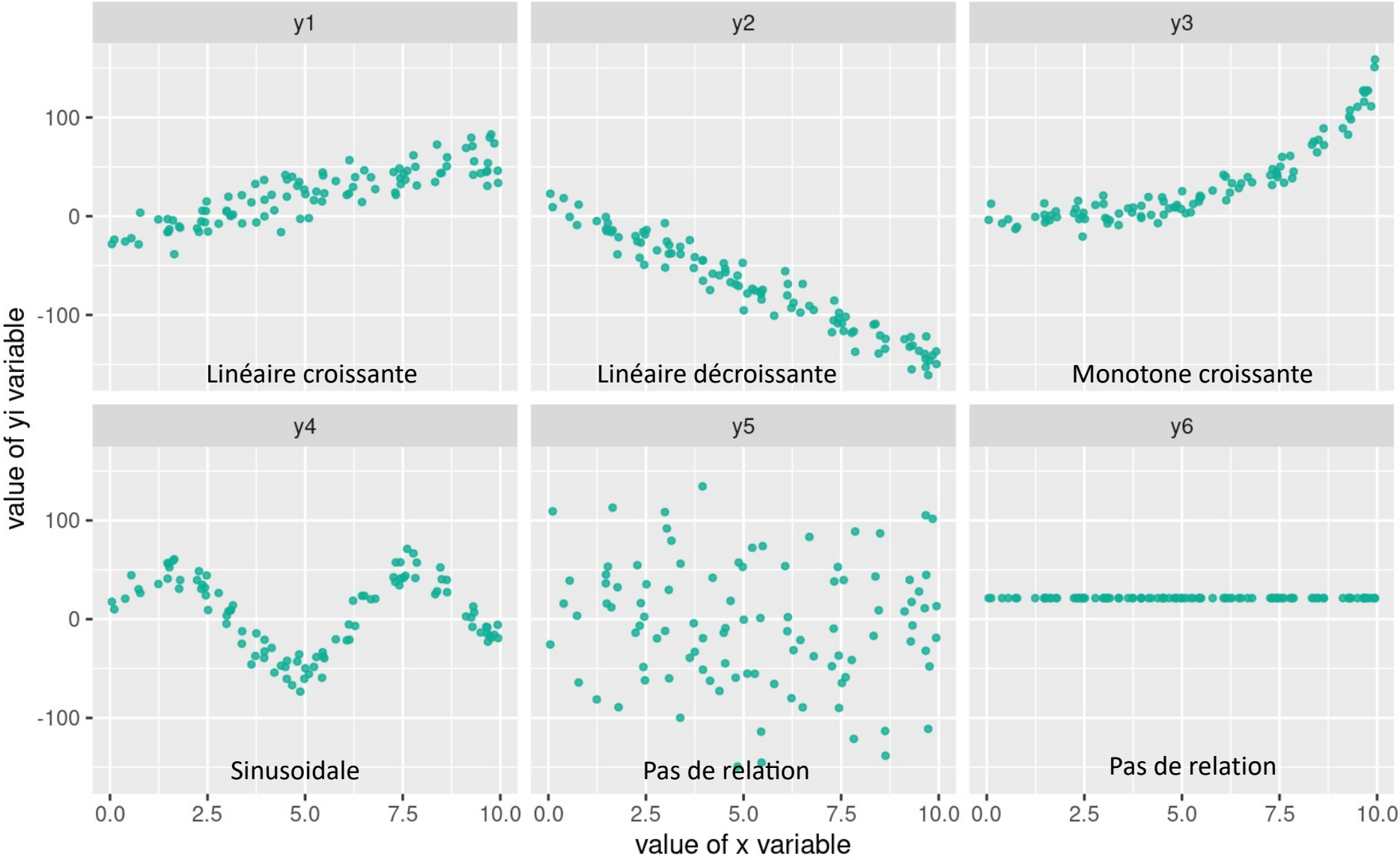
**Link/relationship/dependence** between the variables

→ The values of two variables **do not evolve independently** but on the contrary, present a certain form, a certain regularity

→ Intensity of the association does not indicate causality ...



# What are the relationship between the variables in each graph?

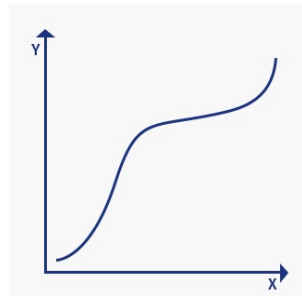


# About Correlation Coefficient r

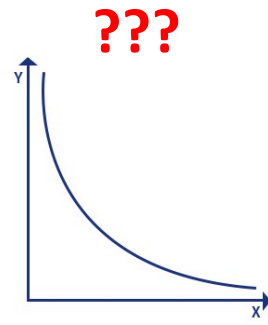
## Intensity & Direction of the association between two variables

### Methods available:

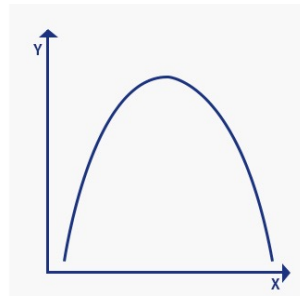
- **Strict Linear Relationship** : **Pearson** ( $r$ , parametric)
- **Monotonous relationship, non-parametric** : **Spearman** ( $Rho$ , rank-based)  
**Kendall** ( $Tau$ ), Alternative to Spearman (small sampling)



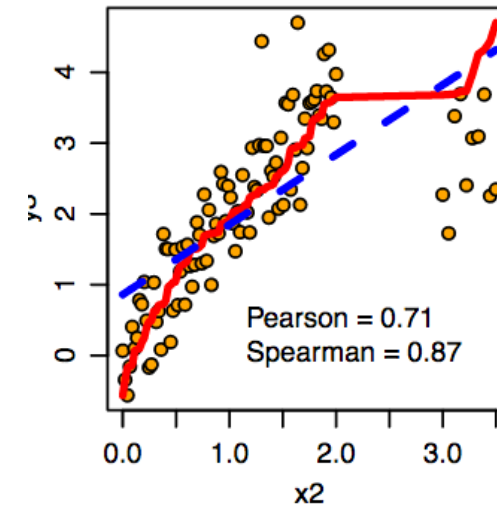
Monotone  
croissante



Monotone  
décroissante



point d'inflection :  
Unimodale

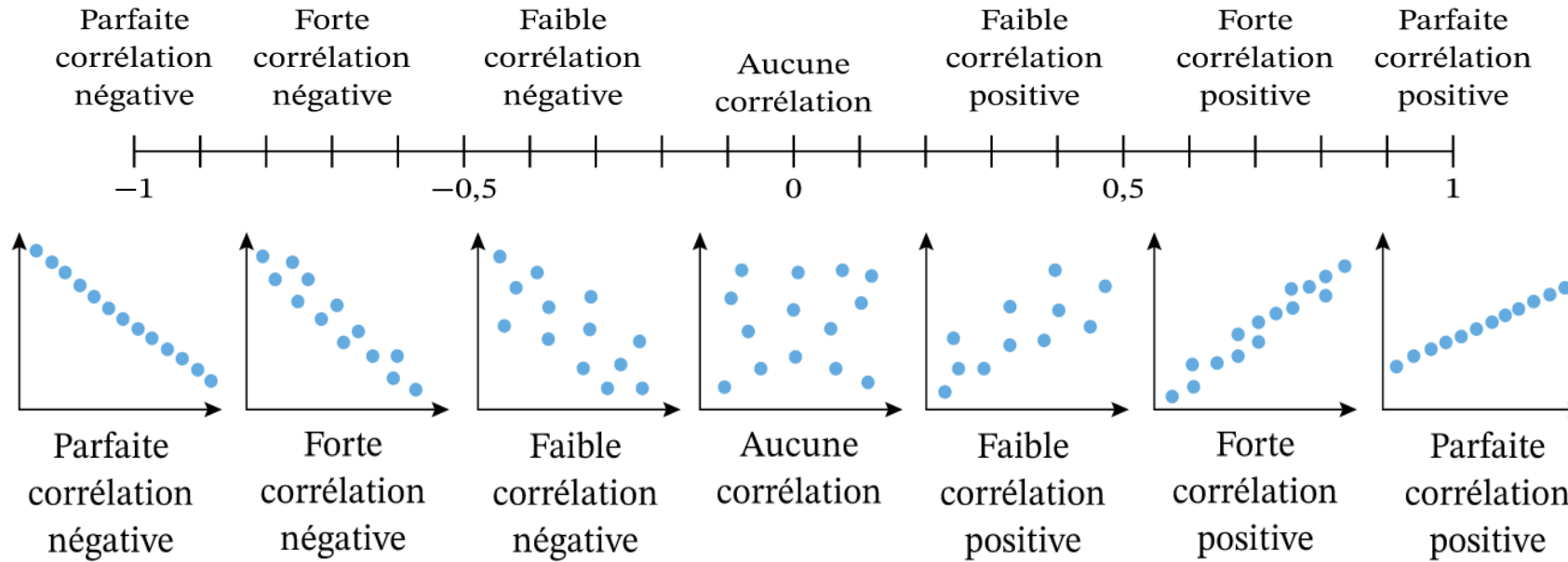


### Coefficient r range between -1 et 1

- **Positive correlation** : The values of both variables tend to increase together
- **Negative correlation** : The values of one variable tend to increase and the values of the other variable decrease
- **Zero** : no **LINEAR** association (**Pearson**)

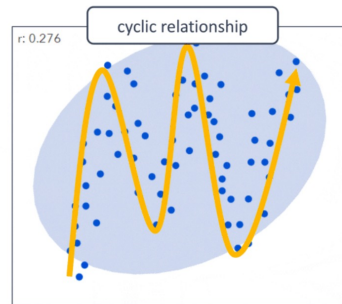
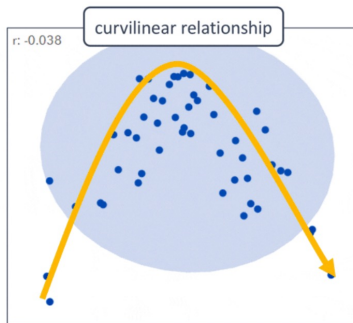
Distribution law of  $r$  under the  $H_0$  hypothesis (No statistical link between X and Y) → Access to p-values to check significance

# For information!!!

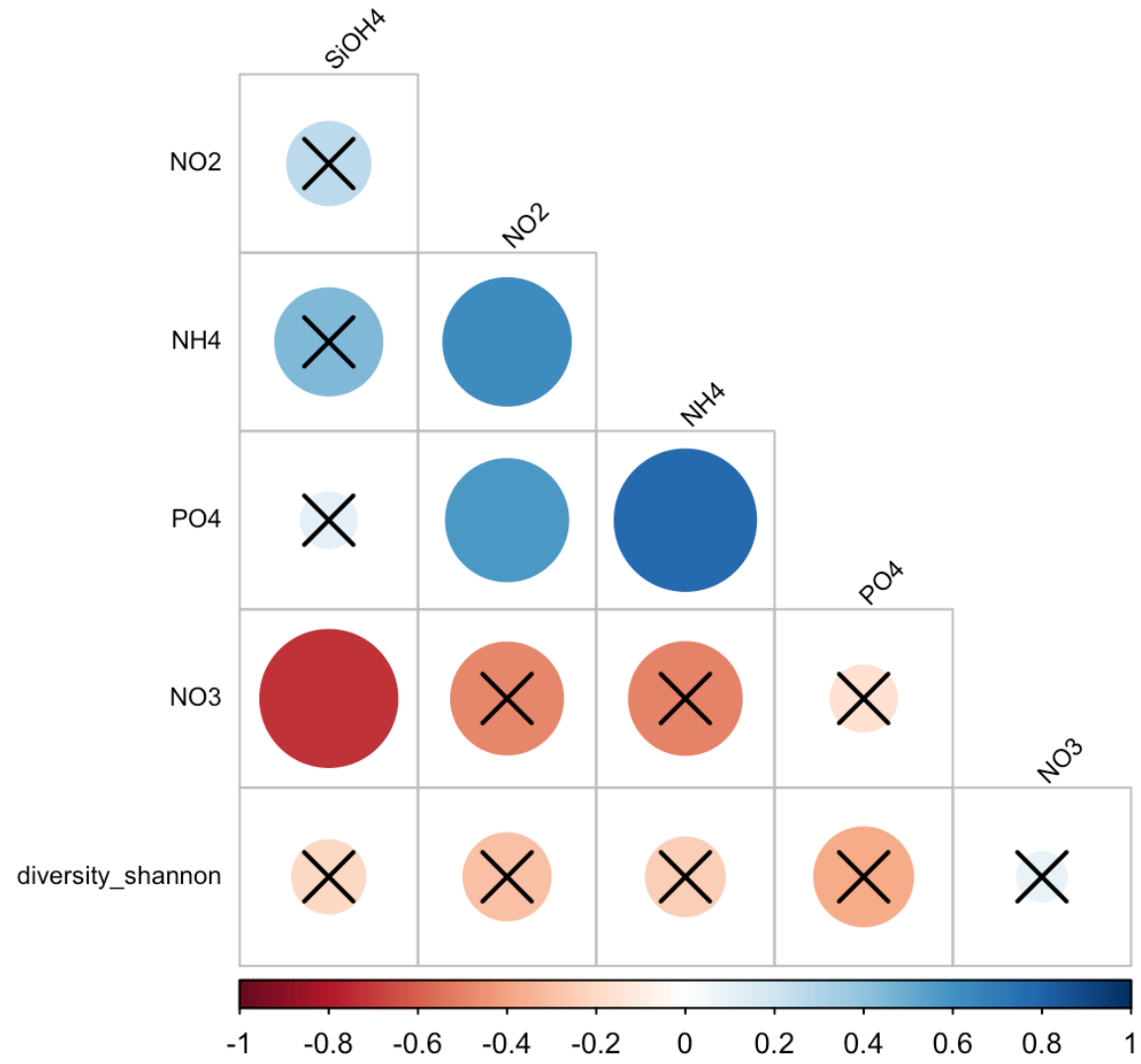


## Because inspecting your results graphically is never useless...

- $r$  close to Zero: no association??



Example of bivariate correlation plot  
(with significant test)



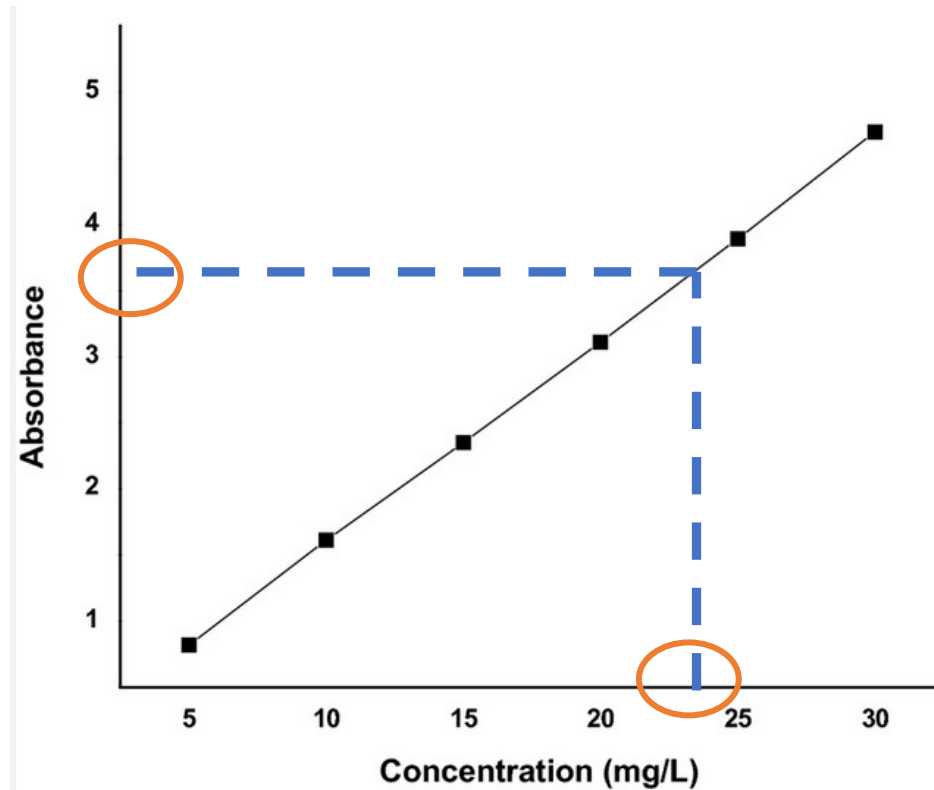
# About simple Linear Regression

- Only for quantitative variables
- Plot the scatter plot Is there a **relationship**?
- Is it **linear**?
- What **orientation** (positive, negative)?
- If the association is **linear** → Make a **regression**

## **Requieurement**

- Normal distribution
- Variance homogeneity

# Your favorite linear regression... calibration curve!!!

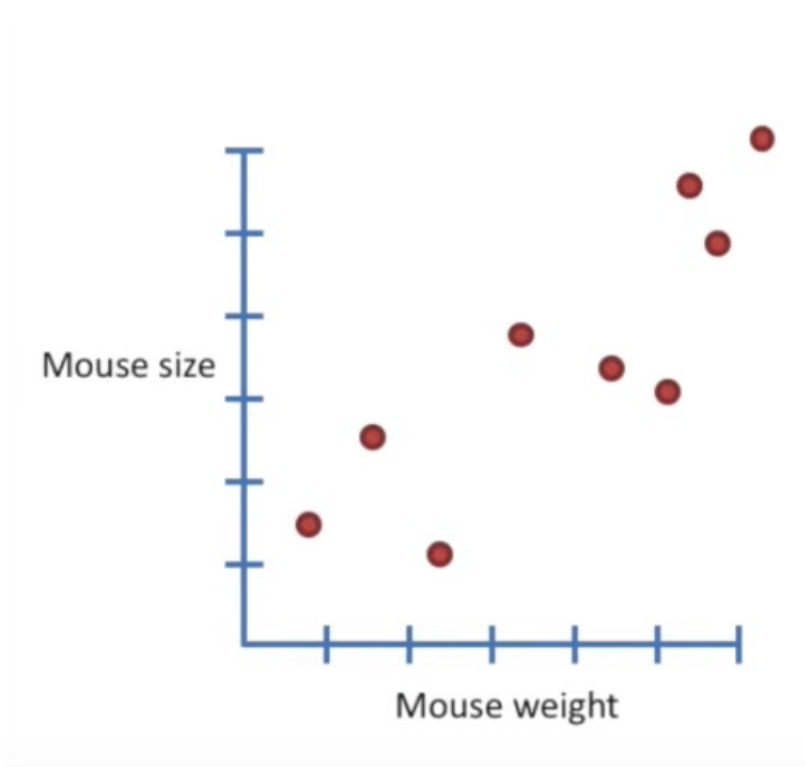


Explain and predict!

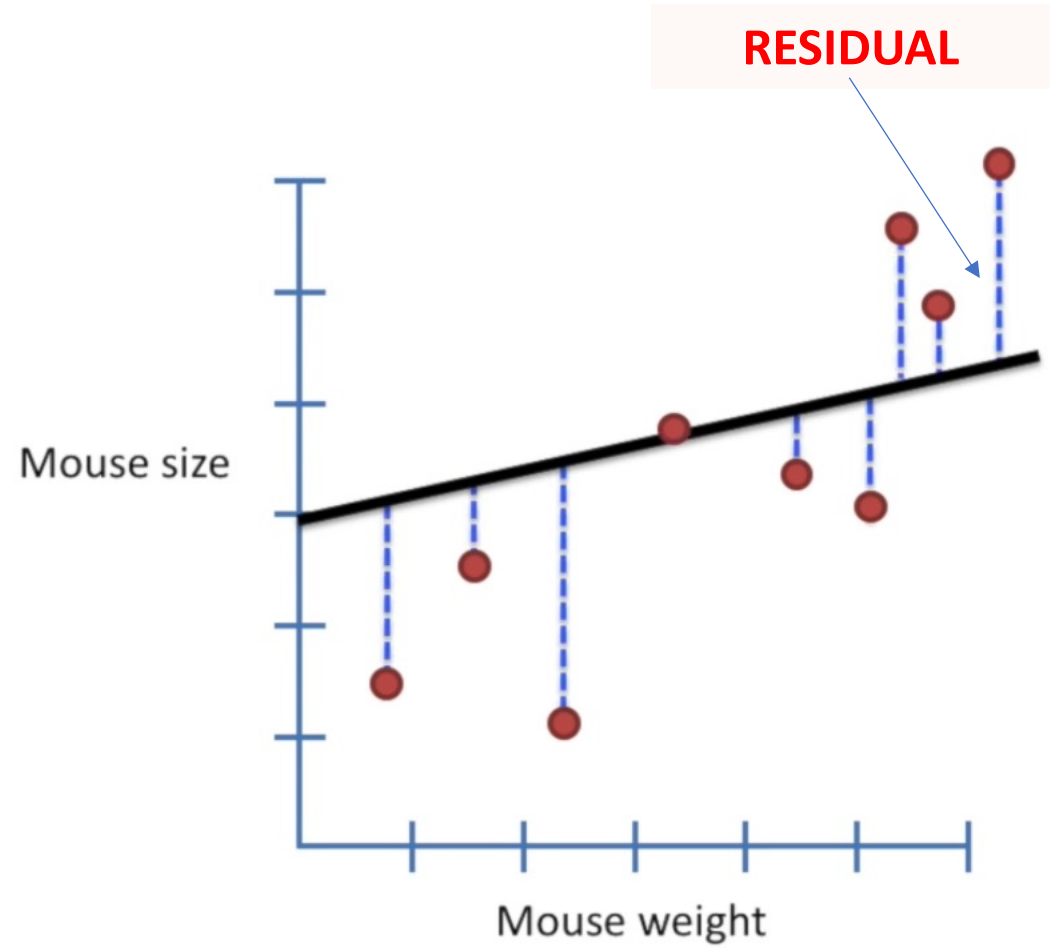
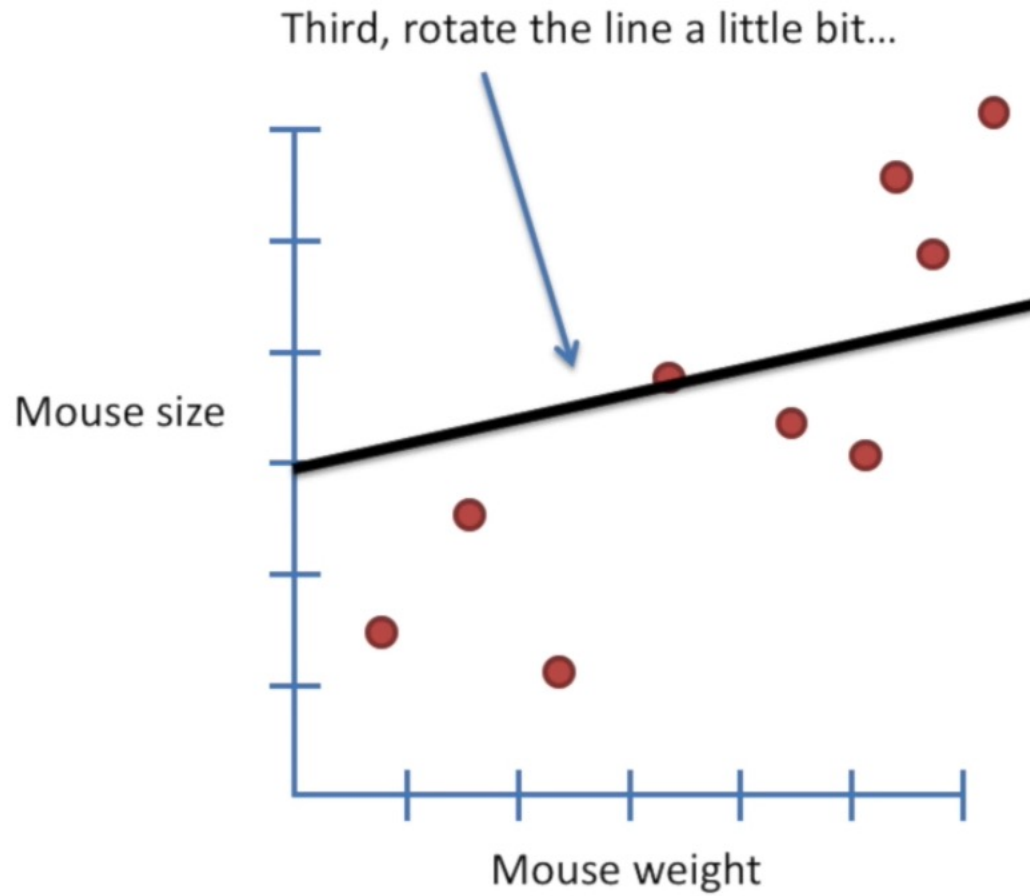
Models a linear type relationship ( $Y=aX+b$ )

Model seeking to establish a **linear relationship** between a variable, called **explained/dependent (Y)**, and another called **explanatory/independent (X)**

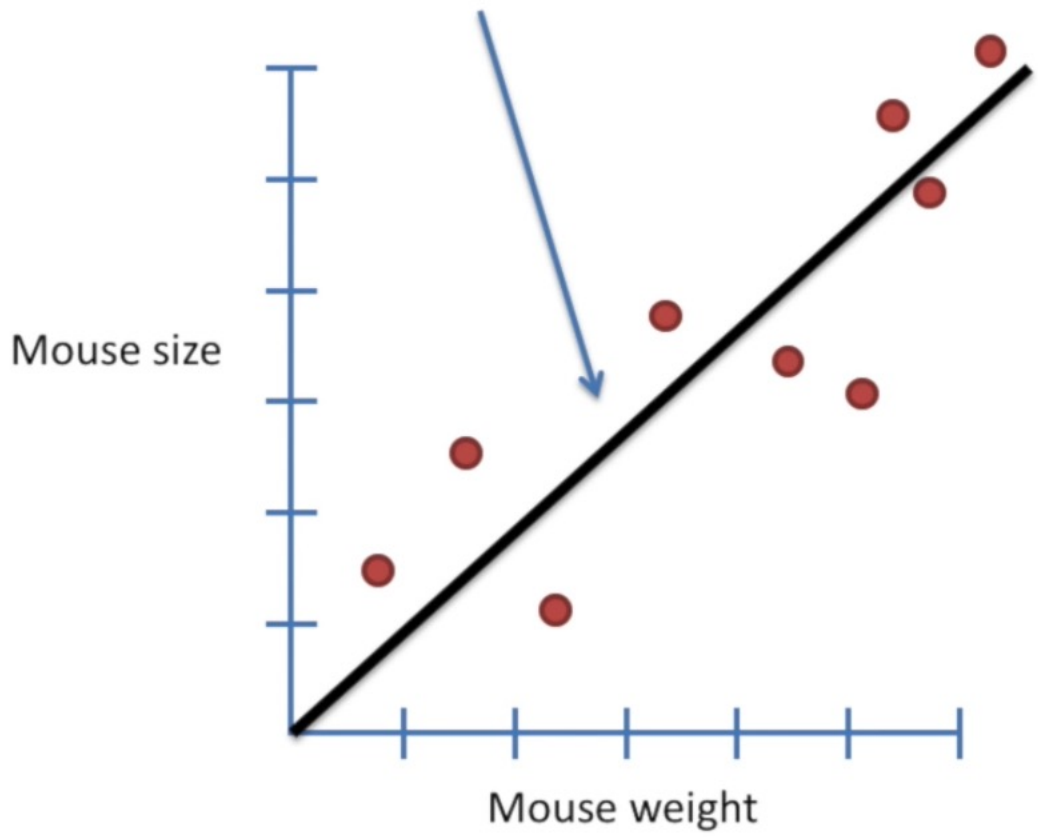
**Can mouse **Weight** predict **Size** correctly? ( $R^2$ )**  
**Relationship is due to chance? (p-value)**



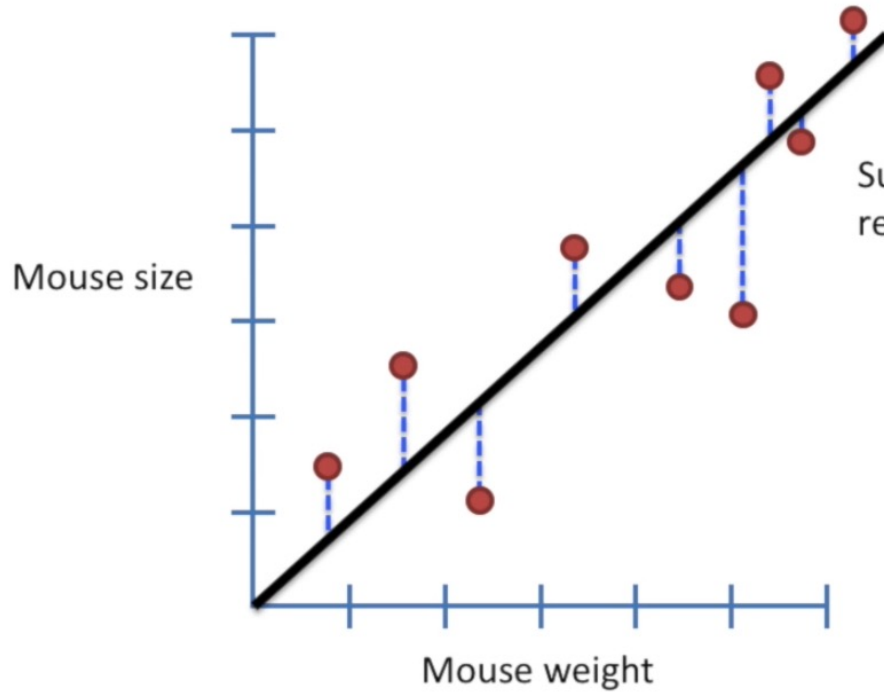
# Least square method



Rotate the line a little bit more...

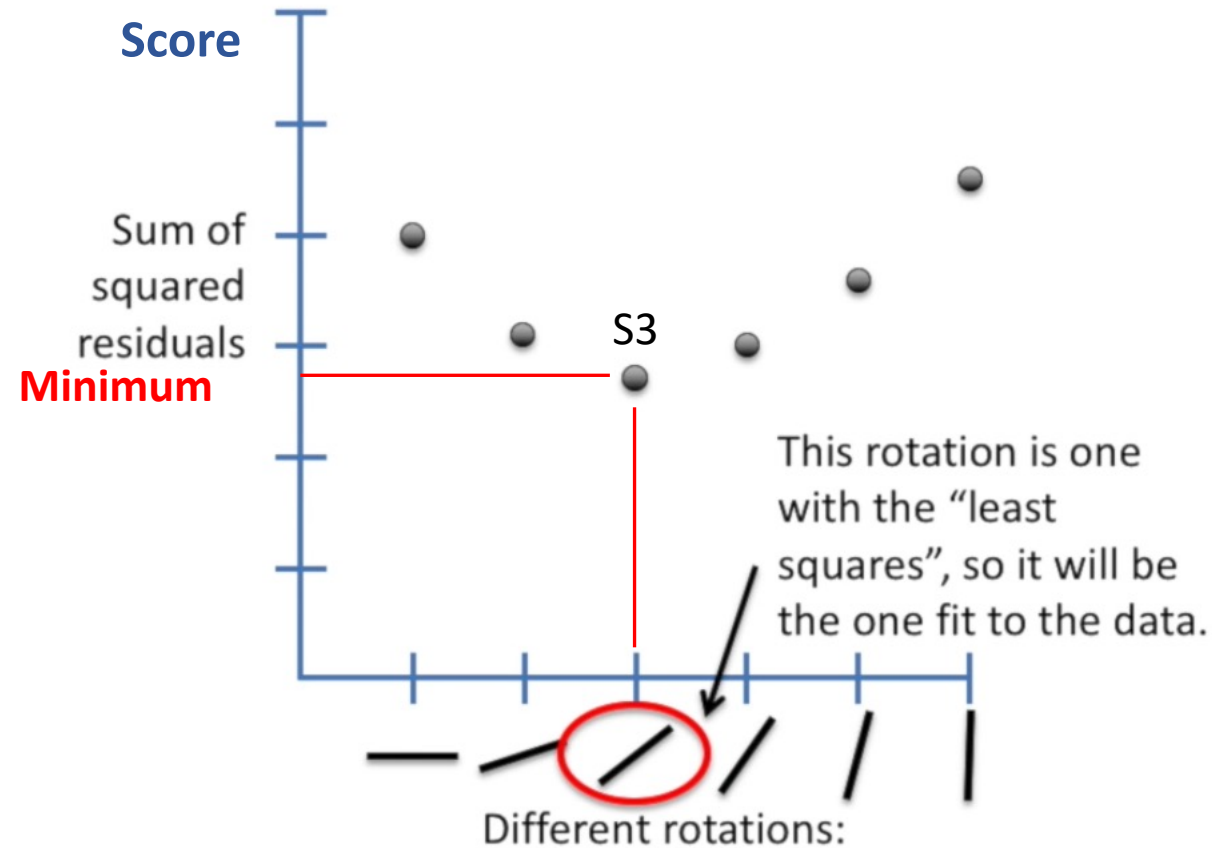


Sum up the squared residuals...

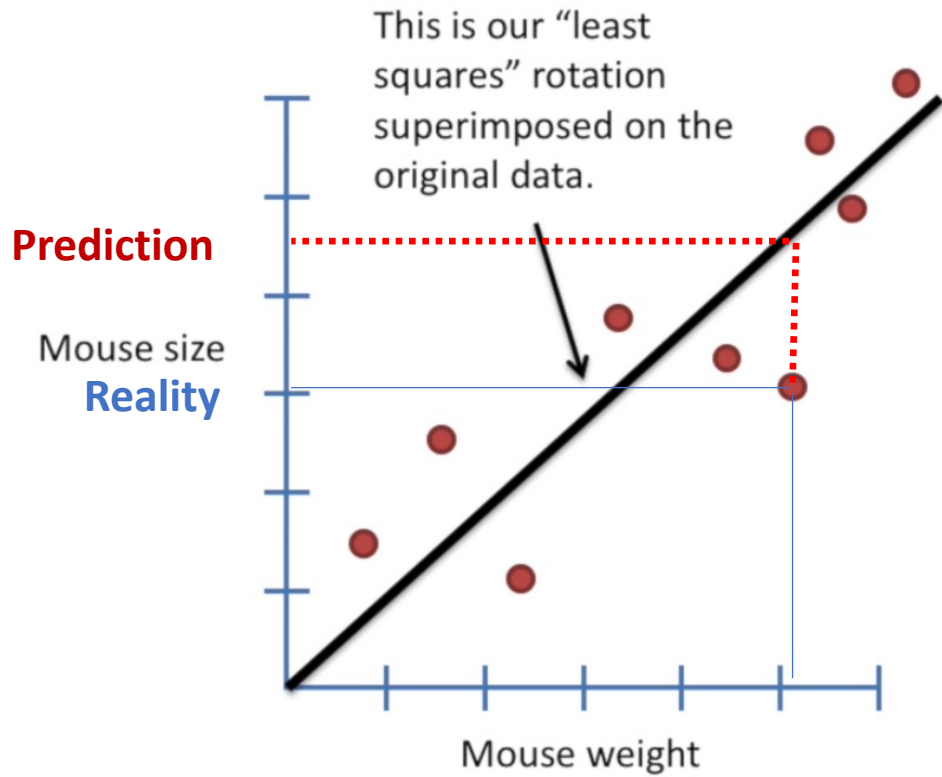


**Again & again, recalculate**

## Resume : Sums of squared residuals for each rotation



**Best rotation (=line position), the one which minimize the score of Sums of squared residuals !!!!**



$$y = 0.1 + 0.78x$$

Dependence to « Mouse weight »

**Coefficient  $R^2$  = prediction quality**

**how good is the linear regression model to predict Mouse size taking into account Mouse weight!!**

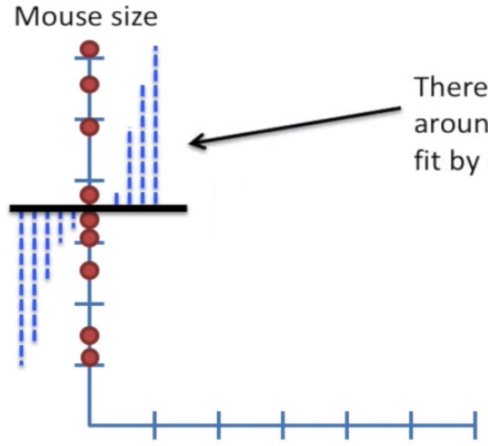
# R<sup>2</sup> : Determination Coefficient

$$R^2 = \frac{\text{Var}(\text{mean}) - \text{Var}(\text{fit})}{\text{Var}(\text{mean})}$$

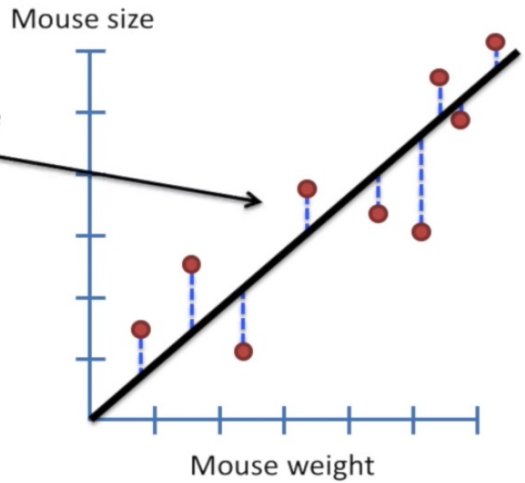
$$R^2 = \frac{\text{Variation expliquée}}{\text{Variation totale}}$$

$$\text{Var}(\text{mean}) = \frac{\text{SS}(\text{mean})}{n}$$

$$\text{Var}(\text{mean}) = 11.1$$



There is less variation around the line that we fit by least-squares.



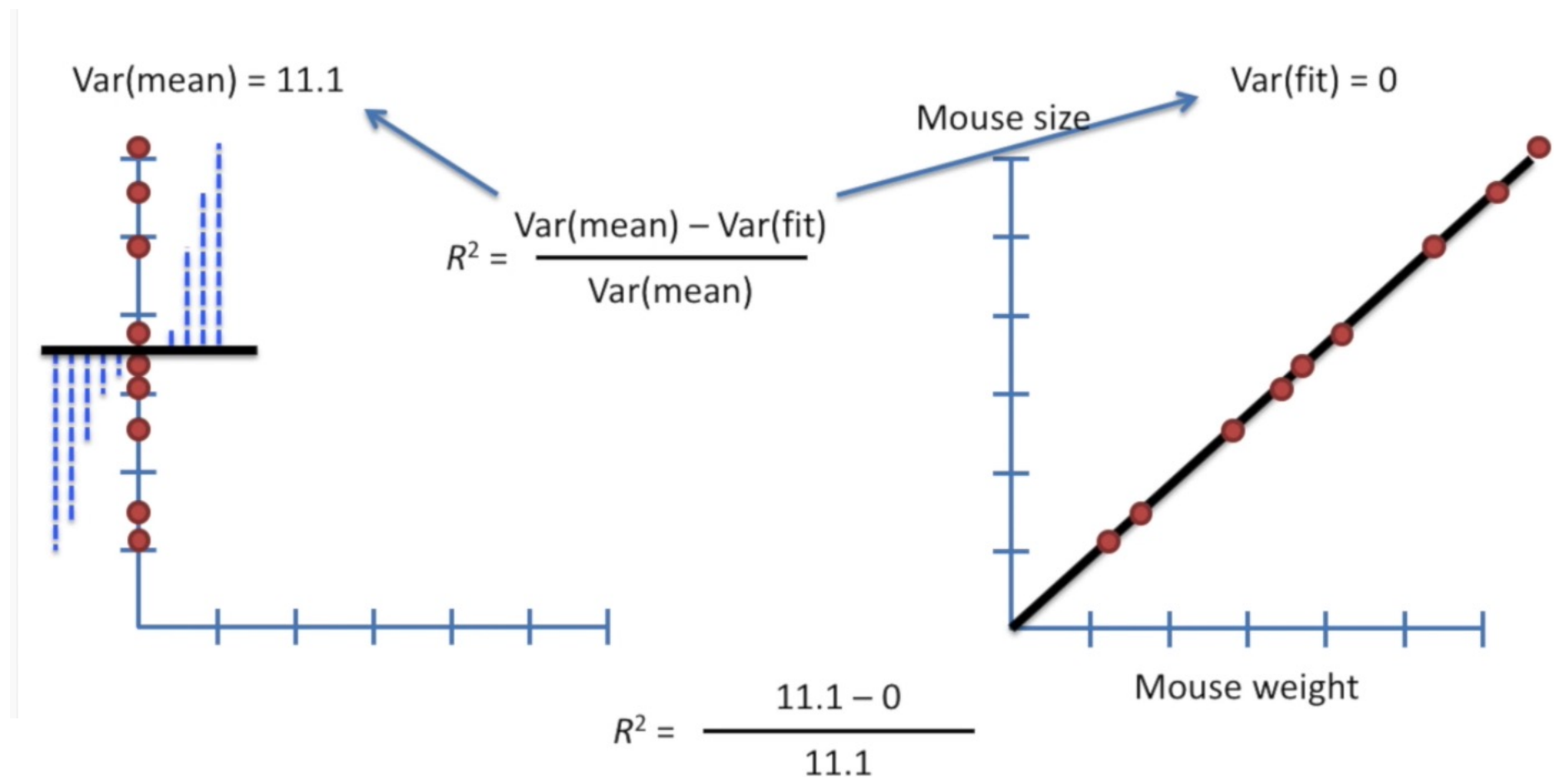
$$\text{Var}(\text{fit}) = \frac{(\text{data} - \text{line})^2}{n}$$

$$\text{Var}(\text{fit}) = 4.4$$

$$R^2 = \frac{11.1 - 4.4}{11.1} = 0.6 = 60\%$$

- R<sup>2</sup> provides percentage variation in Y which is explained by all th X together = % variation of the response variable explained by a linear model (weight variable)
- R<sup>2</sup> between 0 and 1
- Here the established model explains 60% of the variability/variance of the "Mouse size »

# TO be sure ...

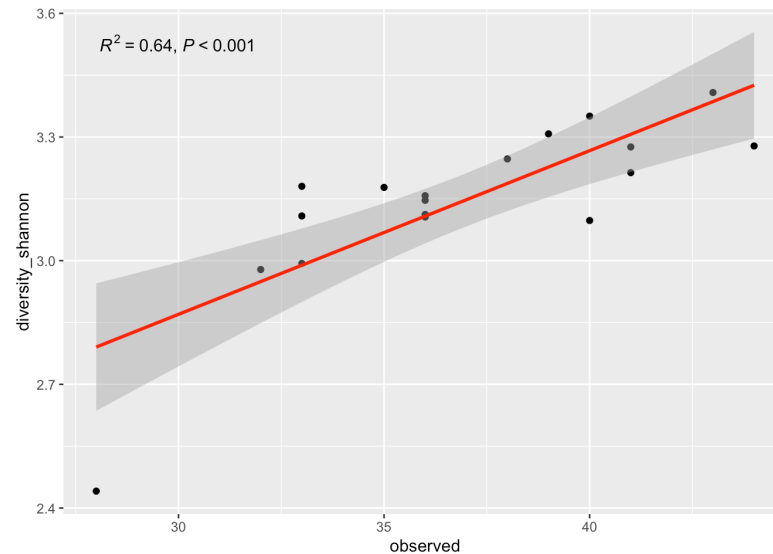


**$R^2 = 1$  = max strength of linear regression model  
(Lesser  $R^2$  the value, the more scattered are the data points)**

# $R^2$ & significance?

$R^2$  based on variance comparison ... so p-value is given by the ratio F & distribution F

$$F = \frac{\text{The variation in mouse size explained by weight}}{\text{The variation in mouse size not explained by weight}}$$



# Relation between $r$ & $R^2$

**Correlation coefficient of Pearson  $r$**  can be linked to linear regression  $R^2$   
**Its square is the explained variance by the regression ( $R^2$ )**

$r = 0.5 \rightarrow R^2 = 0.25 \rightarrow 25\%$  of the Y variance explained by X variable... 😞