



# POWER ANALYSIS

Bioinformatics Awareness Days @ TIGEM  
July 2nd, 2025



Eugenio Del Prete, M. Eng., Ph.D.  
BIOINFORMATICS CORE  
*e.delprete@tigem.it*



## Bioinformatics Core: Tasks

- **STATISTICAL DATA ANALYSIS**  
Experimental Design, Hypothesis Testing, Power Analysis 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...

### TIGEM FACILITY

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

### BIOINFORMATICS CORE

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

### BIOINFO HELPDESK

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

### STATISTICAL COMPENDIUM

[https://bioinformatics.tigem.it/materials/Statistical\\_Small\\_Compndium\\_ver1.1.pdf](https://bioinformatics.tigem.it/materials/Statistical_Small_Compndium_ver1.1.pdf)



DIEGO DI BERNARDO



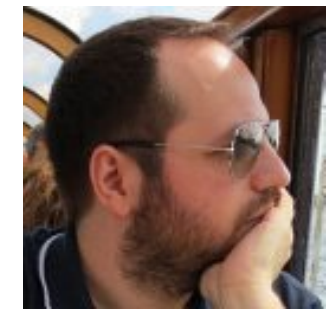
DIEGO CARRELLA



ROSSELLA DE CEGLI



XAVIER BUJANDA CUNDIN



EUGENIO DEL PRETE



## Bioinformatics Core: ...and Friends

### MASS SPECTROMETRY FACILITY



LUCIA SANTORELLI

*<https://proteomics.tigem.it/>*

### STAFF SCIENTIST



ANDREA PASQUADIBISCEGLIE



## 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 SCIENTIST @ TIGEM**



## Outline

### ● PRELIMINARIES

- PREPARE Guidelines
- ARRIVE Guidelines

### ● DEFINITIONS

- Sample Size
- Replicates
- Power
- Significance Level
- Effect Size
- Power Analysis

### ● G\*POWER

- Tool
- Graphical User Interface

### ● CONCLUSION

- Take Home Message
- Final Remarks

### ● EXAMPLES

- One Way ANOVA
- Repeated Measure ANOVA
- Multi Way ANOVA



## Guidelines: PREPARE

### ● Planning Research and Experimental Procedures on Animals: Recommendations for Excellence

- Guidelines for planning animal research and testing (2017)
- Improve the quality of the preparation for animal studies

### ● Checklist

#### a) Formulation of the study

- Literature searches
- Legal issues
- Ethical issues, harm-benefit assessment and humane endpoints
- Experimental design and statistical analysis

#### b) Dialogue between scientists and the animal facility

- Objectives and timescale, funding and division of labor
- Facility evaluation
- Education and training
- Health risks, waste disposal and decontamination

#### c) Quality control of the components in the study

- Test substances and procedures
- Experimental animals
- Quarantine and health monitoring
- Housing and husbandry
- Experimental procedures
- Human killing, release, reuse or rehoming
- Necropsy



## Guidelines: ARRIVE

### ● Animal Research: Reporting In Vivo Experiments

- Recommendations for full and transparent reporting of research involving animals (2010, 2020)
- Interconnection with the PREPARE guidelines

### ● Checklist

#### a) Essential 10

- Study design
- Sample size
- Inclusion and exclusion criteria
- Randomization
- Blinding
- Outcome measures
- Statistical methods
- Experimental animals
- Experimental procedures
- Results

#### b) Recommended Set

- Abstract
- Background
- Objectives
- Ethical statement
- Housing and husbandry
- Animal care and monitoring
- Interpretation and scientific implications
- Generalizability and translation
- Protocol registration
- Data access
- Declaration of interests



## Focus on: Pilot Studies

### ● Pre-experimental Design

- Small-scale **preliminary** experiments conducted before launching a full research study

### ● Main goals

- Test the **feasibility** of the experiment
- **Fine-tune** the experimental conditions

### ● Essential

- Checking the **clarity and effectiveness** of the experimental protocol
- Identifying and correcting technical or methodological **issues**
- Estimating data **variability**
- **Determining the appropriate sample size for future studies**
- Optimizing the use of **resources**
- Evaluating the **logistics** and **timing** of the full-scale experiment



## Sample Size

### Question

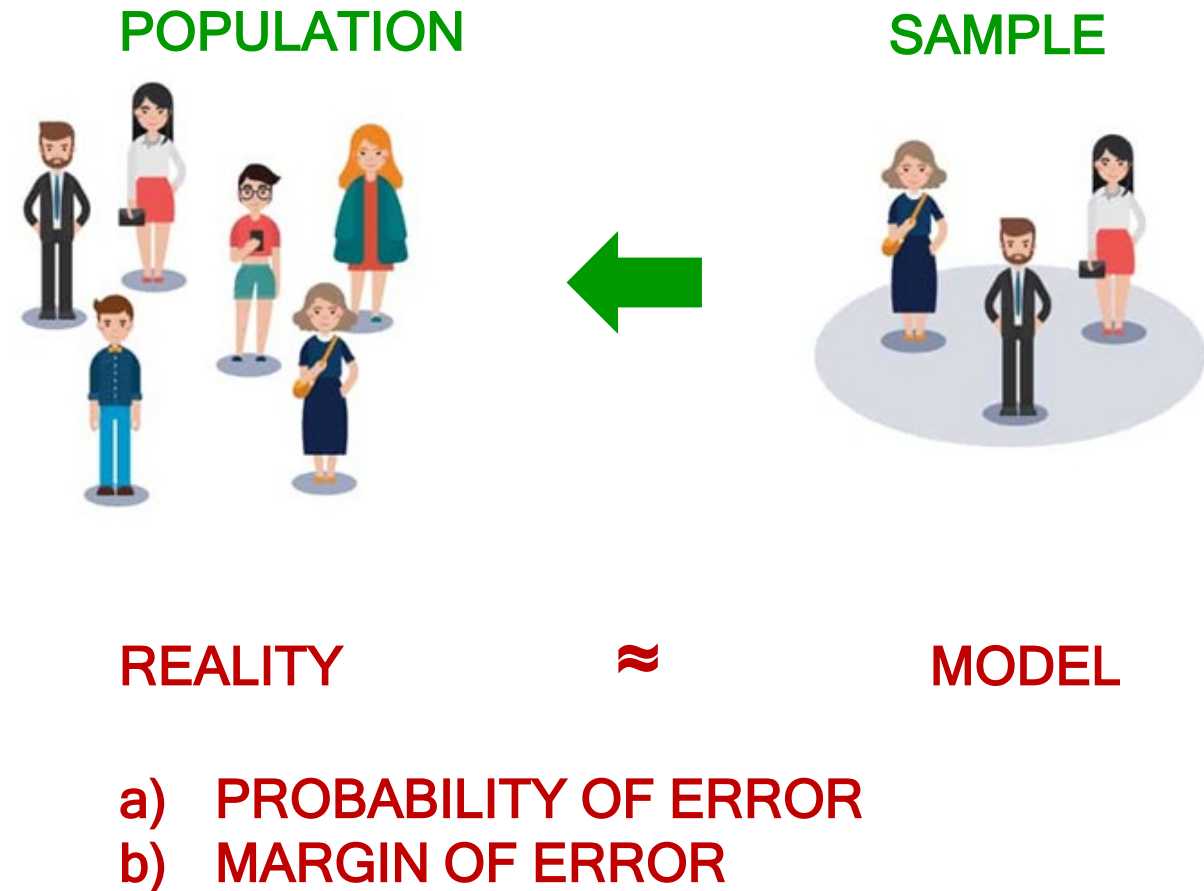
- How many samples do I need for my experiment?

### Problems

- **Impossibility** to collect data from an entire population of interest
- **Too small**: under-detection of effect of interest
- **Too large**: unnecessary wasting of resources and/or animals

### Goal

- Enough samples to **reasonably detect an effect** (if it really exists) **without wasting limited resources** on too many samples





## Replicates

### Technical Replicates

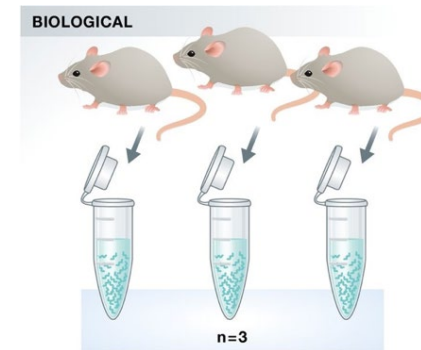
- Use the same biological sample to repeat the technical or experimental steps in order to accurately measure technical variation and remove it during analysis



WITHIN VARIABILITY

### Biological Replicates

- Use different biological samples of the same condition to measure the biological variation between samples



BETWEEN VARIABILITY

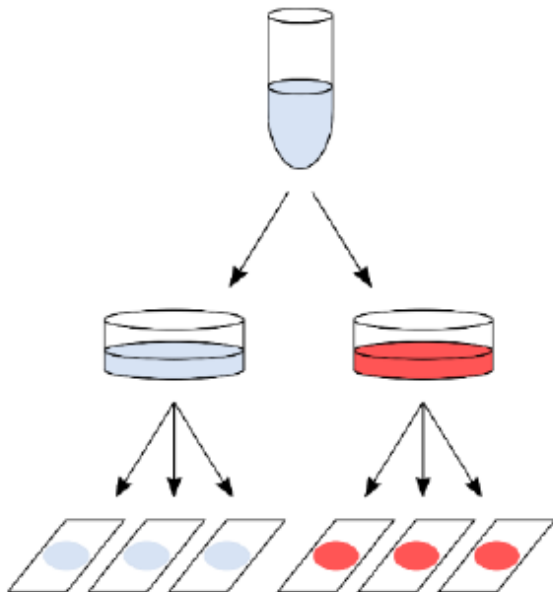
**WHY THREE (3) IS THE 'MAGIC' NUMBER?**



## Focus on: Experimental Design and Sample Size

### Design 1

- Single vial, two culture dishes
- Same time for cellular growth
- Cells counting for the glass slides
- Which is the sample size?



ANSWER: N = 1

There is not independence between the three glass slides within the two conditions

AND

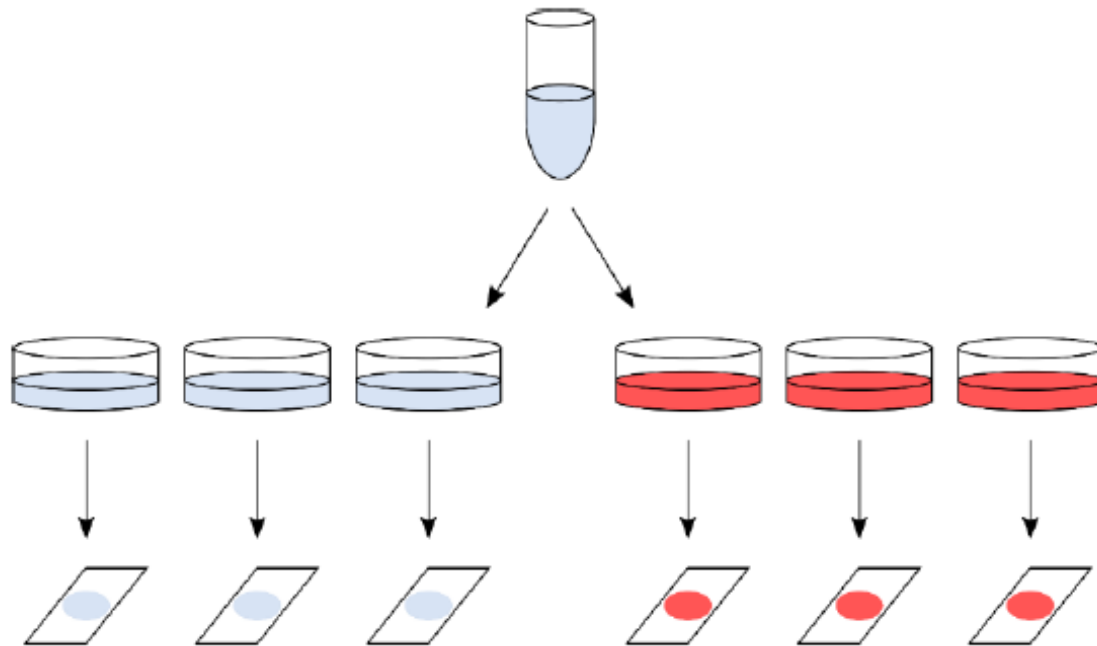
The variability among the glass slides is only due to pipetting error.



## Focus on: Experimental Design and Sample Size

### Design 2

- Single vial, six culture dishes
- Same time for cellular growth
- Cells counting for the glass slides
- Which is the sample size?



ANSWER: N = 1

The glass slides are processed in the same day, same medium, same incubator, and same time.

AND

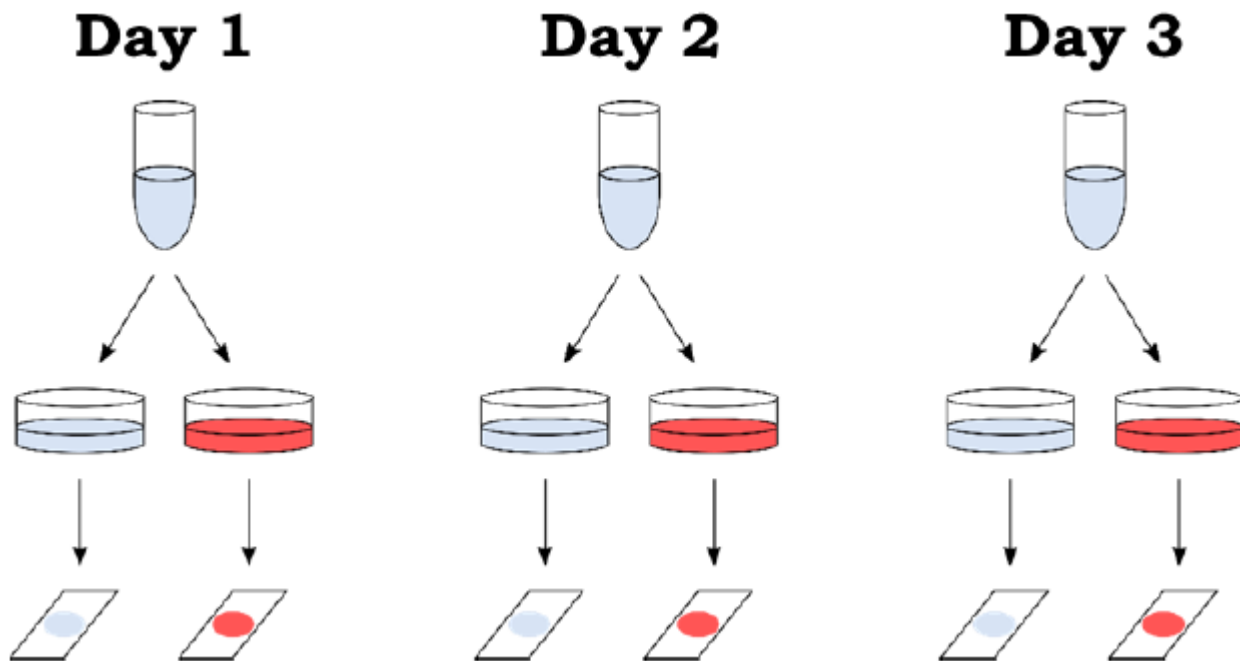
Different plastic does not provide independence.



## Focus on: Experimental Design and Sample Size

### Design 3

- Three vials, one sample, six culture dishes
- Three time for cellular growth
- Cells counting for the glass slides
- Which is the sample size?



ANSWER: N = 3

Three different times introduce the independence, even if the variability is still more technical than biological

AND

The glass slides for each time are paired



## Focus on: Experimental Design and Sample Size

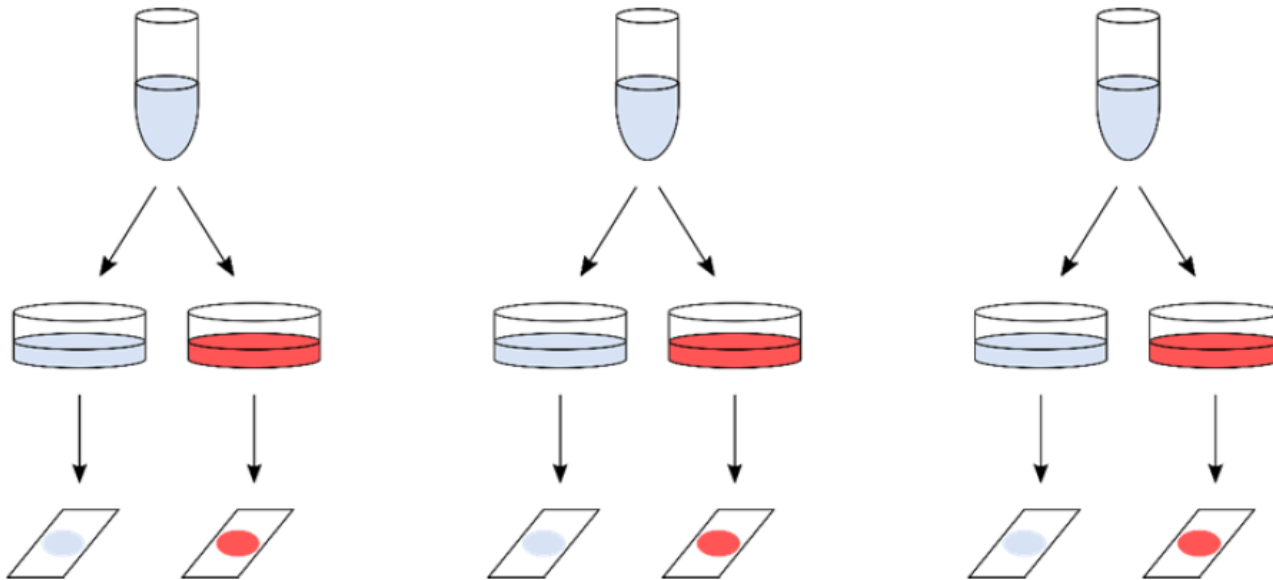
### Design 4

- Three vials, three samples, six culture dishes
- Same time for cellular growth
- Cells counting for the glass slides
- **Which is the sample size?**

person/animal/plant 1

person/animal/plant 2

person/animal/plant 3



ANSWER: N = 3

Three independent samples provides biological replicates

AND

The glass slides for each sample are paired

AND

Pay attention to batch effects (e.g., litter)



# Power and Significance Level

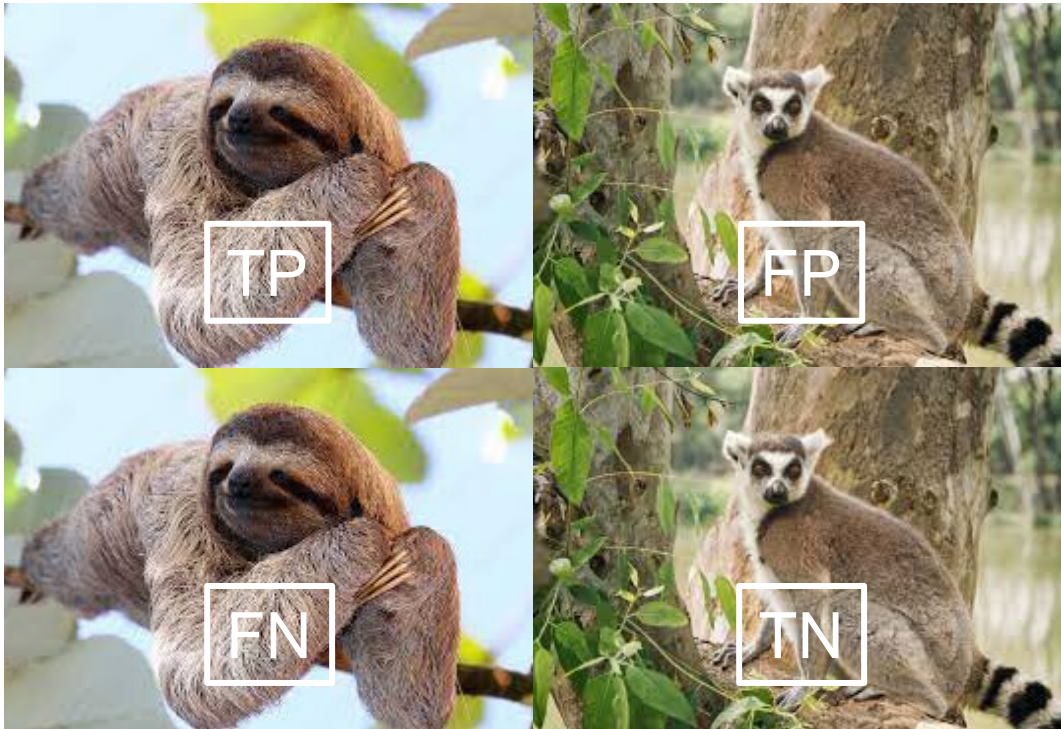
● H0: LEMUR (NOT SLOTH)

## 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



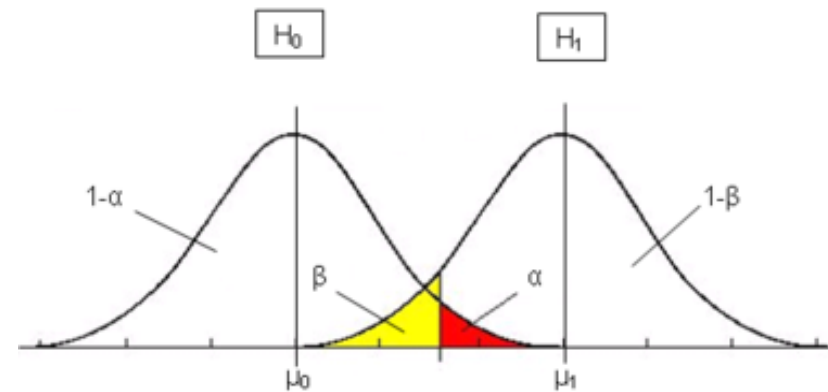
## Power and Significance Level

### ● Power ( $1-\beta$ )

- Probability to reject a **false** null hypothesis ( $H_0$ ) when the alternative hypothesis ( $H_1$ ) is **true**
- Probability of detecting a **true difference** (there is a difference to detect)
- Direct relation with the **Type II Error**

### ● Significance Level ( $\alpha$ )

- Probability to reject a **false** null hypothesis ( $H_0$ ) when the null hypothesis ( $H_0$ ) is **true**
- Probability of detecting a **false difference** (there is not a difference to detect)
- It is the **Type I Error**





# Effect Size

## Considerations

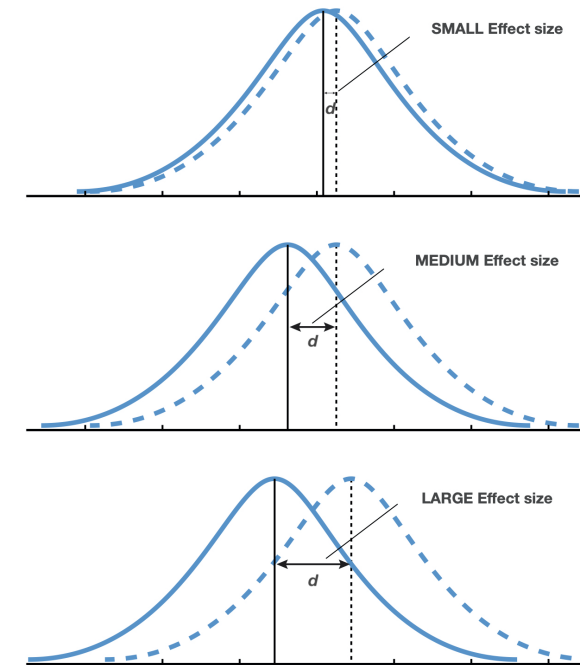
- Property of the sample data (unlike power and significance level)
- To be **estimated**
- Different **statistical tests** have **different values** for each category

## Estimation

- Background information from **preliminary data** or **trial data** or similar study
- **Convention** (forced or not)

## Formula (Generic)

$$Effect\ Size \approx \frac{Mean\ of\ H1 - Mean\ of\ H0}{Pooled\ Standard\ Deviation}$$



Test	Relevant effect size	Effect Size Threshold		
		Small	Medium	Large
t-test for means	d	0.2	0.5	0.8
F-test for ANOVA	f	0.1	0.25	0.4
t-test for correlation	r	0.1	0.3	0.5
Chi-square	w	0.1	0.3	0.5
2 proportions	h	0.2	0.5	0.8



## Power Analysis

### Hypothesis

- Biological question
- One Tail or Two Tails test

### Experimental Design

- Variables (Treatment, Timepoint, Dose, Sex, ...)

### Statistical Test

- Comparison of interest
- Type of variables

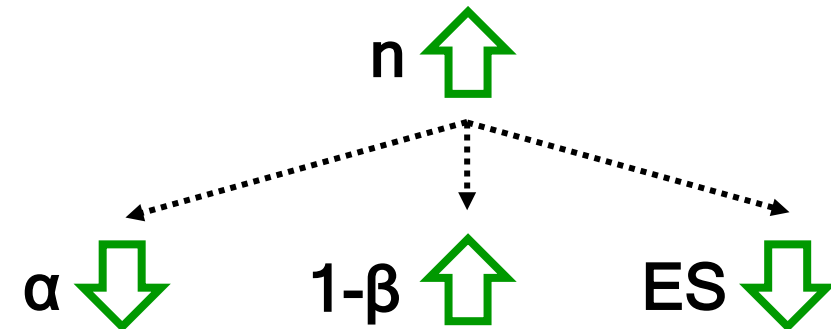
### Parameters

- Power (1-β)
- Significance Level (α)
- Effect Size (ES)
- Sample Size (n)

One Tail:  $\begin{cases} H_0: \mu_0 > \mu_1 \text{ or } H_0: \mu_0 < \mu_1 \\ H_1: \mu_0 < \mu_1 \text{ or } H_1: \mu_0 > \mu_1 \end{cases}$

Two Tails:  $\begin{cases} H_0: \mu_0 = \mu_1 \\ H_1: \mu_0 \neq \mu_1 \end{cases}$

**WHICH IS THE LINK WITH n  
AMONG THE PARAMETERS?**





## Focus on: Type of Power Analysis

### ● A Priori Power Analysis

- Determine how many samples you will need to have a good chance of detecting an effect of a specified size with the desired amount of power



**GIVEN  $\alpha$ ,  $1-\beta$ , ES  
THEN COMPUTE  $n$**

A priori analyses are performed as **part of the research planning process**. They allow you to determine the sample size you need in order to reach a desired level of power.

### ● A Posteriori Power Analysis

- Allow to find how much power you would have if you had a specified and sufficient number of samples when you had already conducted a research



**GIVEN  $\alpha$ ,  $n$ , ES  
THEN COMPUTE  $1-\beta$**

Post hoc analyses are performed after your study has been conducted and can be used to assist in **explaining any potential non-significant results**.



# G\*Power

## Availability

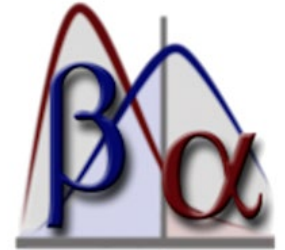
- **WINDOWS:** 3.1.9.7 (March 2020)
- **MAC:** 3.1.9.6 (February 2020)

## Statement

- **Free** software for everyone
- **Downloadable** from the webpage
- **Screenshot** without asking permissions

## Analysis Type

- Selection for the **parameter of interest**
- Selection for the **test of interest**



Analysis Type	Calculated Value
A priori	Required sample size
Compromise	Implied power and alpha values
Criterion	Required alpha value
Post hoc	Achieved power
Sensitivity	Required effect size

**EXACT: 8 specific + 1 general**  
**F-TEST: 16 specific + 1 general**  
**T TEST: 11 specific + 1 general**  
**X<sup>2</sup> TEST: 2 specific + 1 general**  
**Z TEST: 7 specific + 1 general**



# G\*Power GUI

The screenshot shows the G\*Power 3.1 interface. Callout 1 points to the 'Protocol of power analyses' tab, which displays a graph of central and noncentral distributions with a critical F value of 2.4233. Callout 2 points to the 'Input parameters' section, which includes fields for Effect size f (0,3261901),  $\alpha$  err prob (0,05), Power (1- $\beta$  err prob) (0,95), and Number of groups (5). The 'Output parameters' section shows Noncentrality parameter  $\lambda$  (19,1519966), Critical F (2,4232862), Numerator df (4), Denominator df (175), Total sample size (180), and Actual power (0,9507614). A table on the right shows group means and sizes.

Group	Mean	Size
1	10	50
2	13	50
3	14	50
4	15	50
5	20	50

a) Type of power analysis

b) Test family

c) Statistical test

d) Input parameters (with values)

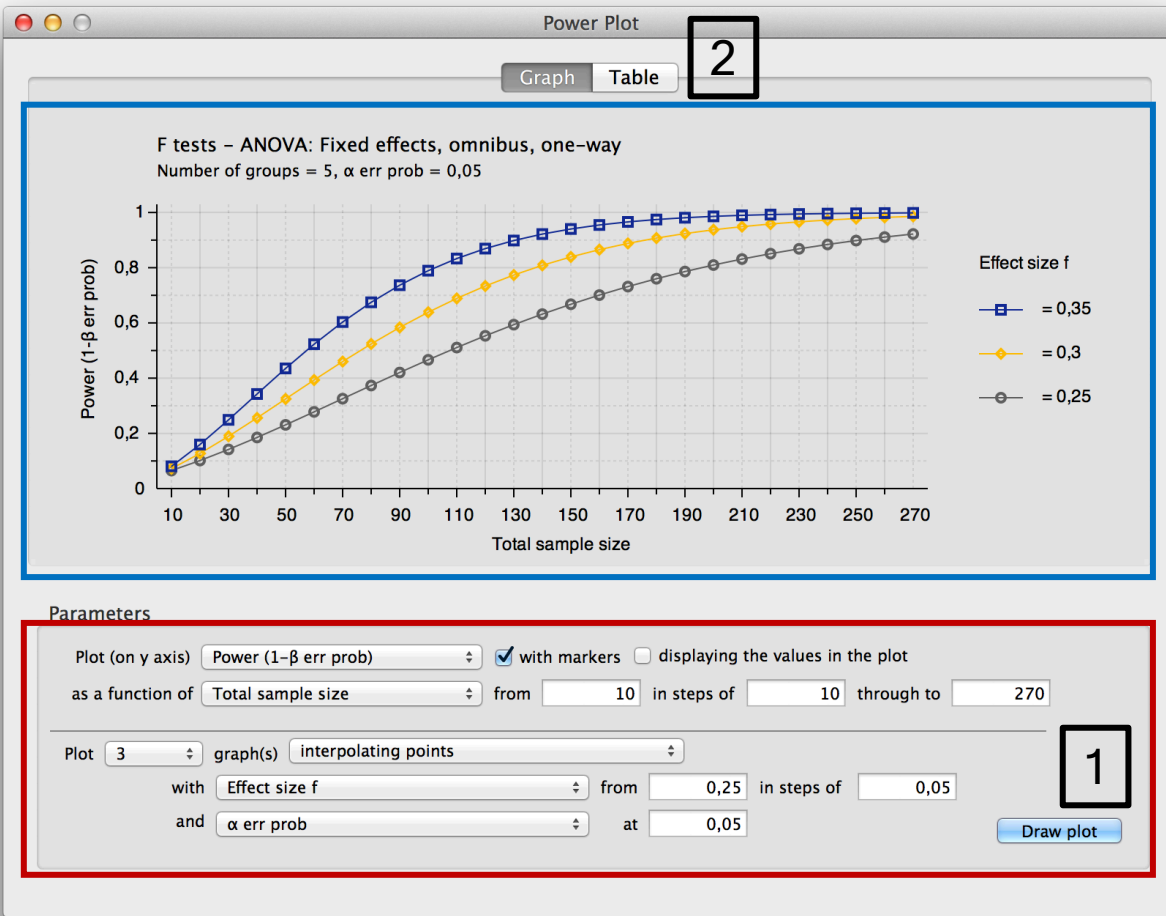
e) Output parameters with density plots

If you click on:

- 1) Protocol of power analysis: Table with I/O parameters
- 2) X-Y plot for a range of values: Plot for parameters



# G\*Power GUI



a) Parameters and attributes

b) Power Plot

If you click on:

1) Draw plot: Power Plot

2) Table: X/Y values in tabular format



# One Way ANOVA

## Description

- Check if **at least one mean** is different among groups
- **Normally distributed (numeric) variables**
- Extension of unpaired t-test for more than two groups

## Conditions

- **Numeric variables:** one **MEASUREMENTS**
- **Categorical variables:** one **E.G. TREATMENT, TIMEPOINT, DOSE**
- **Number of group per categorical variable:** more than two
- **Details:** parametric, unpaired **NORMAL DISTRIBUTION AND DIFFERENT SAMPLES**

## Parameters

- Significance level: 0.05
- Power: 0.80



## One Way ANOVA

### ● Example

- Five different medication regimes on patients after surgery
- Differences in white cell counts
- $H_0$ : no difference among medication regimes;  $H_1$ : difference among medication regimes

### ● What we know

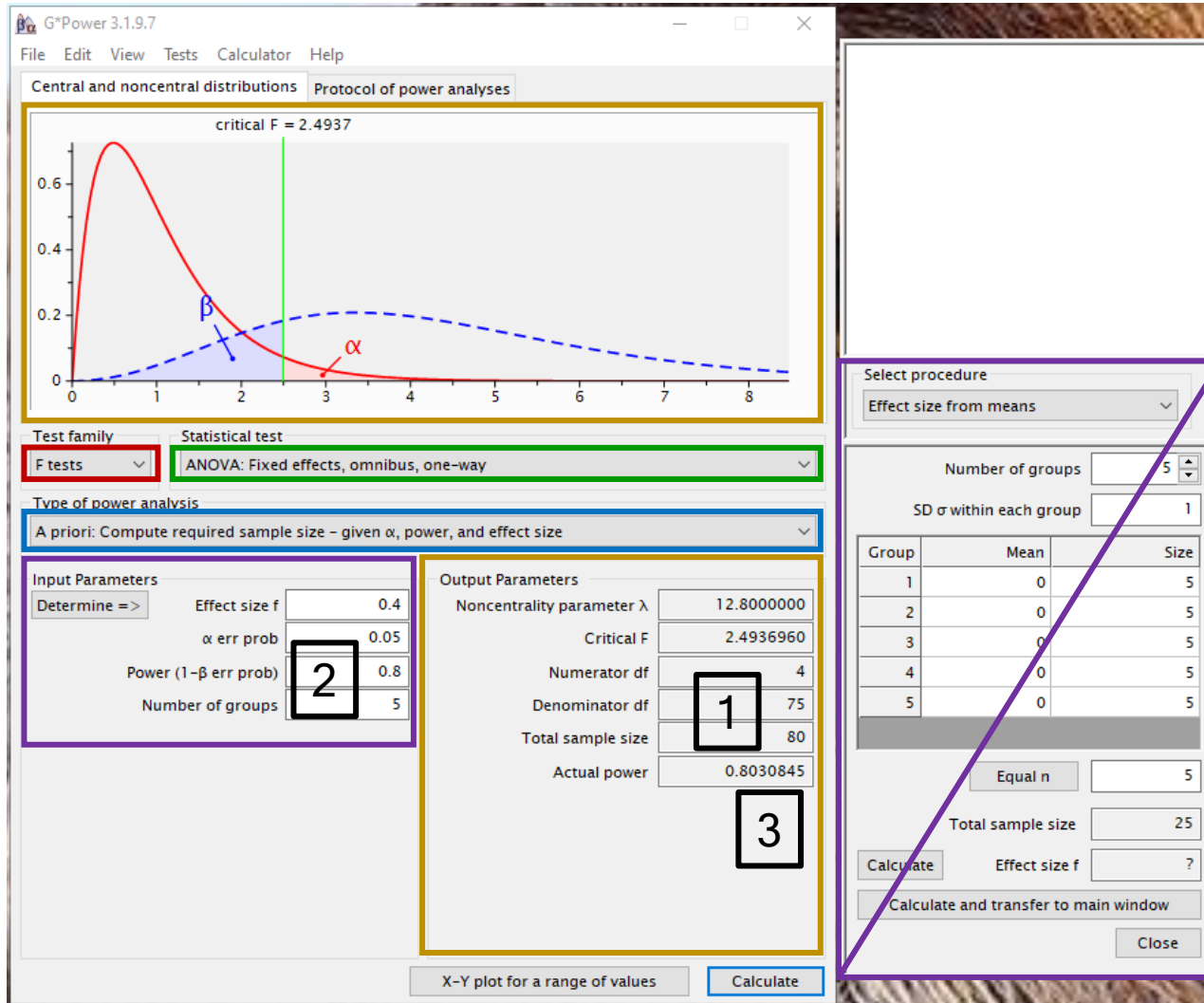
- Number of groups: five
- Number of measurements: one per sample
- Significance level: 0.05
- Power: 0.80

### ● What we guess

- Effect Size  $f$ : 0.4 (large)



# One Way ANOVA



**Input Parameters:**

- Determine => Effect size f: 0.4
- α err prob: 0.05
- Power (1 - β err prob): 0.8
- Number of groups: 5

**Output Parameters:**

- Noncentrality parameter λ: 12.8000000
- Critical F: 2.4936960
- Numerator df: 4
- Denominator df: 75
- Total sample size: 80
- Actual power: 0.8030845

Group	Mean	Size
1	0	5
2	0	5
3	0	5
4	0	5
5	0	5

a) A priori: compute required sample size

b) F tests

c) ANOVA: Fixed effects, omnibus, one-way

d) Input parameters (without values)

e) Output parameters with density plots

RESULTS:

- 1) Total sample size ( $T_{ss}$ ): 80
- 2) Sample size per group:  $T_{ss} / n_g = 16$
- 3) Actual power: 0.803 (check)



## Repeated Measure ANOVA

### Description

- Check if **at least one mean** is different among groups
- **Normally distributed (numeric) variables**
- Extension of paired t-test for more than two groups

### Conditions

- **Numeric variables:** one **MEASUREMENTS**
- **Categorical variables:** one **E.G. TREATMENT, TIMEPOINT, DOSE**
- **Number of group per categorical variable:** more than two
- **Details:** parametric, paired **NORMAL DISTRIBUTION AND SAME SAMPLES**

### Parameters

- Significance level: 0.05
- Power: 0.80



## Repeated Measure ANOVA

### ● Example

- One treatment on patients
- Four measurements of blood pressure after treatment (1, 2, 3, 4 months)
- H0: no difference during time; H1: difference during time

### ● What we know

- Number of groups: one
- Number of measurements: four per sample
- Significance level: 0.05
- Power: 0.80

### ● What we guess

- Correlation: 0.5
- Non-sphericity correction: 1
- Partial  $\eta^2$ : 0.14 (large)

**CORRELATION:** relationships among measurements  
**SPHERICITY:** same differences of variances  
**PARTIAL  $\eta^2$ :** ratio between variances (EFFECT SIZE)



# Repeated Measure ANOVA

a) A priori: compute required sample size

b) F tests

c) ANOVA: Repeated measures, within factors

d) Input parameters (with Partial  $\eta^2$ )

e) Output parameters with density plots

RESULTS:

- 1) Total sample size: 10
- 2) Actual power: 0.814 (check)



## Multi Way ANOVA

### Description

- Define the number of **categories of interest** (one or more than one)
- **Normally distributed (numeric) variables**
- Extension of one way ANOVA for more than one category

### Conditions

- **Numeric variables:** one **MEASUREMENTS**
- **Categorical variables:** more than one **E.G. TREATMENT, TIMEPOINT, DOSE**
- **Number of group per categorical variable:** more than one
- **Number of categories of interest:** one or more than one
- **Details:** parametric, unpaired **NORMAL DISTRIBUTION AND DIFFERENT SAMPLES**

### Parameters

- Significance level: 0.05
- Power: 0.80



## Multi Way ANOVA

### ● Example

- Different treatments on patients (a, b, c)
- Different ages (child, adult, elder) and different cancer stages (I, II, III, IV, V)
- $H_0$ : no difference in treatment across ages/stages;  
 $H_1$ : difference in treatment across ages/stages

### ● What we know

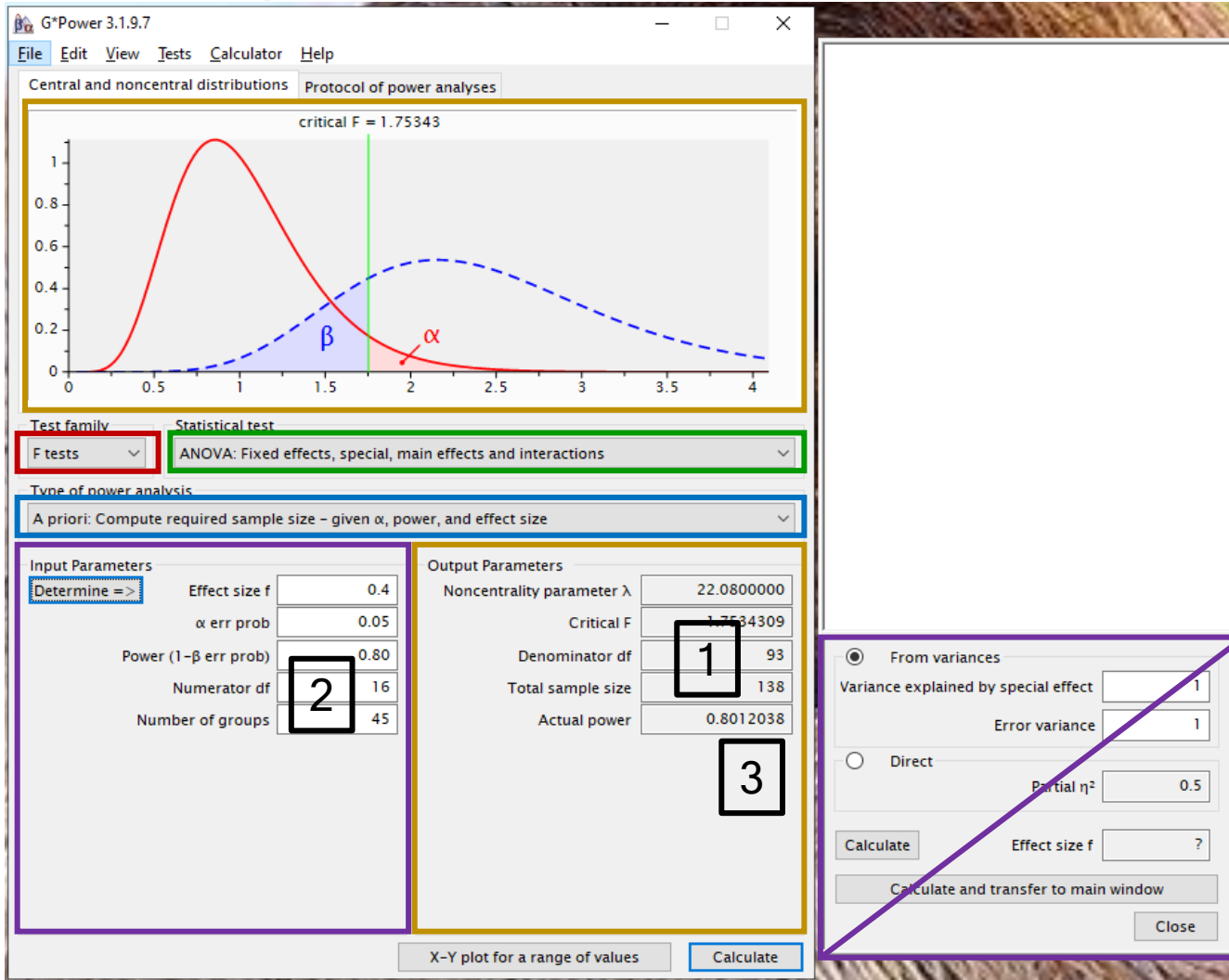
- **Number of groups:** three treatments ( $T$ ), three ages ( $A$ ), five stages ( $S$ ),  $n_g = TAS = 45$
- **Number of degrees of freedom:**  $df = (T - 1)(A - 1)(S - 1) = 16$
- **Significance level:** 0.05
- **Power:** 0.80

### ● What we guess

- **Effect Size f:** 0.4 (large)



# Multi Way ANOVA



The screenshot shows the G\*Power 3.1.9.7 interface. At the top, a graph displays two density curves: a solid red curve for the null distribution and a dashed blue curve for the alternative distribution. A vertical green line marks the critical F value at 1.75343. The area under the red curve to the right of this line is shaded red and labeled  $\alpha$ . The area under the blue curve to the left of this line is shaded blue and labeled  $\beta$ .

Below the graph, the 'Test family' is set to 'F tests' and the 'Statistical test' is 'ANOVA: Fixed effects, special, main effects and interactions'. The 'Type of power analysis' is 'A priori: Compute required sample size - given  $\alpha$ , power, and effect size'.

The 'Input Parameters' section includes:

- Effect size f: 0.4
- $\alpha$  err prob: 0.05
- Power (1- $\beta$  err prob): 0.80
- Numerator df: 16
- Number of groups: 45

The 'Output Parameters' section includes:

- Noncentrality parameter  $\lambda$ : 22.0800000
- Critical F: 1.7534309
- Denominator df: 93
- Total sample size: 138
- Actual power: 0.8012038

At the bottom right, the 'From variances' section is visible, with 'Variance explained by special effect' set to 1 and 'Error variance' set to 1. The 'Direct' section is also visible with 'Partial  $\eta^2$ ' set to 0.5. A purple box highlights the 'From variances' section, and a diagonal line is drawn across it.

a) A priori: compute required sample size

b) F tests

c) ANOVA: Fixed effects, special, main effects and interactions

d) Input parameters (without values)

e) Output parameters with density plots

RESULTS:

- 1) Total sample size ( $T_{ss}$ ): 138
- 2) Sample size per group:  $T_{ss}/n_g \approx 4$
- 3) Actual power: 0.801 (check)



## Focus on: Non-Parametric Tests

- **Not normally distributed (numeric) variables**
  - Peculiar distribution and/or low number of samples
  - Information obtained from a preliminary/previous/similar experiment
  - No relaxation of the hypothesis
- **Rule of Thumb**
  - Compute the **parametric test** and add **15%** to total sample size
  - **Formula:**  $TSS_{NP} = TSS + 0.15TSS = 1.15TSS$
- **Previous Examples**
  - **One Way ANOVA:**  $1.15 \times 80 = 92$
  - **Repeated Measure ANOVA:**  $1.15 \times 10 \approx 12$
  - **Multi Way ANOVA:**  $1.15 \times 138 \approx 159$

**GENERALIZED LINEAR MIXED MODELS**



## Take Home Message

- **Sample size** is the most important parameter to detect a possible effect among different conditions
- **Sample size, significance level, power, and effect size** are strictly connected each other (considering the distribution of variables)
- Different **types** of a priori power analysis exist, based on the **parameter of interest** and on the **statistical test** to perform
- **G\*Power** is a free, downloadable, and user-friendly tools to perform a priori and a posteriori power analysis
- **Guidelines** for planning, testing and reporting experiments are available online

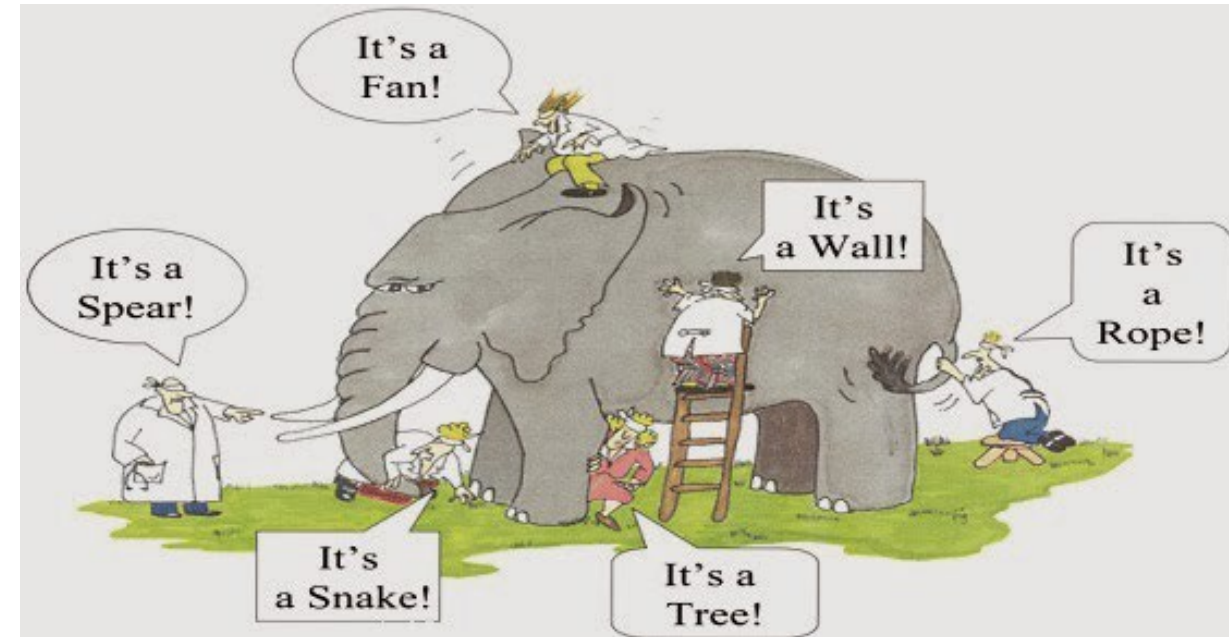


## Final Remarks

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



Eugenio Del Prete, M.Eng., Ph.D.  
Biostatistician and Data Scientist  
Telethon Institute of Genetics and Medicine (TIGEM)  
Pozzuoli (NA), Italy  
e-mail: e.delprete@tigem.it

## References

- [1] Smith, A.J. **PREPARE: guidelines for planning animal research and testing**, Laboratory Animals 2018 (2017).
- [2] du Sert, N.P. **Reporting animal research: Explanation and elaboration for the ARRIVE guidelines 2.0**, PLoS Biology (2020)
- [3] Faul, F. **Statistical power analyses using G\*Power 3.1: Tests for correlation and regression analyses**, Behavior Research Methods (2009).
- [4] Uttley, J. **Power Analysis, Sample Size, and Assessment of Statistical Assumptions - Improving the Evidential Value of Lighting Research**, LEUKOS (2019).
- [5] Kang, H. **Sample size determination and power analysis using G\*Power software**, Journal of Educational Evaluation of Health Professions (2021).
- [h1] *<https://norecopa.no/PREPARE>*
- [h2] *<https://arriveguidelines.org/>*
- [h3] *<https://www.bioinformatics.babraham.ac.uk/training/>*
- [h4] *<https://med.und.edu/research/daccota/berdc-resources.html>*
- [h5] *<https://www.psychologie.hhu.de/arbeitsgruppen/allgemeine-psychologie-und-arbeitspsychologie/gpower>*