



HYPOTHESIS TEST

Bioinformatics Awareness Days @ TIGEM
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Bioinformatics Core: Tasks

- **STATISTICAL DATA ANALYSIS**
Experimental Design, Hypothesis Testing, Differential Expression Analysis, Cluster Analysis, Time Series Data Analysis, Survival Analysis, Correlation Analysis
- **OMICS**
Microarray Analysis, Gene Networks, Pathway Analysis, TFBS Identification, Gene Annotation, Integration, Protein Analysis, Drug Networks
- **NEXT GENERATION SEQUENCING**
Whole Exome, Targeted Gene, RNA, miRNA, ChIP, Visualization, Interpretation
- **DATABASE AND SOFTWARE**
DB Creation, DB Maintenance, Web Sites Creation, Web Service Support
- **BIOINFORMATICS AND (BIO)STATISTICS TRAINING**



Bioinformatics Core: People



<https://www.tigem.it/research/facilities/core-facilities/bioinformatics>

<https://bioinformatics.tigem.it/>

DIEGO DI BERNARDO



DIEGO CARRELLA



ROSSELLA DE CEGLI



SERGIO SARNATARO



EUGENIO DEL PRETE



Bioinformatics Core: Something about Me

- **TLC ENGINEER @ UNIVERSITY OF ROME 'SAPIENZA'**
MAIN TOPICS: Signal Processing, Remote Sensing, Bioinformatics
THESIS: miRNA Analysis, Genomic Data Mining, Consensus Analysis, PSSM Creation
- **BIOINFORMATICS RESEARCH FELLOW @ INSTITUTE OF FOOD SCIENCES (CNR)**
Protein Prediction and Classification, Protein Analysis, Proteomic Mass Spectra Analysis, Sequence Alignment and Phylogenetic Tree, Docking
- **PHD IN APPLIED BIOLOGY @ UNIVERSITY OF BASILICATA**
Celiac Disease and Comorbidities, Microarray Data Analysis, Ontologies, Gene Set Enrichment Analysis, Semantic Similarity, Proteomic Mass Spectra Analysis
- **BIOINFORMATICS RESEARCH FELLOW @ INSTITUTE OF APPLIED MATHEMATICS (CNR)**
Proteomic Mass Spectra Analysis, Metabolomic (Lipidomic) Data Analysis, Web Tools Developer, Hypothesis Tests, Omics Data Integration
- **BIostatistician and Data Analyst @ TIGEM**



Outline

● UNCERTAINTY AND VARIABILITY

- Descriptive Statistics
- Uncertainty and Variability
- Measurement

● HYPOTHESIS TESTING

- Inferential Statistics
- Hypothesis Testing: What, How, Errors, Which
- Multiple Test Correction

● EXAMPLES

- Example One
- Example Two
- Example Three

● CONCLUSION

- Take Home Message
- Final Remarks



Not Only Aphorism...



Trilussa (1871 - 1950)

Carlo Alberto C. M. Salustri

Poet, Writer, Journalist

LA STATISTICA

Sai che d'è la statistica? È 'na cosa
che serve pé fa' un conto in generale
de la gente che nasce, che sta male,
che more, che va in carcere e che sposa.

Ma pé me la statistica curiosa
è dove c'entra la percentuale,
pé via che, lì, la media è sempre eguale
puro co' la persona bisognosa.

**Me spiego: da li conti che se fanno
secondo le statistiche d'adesso
risurta che te tocca un pollo all'anno:
e, se nun entra ne le spese tue,
t'entra ne la statistica lo stesso
perché c'è un antro che ne magna due.**

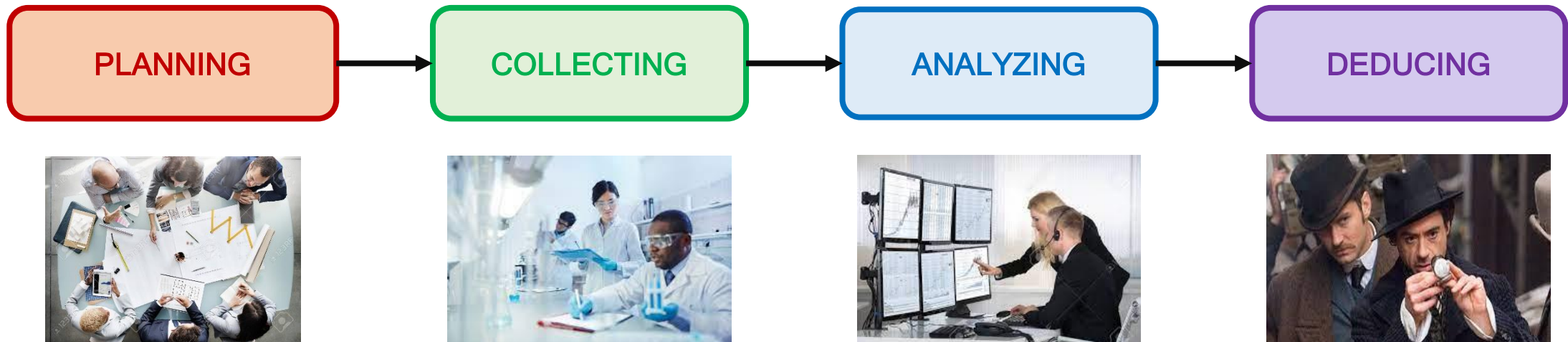


Statistics

Science

- Study of collective and measurable phenomena, with **quantifiable** data
- Answer to a **well-posed** question to find a solution, with a degree of **uncertainty**
- Application of mathematical principles and techniques to **learn** from data

Workflow





Descriptive Statistics

Descriptive Statistics

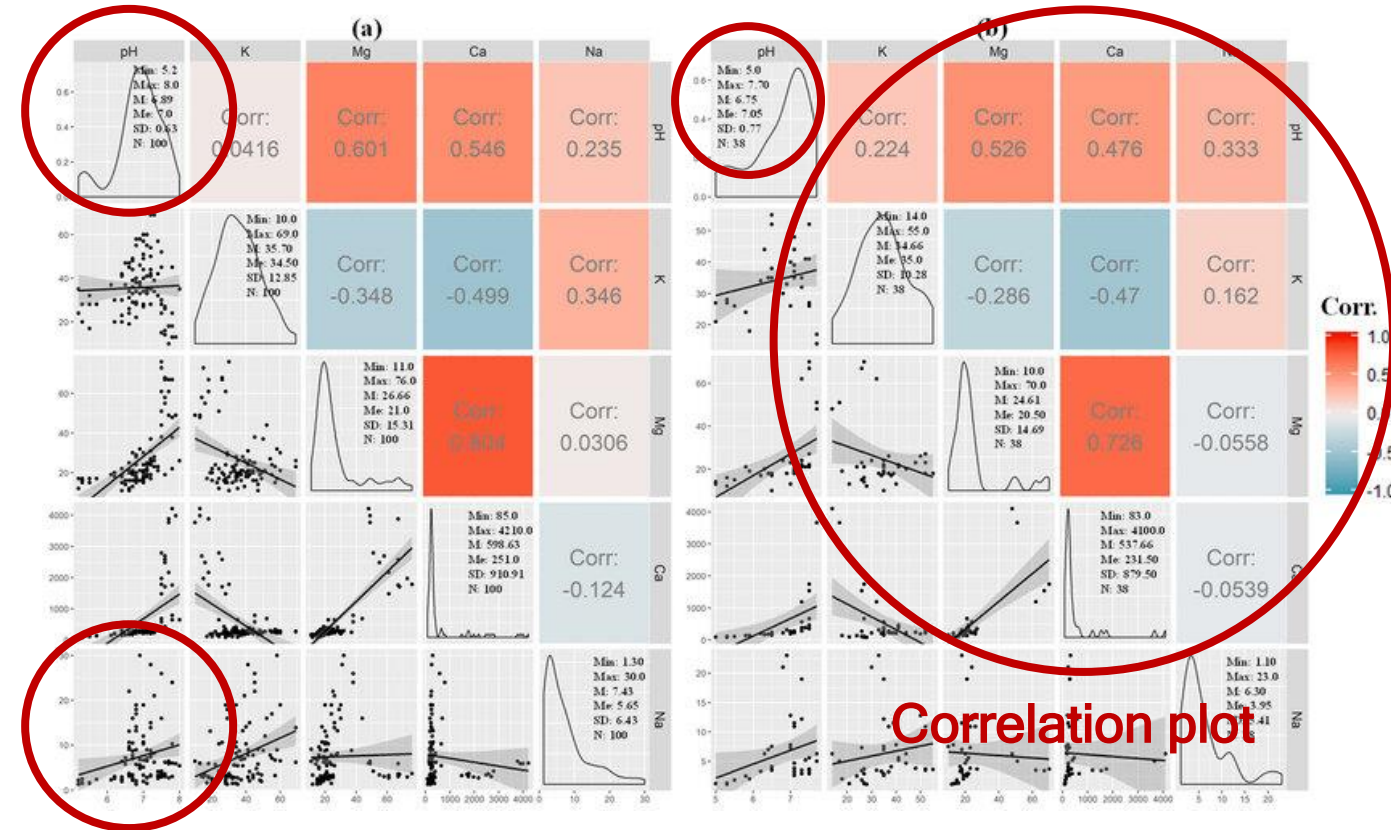
- Description of the features for a specific dataset
- Summary of the information from a specific dataset

Description Tools

- **Plots:** barplot, boxplot, pie chart, scatter plot, density plot, correlation plot
- **Tables:** descriptive table, summary table

Density plot

Summary table



Scatterplot

Correlation plot



Uncertainty

Uncertainty

- COMMON SENSE: not known beyond doubt, not having complete knowledge
- STATISTICAL: probability and repeatability



Example: Coin Flip

- Flip the coin **10 times**: H, H, H, T, T, T, T, H, H, H
- Calculate percentage: H 60%, T 40%
- Flip the coin **1000 times** ($1000 \gg 10$)
- Calculate percentage: H 54%, T 46 %

(Strong) Law of Large Numbers

- X_1, X_2, \dots, X_n is an infinite sequence of independent and identical distributed random variables
- Expected values $E(X_1) = E(X_2) = \dots = E(X_n) = \mu$ and sample average $\bar{X}_n = \frac{1}{n}(X_1 + X_2 + \dots + X_n)$

then

$$\bar{X}_n \rightarrow \mu \text{ when } n \rightarrow \infty$$

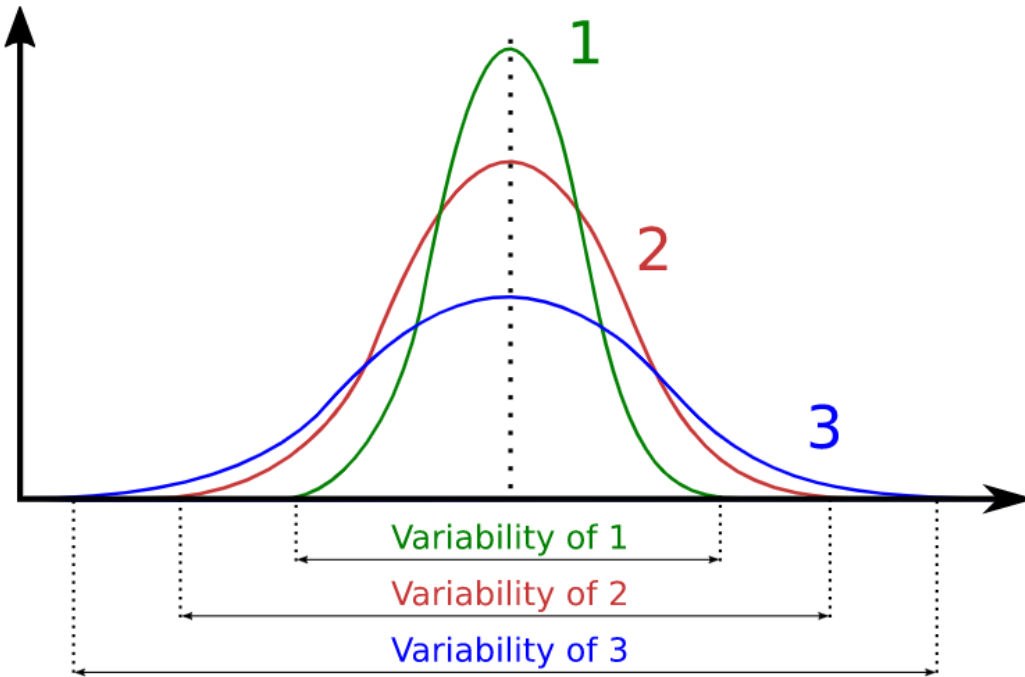


Variability

Variability

- COMMON SENSE: different values in a particular condition
- STATISTICAL: divergence of data from its mean value (spread, dispersion)

Normal distribution



Sample Mean: $\bar{X}_n = \frac{1}{n} \sum_{i=1}^n X_n$

Sample Variance: $\sigma_n^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X}_n)^2$

Sample Error: $\sigma_{\bar{X}} = \frac{\sigma_n}{\sqrt{n}}$

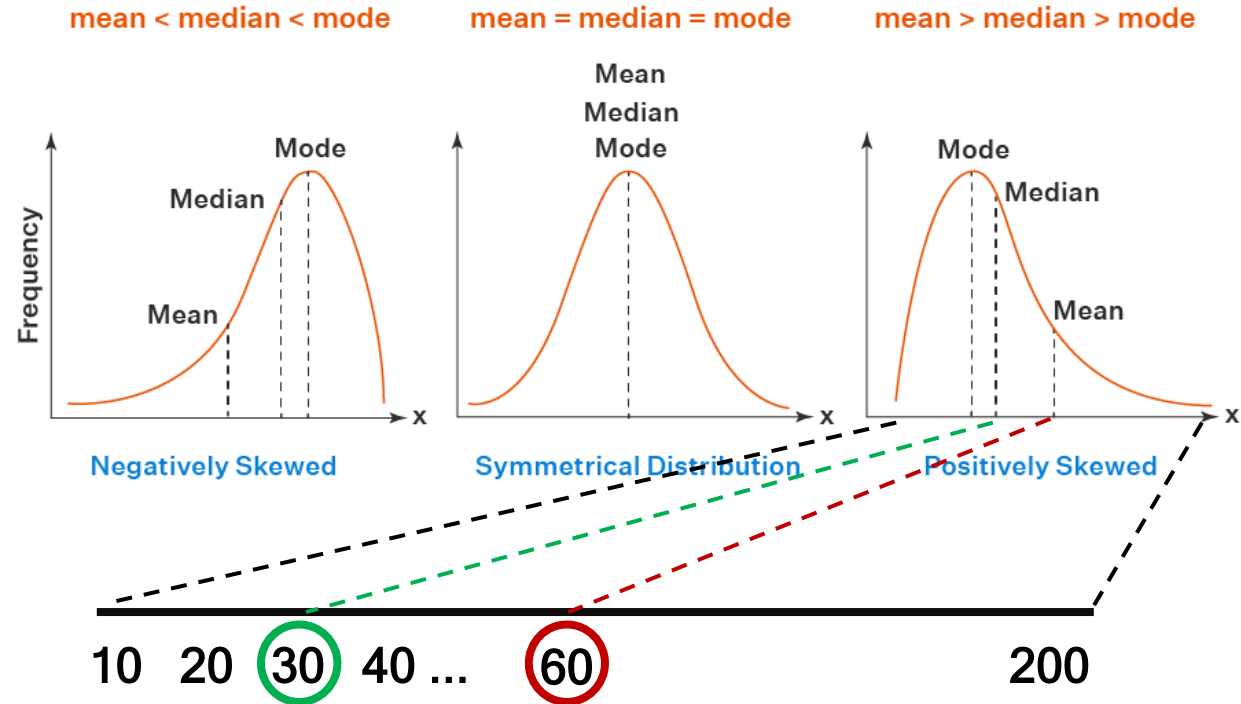
WHAT ABOUT σ_n ?
WHAT ABOUT n ?

(See GALTON'S BOARD)



Measure of Central Tendency

- Mode**
 Most frequent value in the data set
 (nominal data)
- (Arithmetic) Mean**
 Sum of all measurements divided by the number of observations in the data set
- Median**
 Middle value that separates the higher half from the lower half of the data set
 (ordinal data)



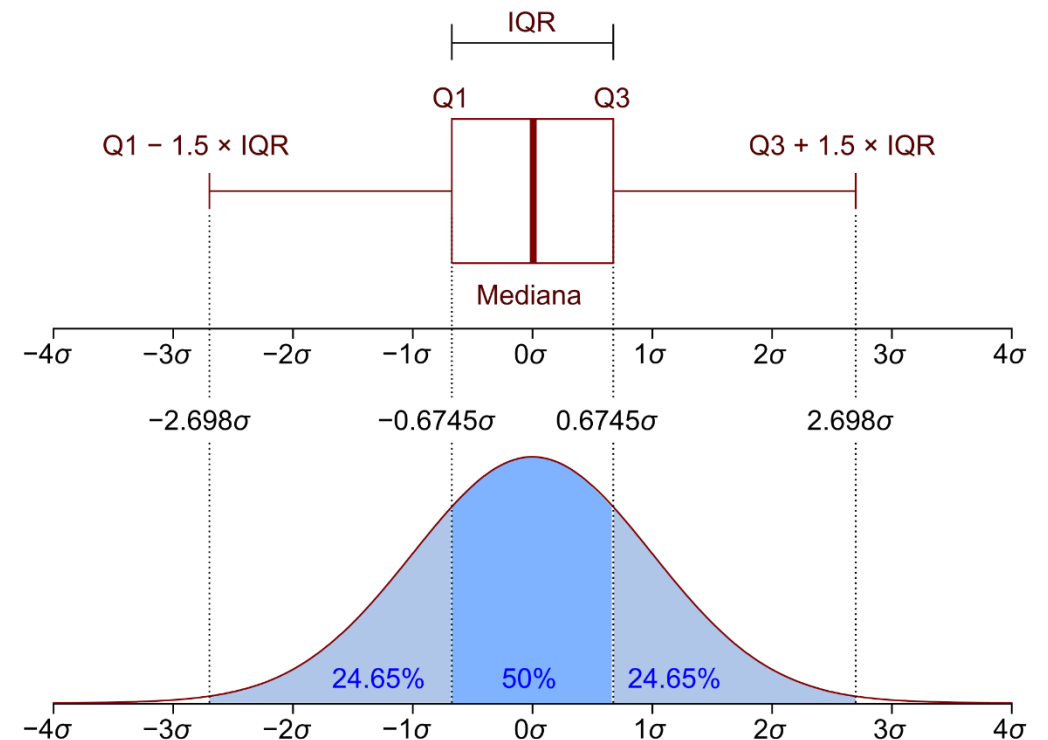


Measure of Variability

- Range**
 Difference between the smallest and the largest value in the data set
- Standard Deviation (SD)**
 How data is spread out going from the mean
- Coefficient of Variation (CV)**
 Relative dispersion of data around the mean

$$c_v = \frac{\sigma}{\mu} (\times 100)$$
- InterQuartile Range (IQR)**
 How widespread the interval is, in which the middle 50 % of all the values lie

- SD is the square root of sample variance
- CV is a normalization (dimensionless)





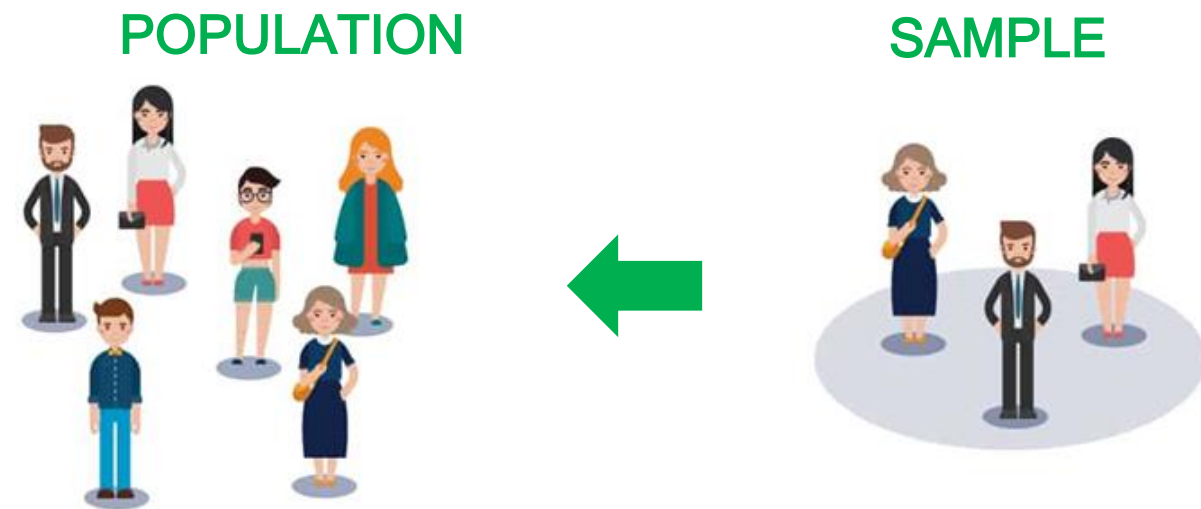
Inferential Statistics

Inferential Statistics

- Assumption from the features of a specific dataset and **validation**
- Statistical methods for inferring the characteristics of a population (**parameter**) from a sample (**statistic**)

Estimation

- Measure a statistic from the sample
- Generalize to the population:
 - a) approximate estimation (**margin of error**)
 - b) sample \neq population (**probability of error**)





Hypothesis Testing: What and How

Hypothesis Testing

- An analyst **tests** an assumption regarding a population parameter
- The methodology employed depends:
 - a) on the **nature of the data** used
 - b) on the **reason for the analysis**

How to test a hypothesis

1. State null hypothesis H_0

Children who take vitamin C are no less likely to become ill during flu season

2. State alternative hypothesis H_1

Children who take vitamin C are less likely to become ill during flu season

3. Determine significance level α

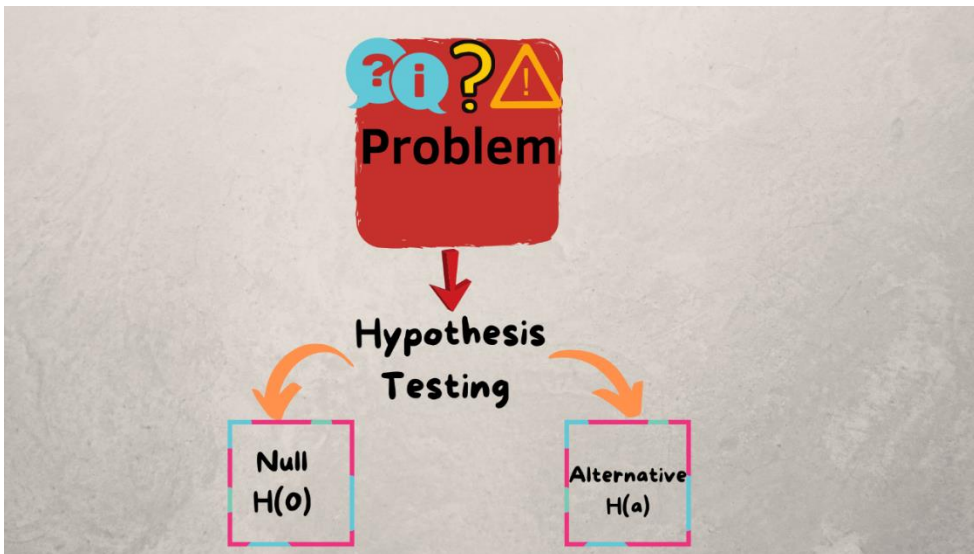
Percentage of error be willing to accept (5%)

4. Calculate H_0 probability p-value

One group with vitamin C during flu season and the other with a placebo. Collecting a p-value of 0.1

5. Reject or not H_0

$P\text{-value} > \alpha$, H_0 cannot be rejected





Hypothesis Testing: Types of Error

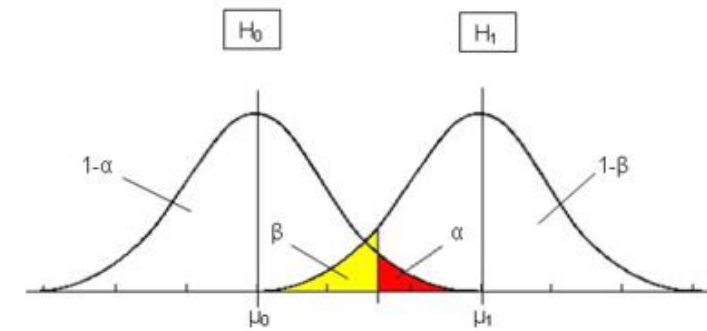
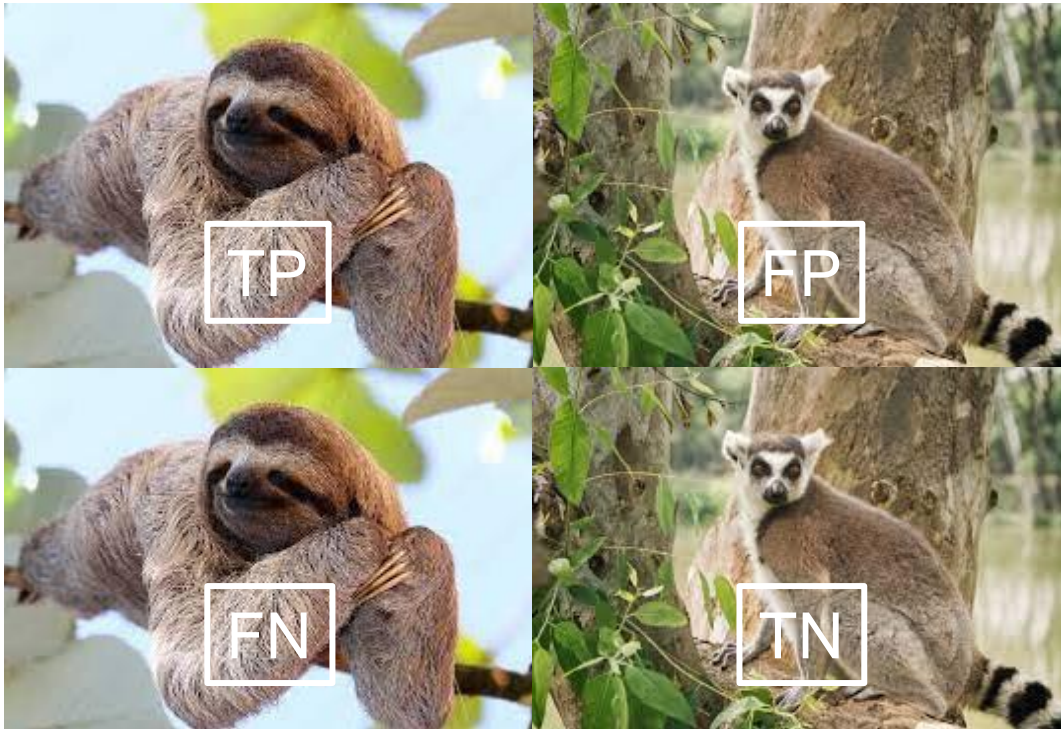
● H0: LEMUR (NOT SLOTH)

REALITY

H0 FALSE

H0 TRUE

STUDY FINDINGS
 REJECT H0
 ACCEPT H0



TRUE POSITIVE (TP) - POWER (1-β)
 Probability to REJECT H0 when H0 is FALSE

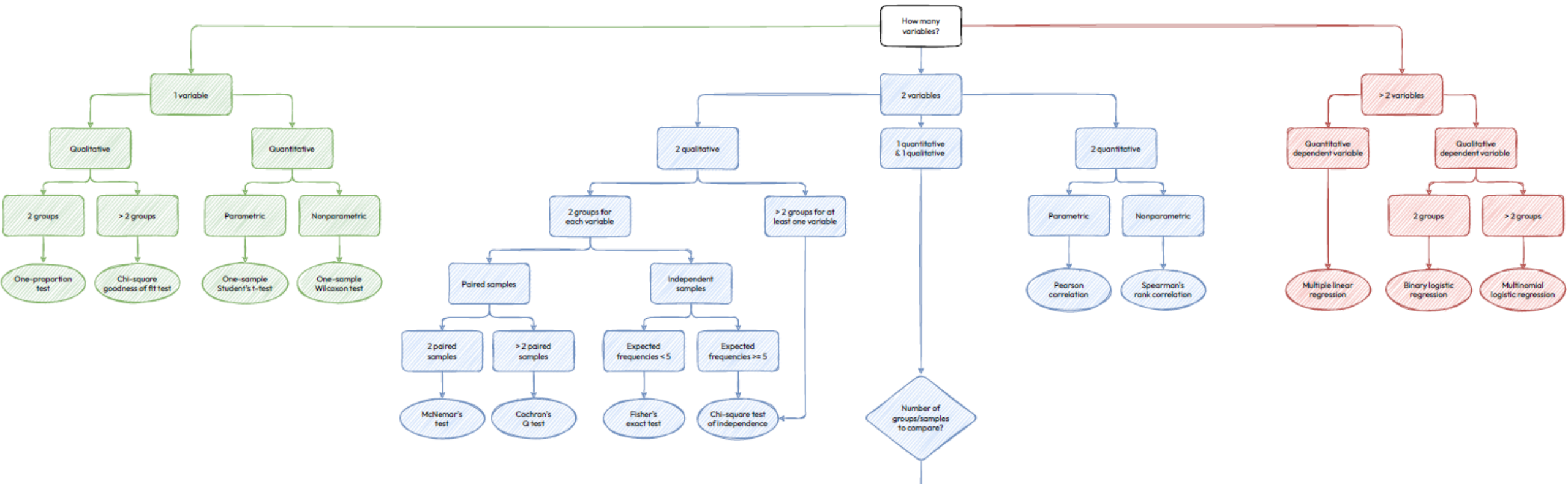
FALSE POSITIVE (FP) - TYPE I ERROR, α
 Probability to REJECT H0 when H0 is TRUE

FALSE NEGATIVE (FN) - TYPE II ERROR, β
 Probability to ACCEPT H0 when H0 is FALSE

TRUE NEGATIVE (TN)
 Probability to ACCEPT H0 when H0 is TRUE

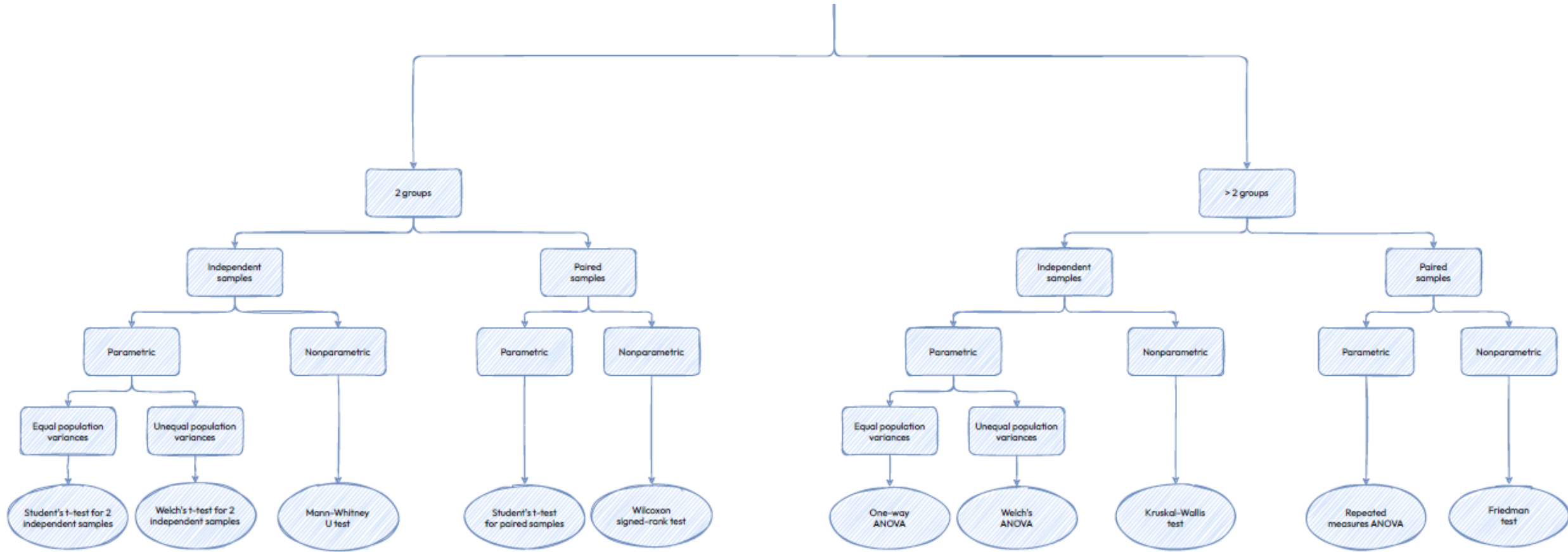


Hypothesis Testing: Which





Hypothesis Testing: Which





Hypothesis Testing: Which

- **QUESTION 1: Which kinds of variable?** CONTINUOUS, DISCRETE, CATEGORICAL
- **QUESTION 2: How many groups per variable?** 1 GROUP, 2 GROUPS, > 2 GROUPS
- **QUESTION 3: Are the samples paired?** UNPAIRED, PAIRED
- **QUESTION 4: Are the distributions normal?** PARAMETRIC, NON-PARAMETRIC
- **QUESTION 5: Have the distributions the same variance?** HOMOSCEDASTICITY



Hypothesis Testing: Multiple Test Correction

● Probability of At Least One Type I Error

$$Pr(\alpha|m) = 1 - (1 - \alpha)^m$$

● Two Methods

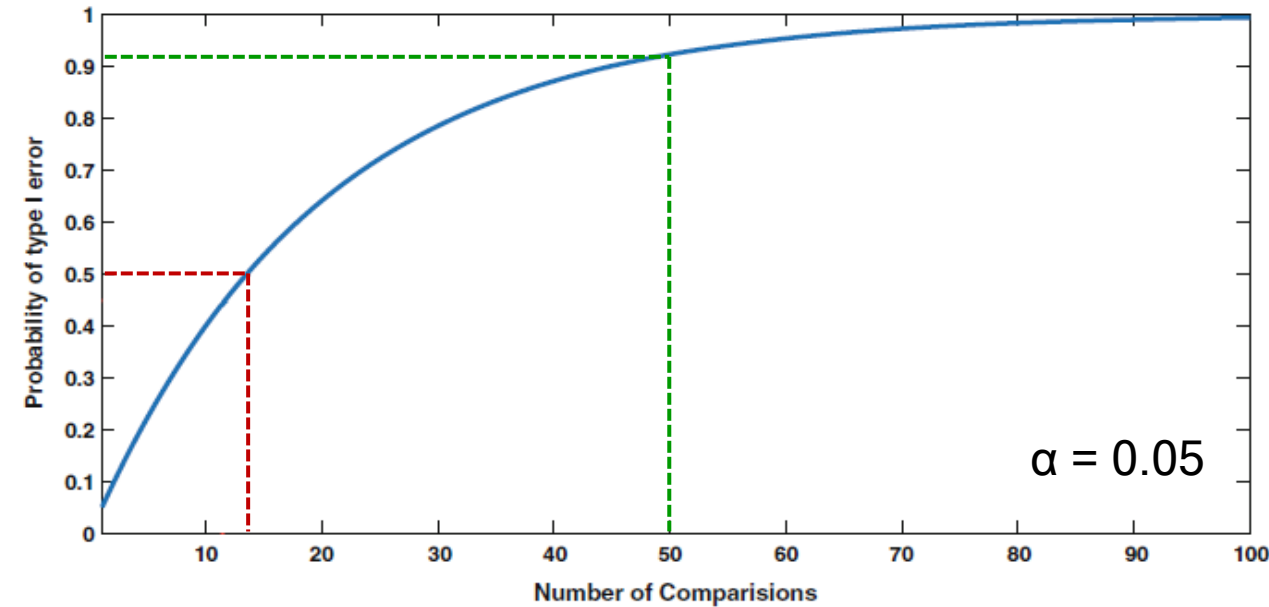
Hard Correction: BONFERRONI

- $p_B = p_j m < \alpha$
- Control Family-Wise Error Rate
- Loss of power due to large number of tests

Soft Correction: BENJAMINI-HOCHBERG

- $p_1 < p_2 < p_j \dots < p_m \rightarrow p_{BH} = \frac{p_j m}{j} < \alpha$
- Control False Discovery Rate
- Flexible procedure

Example: Identification of Differential Expressed Genes from RNASeq Data



$\alpha = 0.05, m = 1 \rightarrow Pr(\alpha = 0.05|m = 1) = 0.05$

$\alpha = 0.05, m = 13 \rightarrow Pr(\alpha = 0.05|m = 13) = 0.49$

$\alpha = 0.05, m = 50 \rightarrow Pr(\alpha = 0.05|m = 50) = 0.92$



Example One

- Suppose to test if the **average weight** of **10 mice** differs from **25 mg**

QUESTIONS

- Which kinds of variable?
- How many groups per variable?
- Are the samples paired?
- Are the distributions normal?
- Have the distributions the same variance?
- Multiple test correction?

ANSWERS

- One continuous variable
- One group
- No (one measurement)
- Yes
- Not relevant
- No (one test)

Dataset	
Name	Weight
M_1	20.6
M_2	20.0
M_3	20.4
M_4	22.0
M_5	19.9
M_6	20.7
M_7	18.8
M_8	20.5
M_9	20.4
M_10	23.3

d) D'Agostino-Pearson test (0.1139), Shapiro-Wilk test (0.1634) → Answer: Yes

- **TEST**: One sample Student's t-test (< 0.0001 , ****) → Answer: Yes



Example Two

● Suppose to test if **two different treatments** affect the weight of the mice

QUESTIONS

- Which kinds of variable?
- How many groups per variable?
- Are the samples paired?
- Are the distributions normal?
- Have the distributions same variance?
- Multiple test correction?

ANSWERS

- Two variables: one continuous, one categorical
- Three groups for categorical
- No (experimental design)
- Yes
- Yes
- Yes (three test)

d) D'Agostino-Pearson test (0.8898, 0.6164, 0.6025),
 Shapiro-Wilk test (0.9567, 0.9304, 0.9410) → Answer: Yes

e) Brown-Forsythe test (0.3412), Bartlett's test (0.2371) → Answer: Yes

● **TEST: One way ANOVA** (0.0159, *)

CTRL vs TRT1 (0.3909) → No, CTRL vs TRT2 (0.1980) → No,
 TRT1 vs TRT2 (0.0120) → Yes

ID	Dataset	Group
M_1	24.17	CTRL
M_2	25.58	CTRL
M_3	25.18	CTRL
M_4	26.11	CTRL
M_5	24.50	CTRL
M_6	24.61	CTRL
M_7	25.17	CTRL
M_8	24.53	CTRL
M_9	25.33	CTRL
M_10	25.14	CTRL
M_11	24.81	TRT1
M_12	24.17	TRT1
M_13	24.41	TRT1
M_14	23.59	TRT1
M_15	25.87	TRT1
M_16	23.83	TRT1
M_17	26.03	TRT1
M_18	24.89	TRT1
M_19	24.32	TRT1
M_20	24.69	TRT1
M_21	26.31	TRT2
M_22	25.12	TRT2
M_23	25.54	TRT2
M_24	25.50	TRT2
M_25	25.37	TRT2
M_26	25.29	TRT2
M_27	24.92	TRT2
M_28	26.15	TRT2
M_29	25.80	TRT2
M_30	25.26	TRT2



Example Three

● Suppose to test if one treatment affect the weight of the (same) mice

QUESTIONS

- Which kinds of variable?
- How many groups per variable?
- Are the samples paired?
- Are the distributions normal?
- Have the distributions same variance?
- Multiple test correction?

ANSWERS

- Two variables: one continuous, one categorical
- Two groups for categorical
- Yes (experimental design)
- No
- Not relevant
- No (one test)

ID	Dataset	
	Before	After
M_1	20.01	39.29
M_2	19.09	39.32
M_3	19.27	34.51
M_4	21.30	39.30
M_5	24.14	43.40
M_6	19.69	42.79
M_7	17.22	42.20
M_8	18.55	38.39
M_9	20.52	39.23
M_10	19.37	35.22

d) D'Agostino-Pearson test (0.0445, 0.8714), Shapiro-Wilk test (0.2768, 0.2894) → Answer: No

● TEST: Paired-sample Wilcoxon test (0.002, **) → Answer: Yes



Take Home Message

- Descriptive statistics lend inferential statistics the quantities of interest
- Inferential statistics is correlated with the concept of **error**, because a sample **approximates the population**
- Type I error (α) and type II error (β) have a reverse trend: if it is possible, **increment the sample size**
- Select the **hypothesis test** corresponding to the actual experimental design, and correct for **multiple comparisons**
- Check the assumptions for selecting a **parametric or non-parametric test**

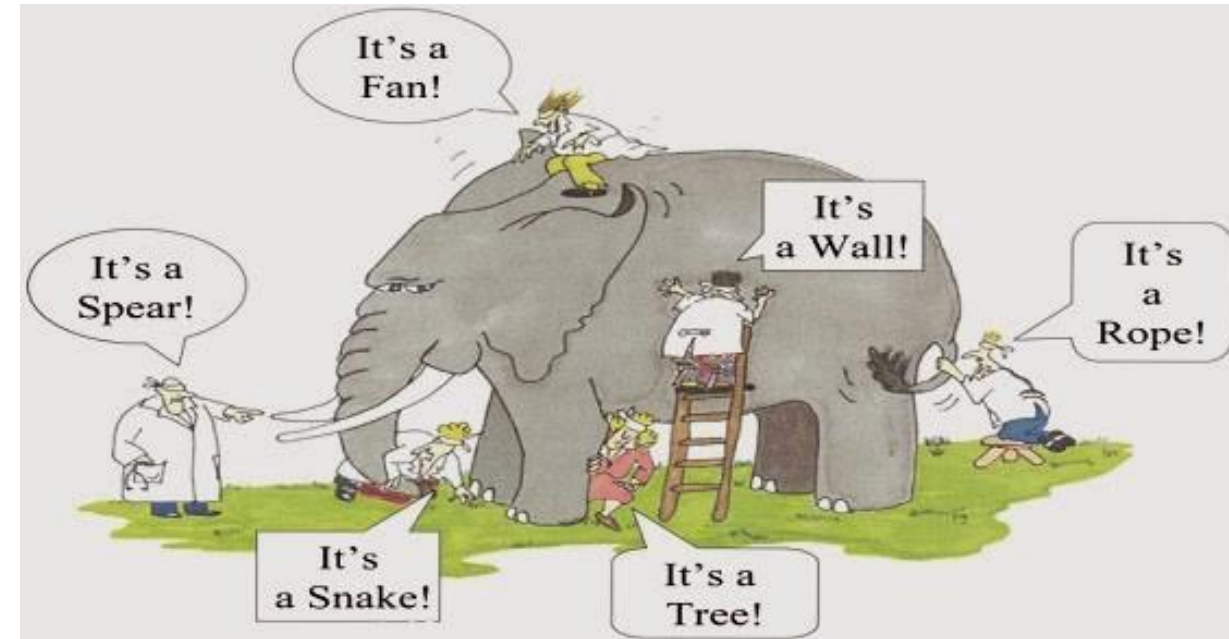


Final Remarks

To consult the statistician after an experiment is finished is often merely to ask him to conduct a postmortem examination. He can perhaps say what the experiment died of.

Sir R. A. Fisher

First Session of the Indian Statistical Conference,
Calcutta, 1938



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- [2] Banerjee, A. Hypothesis testing, type I and type II errors, *Ind. Psychiatry J.* (2009).
- [3] Greenland, S. **Statistical tests, P values, confidence intervals, and power: a guide to misinterpretations**, *Eur J Epidemiol.* (2016).

- [h1] <https://bookdown.org/jgscott/DSGI/>
- [h2] <https://statsandr.com/>
- [h3] <https://youtu.be/EvHiee7gs9Y>