

Hypothesis Testing . Quantitative Variables

Curs d'Estadística Bàsica per a la Recerca Biomèdica

UEB – VHIR

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Syllabus

1. INTRODUCTION

2. TYPE OF TEST

3. NORMALITY TEST

4. TWO GROUPS COMPARISON IN
INDEPENDENT SAMPLES

5. TWO GROUPS COMPARISON IN DEPENDENT
SAMPLES

6. MORE THAN TWO GROUPS COMPARISON IN
INDEPENDENT SAMPLES

7. MORE THAN TWO GROUPS COMPARISON IN
DEPENDENT SAMPLES

Example Data

- A study was designed to compare two distinct hypertension control programs.
- 60 individuals with HTA were randomly assigned to either one or the other group (30 per group)
- Blood pressure was measured each month during a year

A	B	C	D	E	F	G	H	I
numero	sexo	grupo	tas1	tad1	tas2	tad2	tas3	tad3
1	VARON	B	150	100	150	90	170	
2	MUJER	B	160	90	170	90	160	
3	MUJER	B	150	90	110	90	115	
4	VARON	A	120	80	140	90	140	
5	MUJER	A	150	85	145	85	160	
6	MUJER	B	140	75	160	70	135	
7	MUJER	A	150	100	140	90	130	
8	VARON	A	160	90	170	90	170	
9	MUJER	A	145	105	170	95	140	
10	MUJER	A	210	110				
11	MUJER	A	170	100	170	90	170	
12	MUJER	B	140	90	140	90	100	

Questions to solve

- Are samples “comparable” at baseline?
- Has there been a change in BP between month 1 (first measure) and month 12?

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Type of Hypothesis

Confirmation Hypothesis

Aim is to confirm hypothesis about parameters or distributions.

Goodness of fit test to verify hypothesis about the distribution of variable in population

Does arterial pressure in the population follow a normal distribution?

Test to verify values about a parameter.

Is the average "bua" value in our population equal to 70?

Is the proportion of lung cancer cases equal to 2.6%?

Type of Hypothesis

Independence Hypothesis

- Aim is to test hypothesis about
 - relation among variables in a population
 - Is CD4 lymphocytes count related with CD8 count in HIV positive?
 - differences in a variable among two or more populations.
 - Is the average "bua" value the same in menopausal and in non menopausal population?
 - Is the proportion of lung cancer cases the same in people with high or low fruit consumption?

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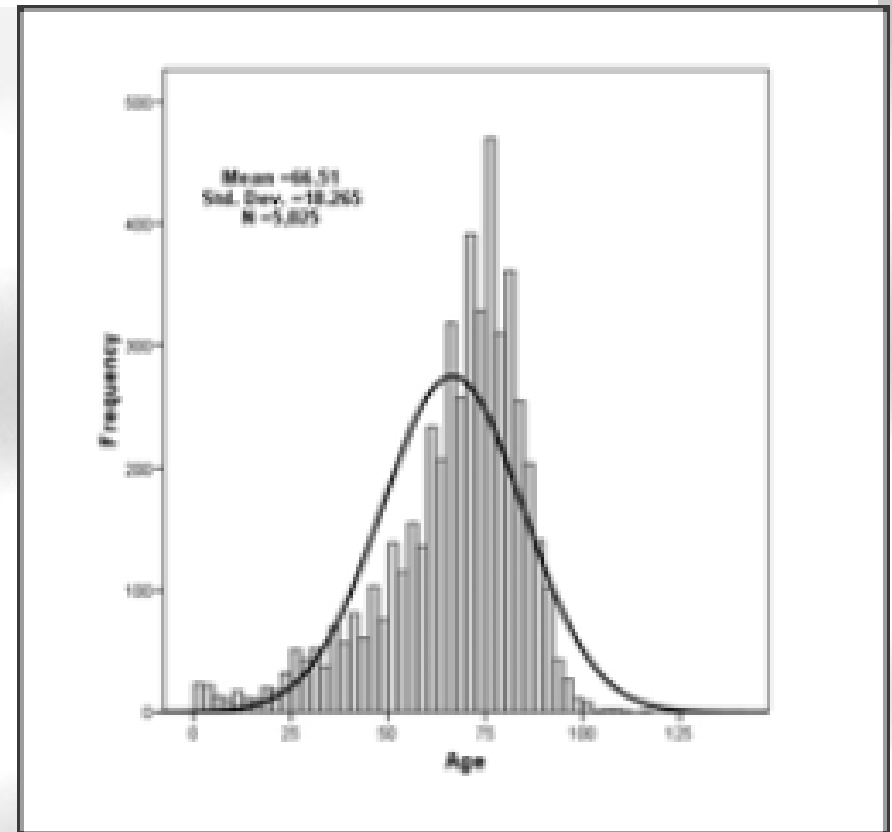
5. **TWO GROUPS** COMPARISON IN DEPENDENT
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Normality test

- Some parametrical test assume data come from a normal population
- How can we check this assumption?
- What can we do if assumption is false?

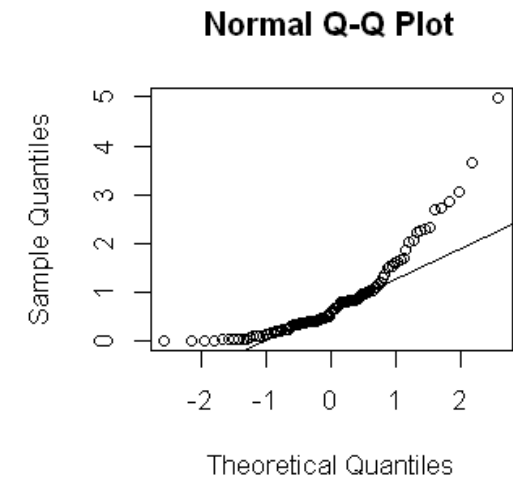
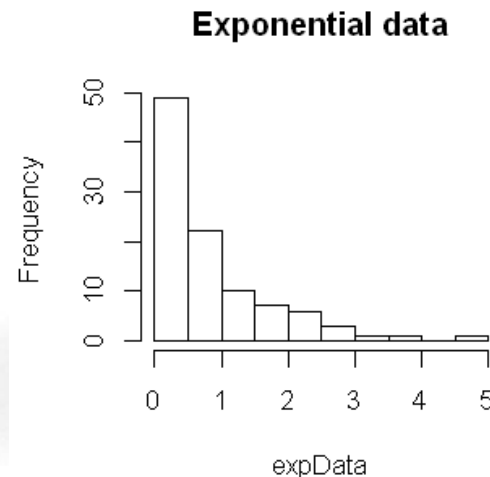
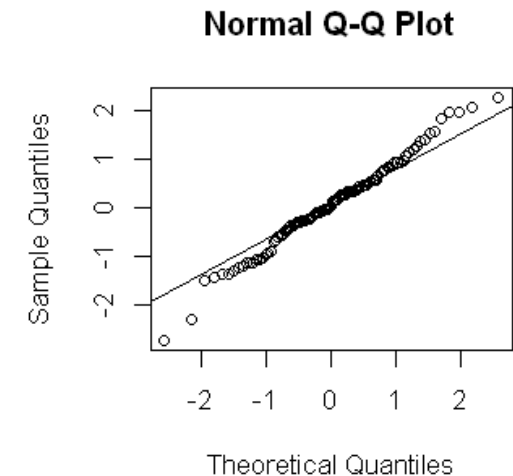
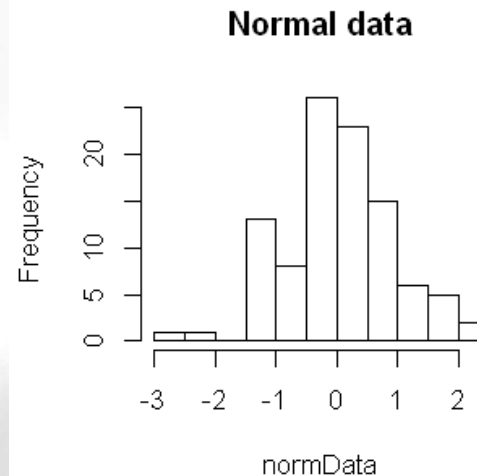


Testing normality

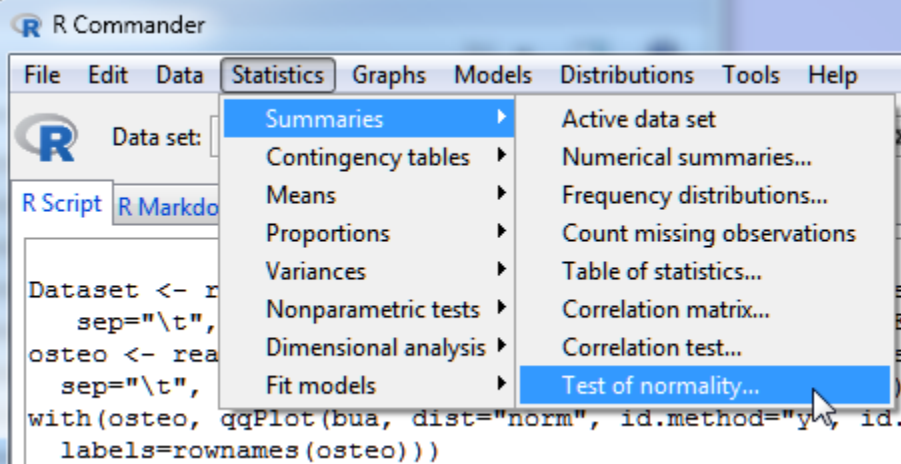
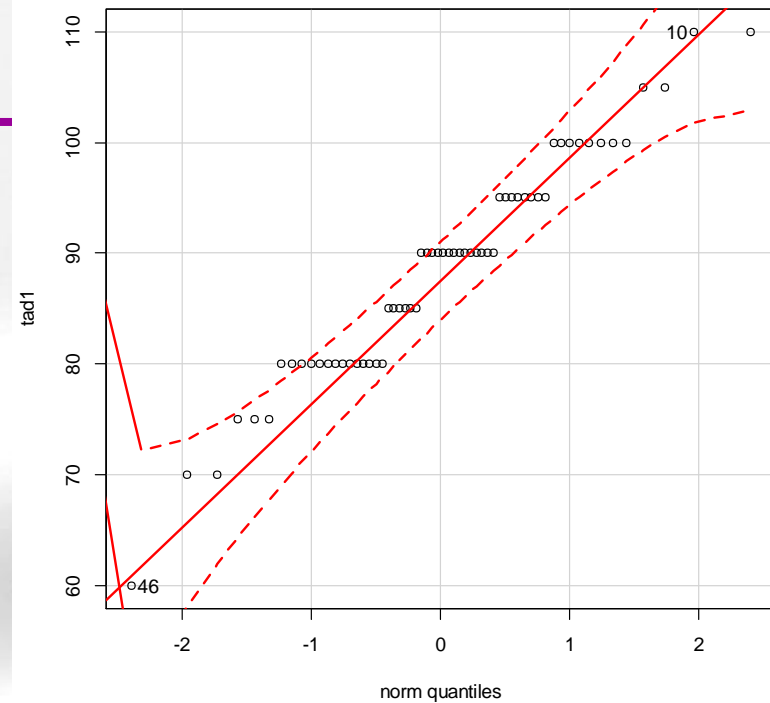
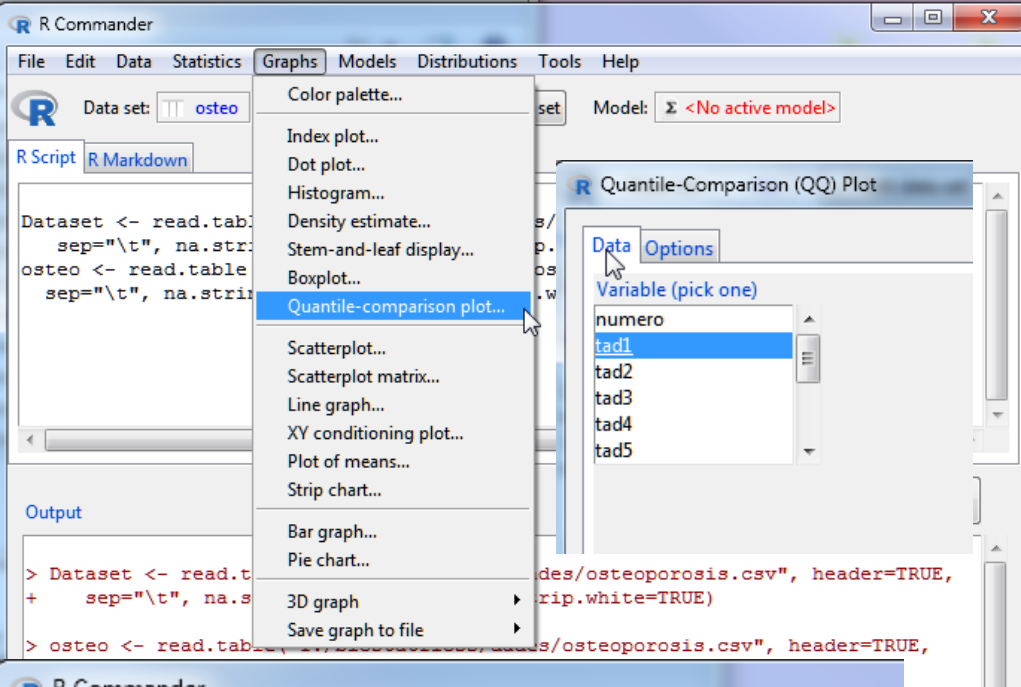
- We can use graphical methods or hypothesis tests
- Graphs
 - Check if it is a symmetric distribution
 - Probability graphs (QQ-plots)
- Hypothesis test (Normality)
 - Kolmogorov-Smirnov test
 - Kolmogorov-Liliefors test
 - Shapiro-Wilks test

Histograms and QQ-plots

- Histogram
 - It should be symmetric with gaussian shape.
- QQ-plot
 - Dots should be over the diagonal line
- Non normal data deviate from normal patterns.
- Difficult to quantify if there are few data



- Statistical normality test are more precise than graphs. It is possible to calculate a p-value.
- The most used tests are Kolmogorov-Smirnov and Shapiro-Wilks test.
- The hypothesis to test are:
 - H_0 : Data follow a normal distribution
 - H_1 : Data do not follow a normal distribution



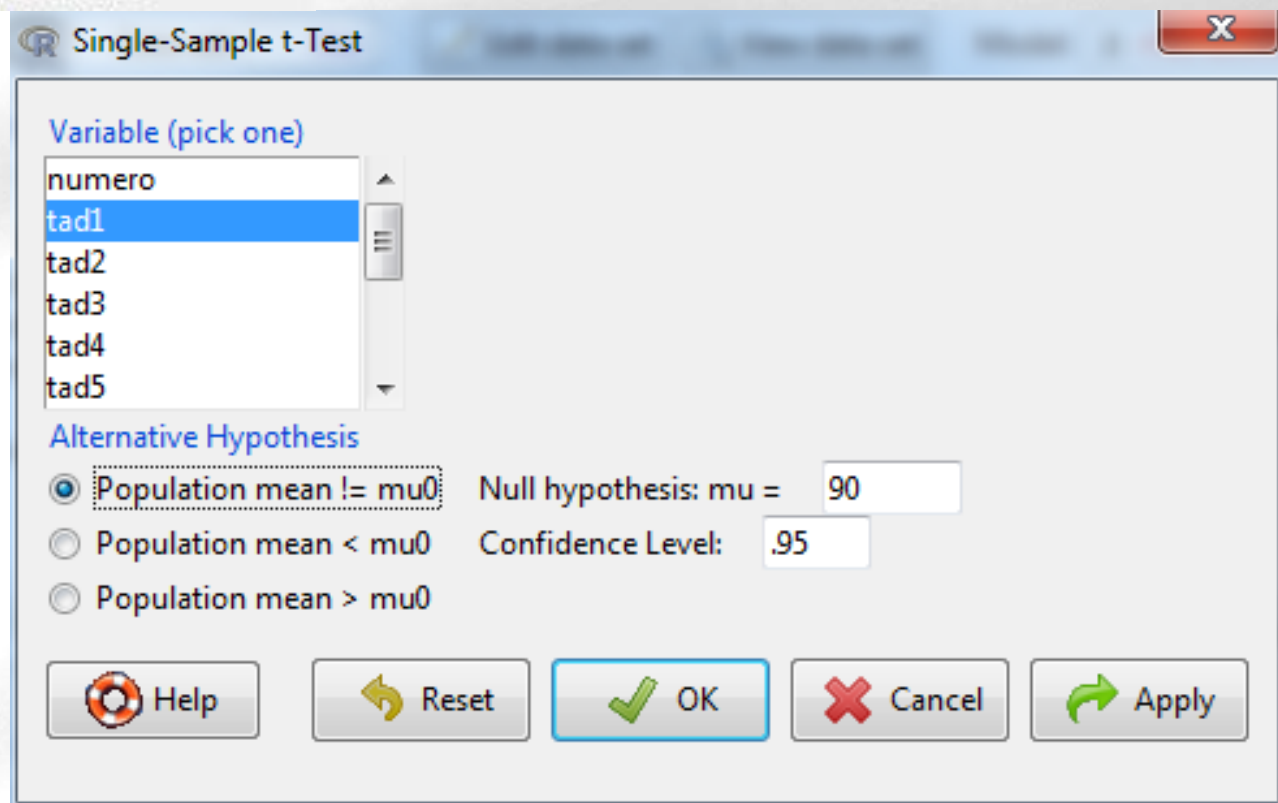
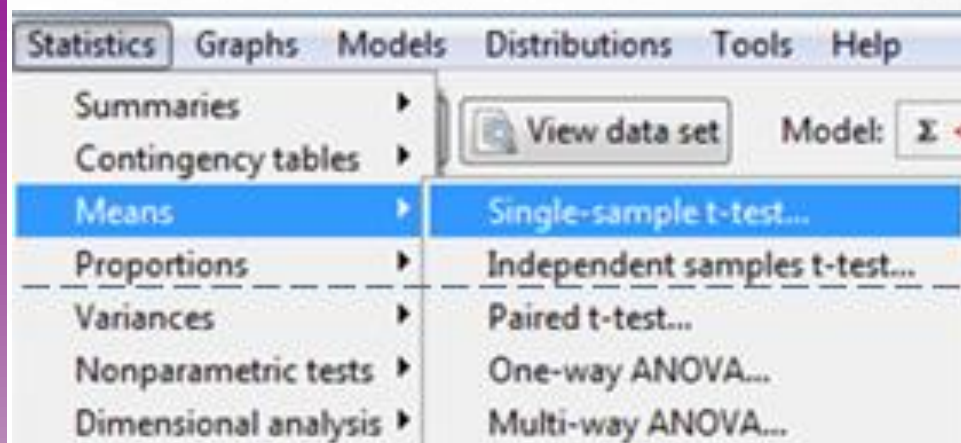
Shapiro-Wilk normality test

data: tad1

W = 0.96622, p-value = 0.09512

One sample t-test

- We do not use it very often.
- Very similar to estimation questions. It can be solved calculating a confidence interval
- Idea: We want to verify from a sample a previous hypothesis about the mean in a population
- *Can it be accepted that the initial TAD is 90 in Hypertensive patients?*



One Sample t-test

```
data:  tad1
t = -1.2137, df = 59, p-value = 0.2297
alternative hypothesis: true mean is not equal to 90
95 percent confidence interval:
 85.80626 91.02707
sample estimates:
mean of x
 88.41667
```


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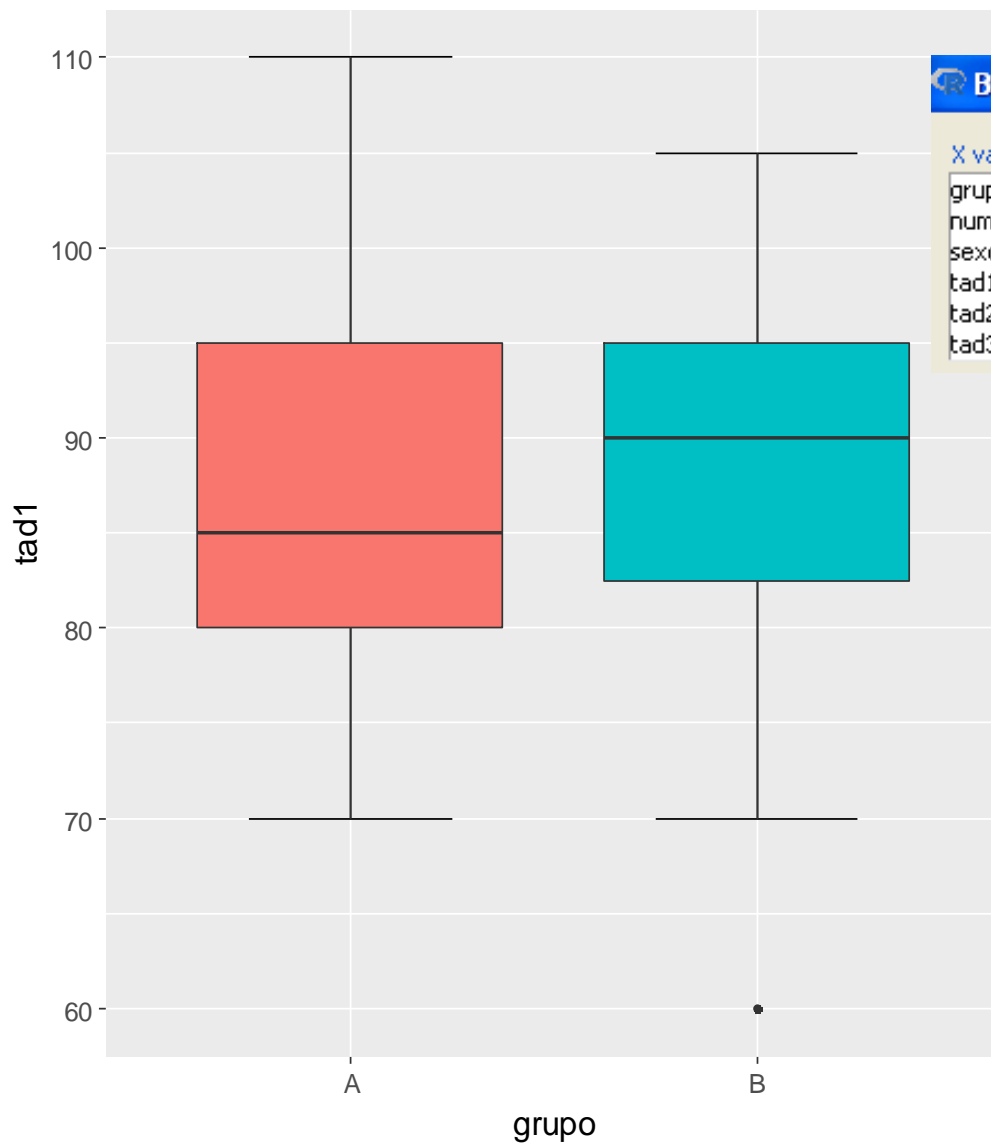
6. MORE THAN TWO GROUPS COMPARISON IN
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Questions to answer

- Are samples comparable at baseline time
- Is blood pressure comparable between first and 12th measures

Boxplot TAD, by group



Box plot / Violin plot / Confidence interval

X variable	Y variable (pick one)	Stratum variable
grupo	numero	grupo
numero	tad1	sexo
sexo	tad2	
tad1	tad3	
tad2	tad4	
tad3	tad5	

grupo

A

B

Compare a Quantitative variable in two groups

Null Hypothesis: There is not difference of the variable in two population or groups

Samples have been generated

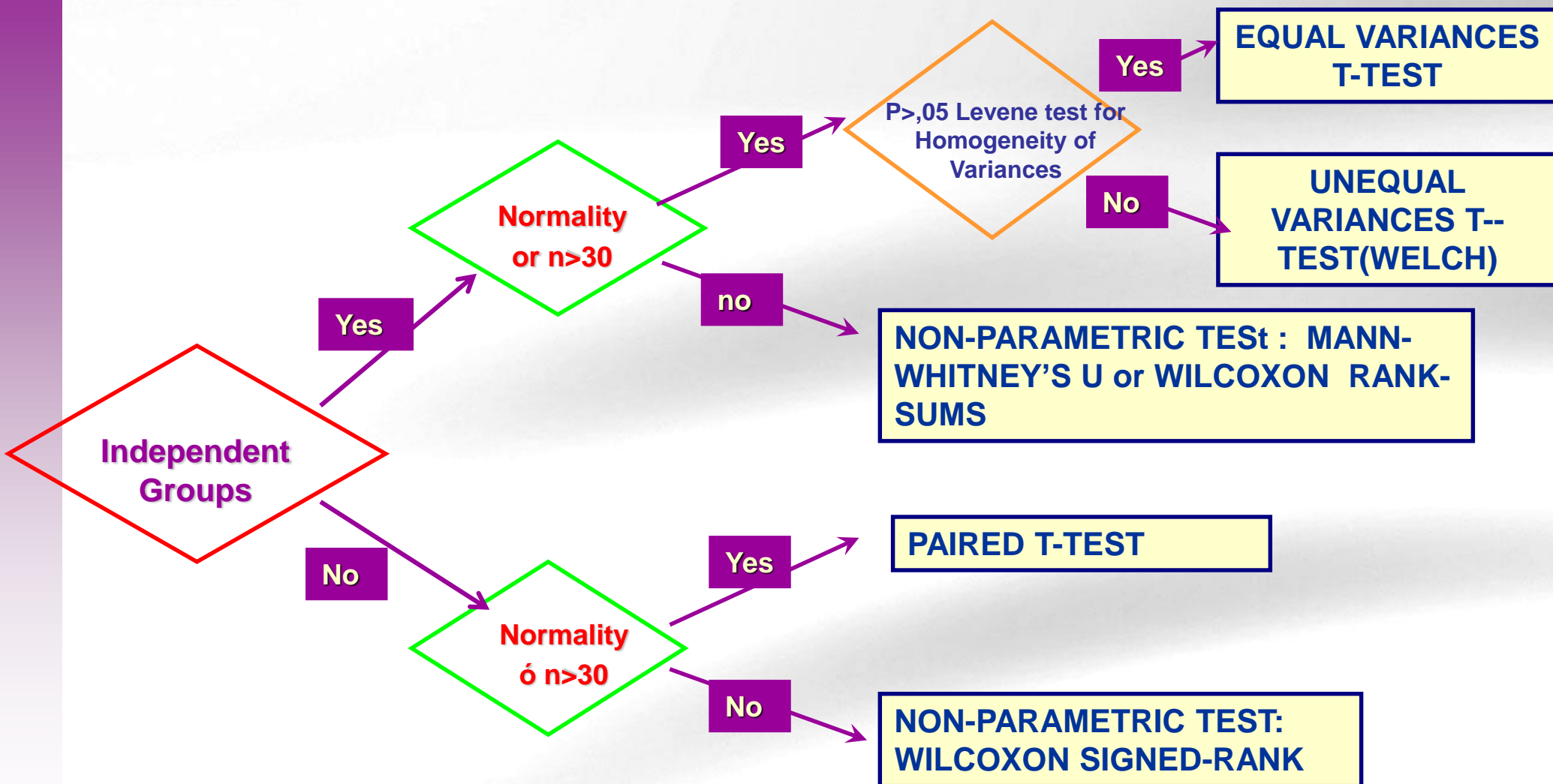
INDEPENDENT

Selected individuals in a group have nothing to do with selected individuals in the other group.

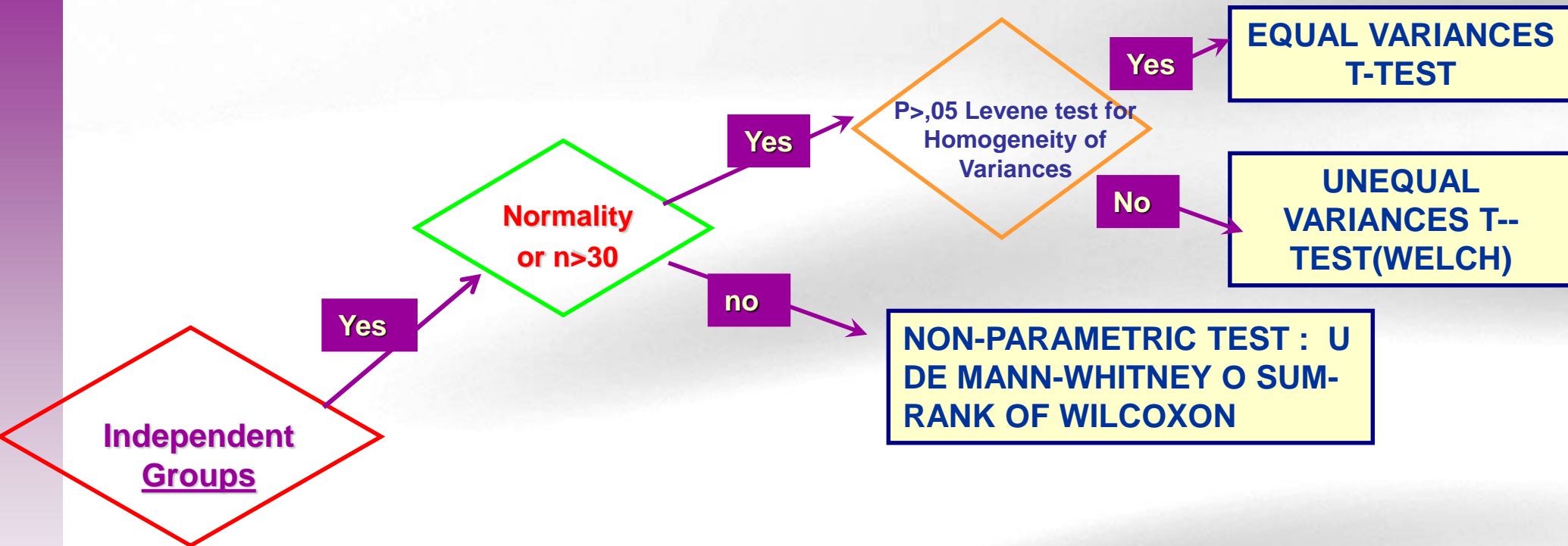
DEPENDENT

Each individual in a group has a correspondent in other group. These are ***paired data***.

Two sample tests

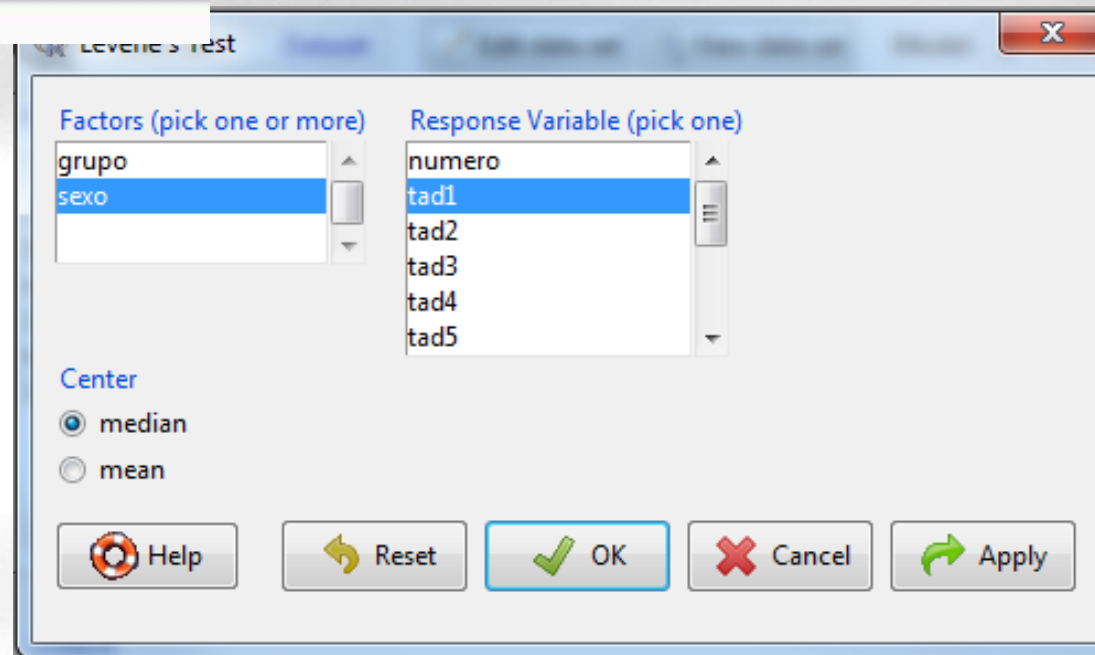
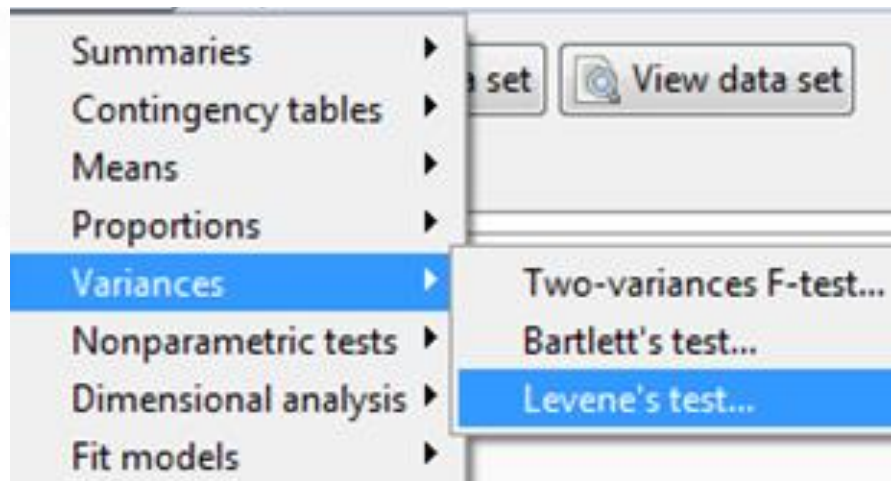


Two sample tests (1)



1. Data is normal (normality test) or sample size > 30 .
2. Mean is a good summary statistic for this problem.
3. Test homogeneity of variances

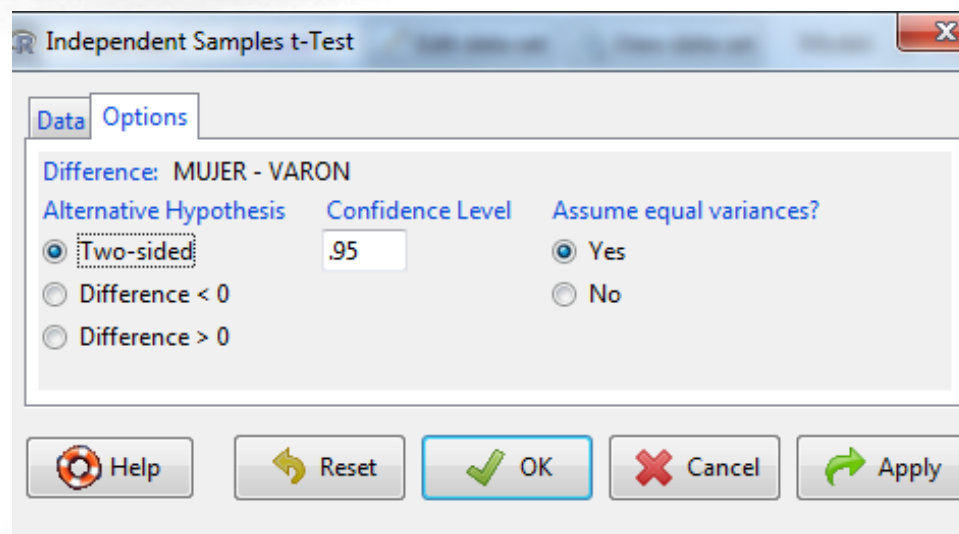
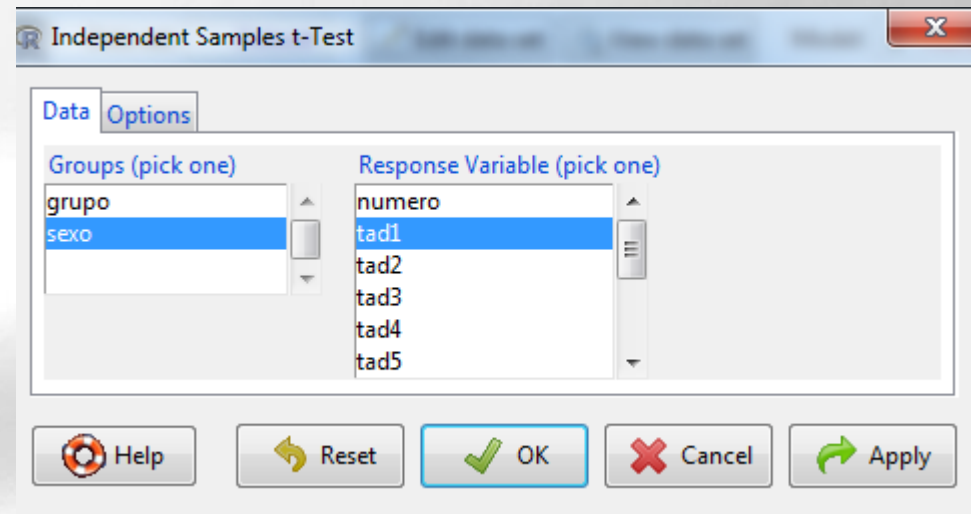
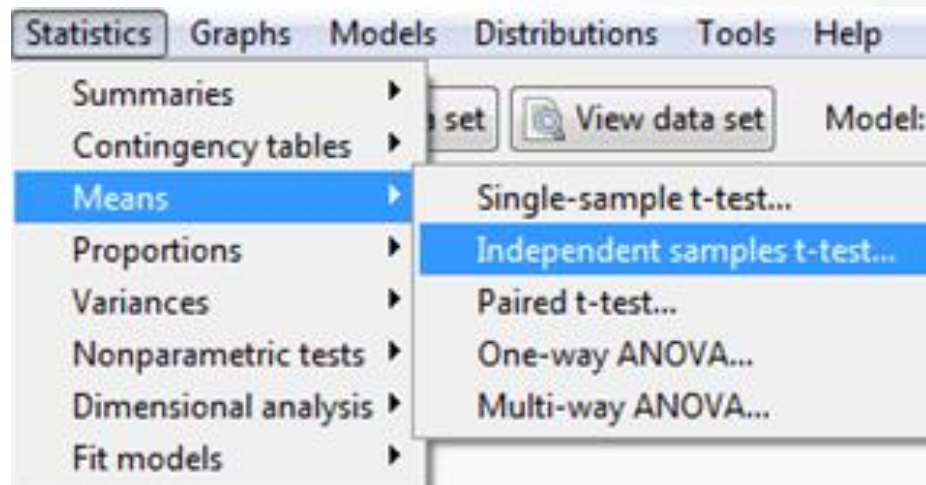
Testing variance homogeneity



```
> with(Dataset, tapply(tad1, sexo, var, na.rm=TRUE))  
      MUJER      VARON  
82.50751 138.24111  
  
> leveneTest(tad1 ~ sexo, data=Dataset, center="median")  
Levene's Test for Homogeneity of Variance (center = "median")  
      Df F value Pr(>F)  
group  1  1.3506 0.2499  
      58
```

- P value is over 0.05
- We can assume homogeneity of variances

T-test when variances are equal



T-test when variances are equal

```
> t.test(tad1~sexo, alternative='two.sided', conf.level=.95, var.equal=TRUE,  
+ data=Dataset)
```

Two Sample t-test

```
data: tad1 by sexo  
t = 0.3543, df = 58, p-value = 0.7244  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 -4.453505  6.368899  
sample estimates:  
mean in group MUJER mean in group VARON  
      88.78378      87.82609
```

Differences are not statistically significant

T-test when variances are not equal (Welch)

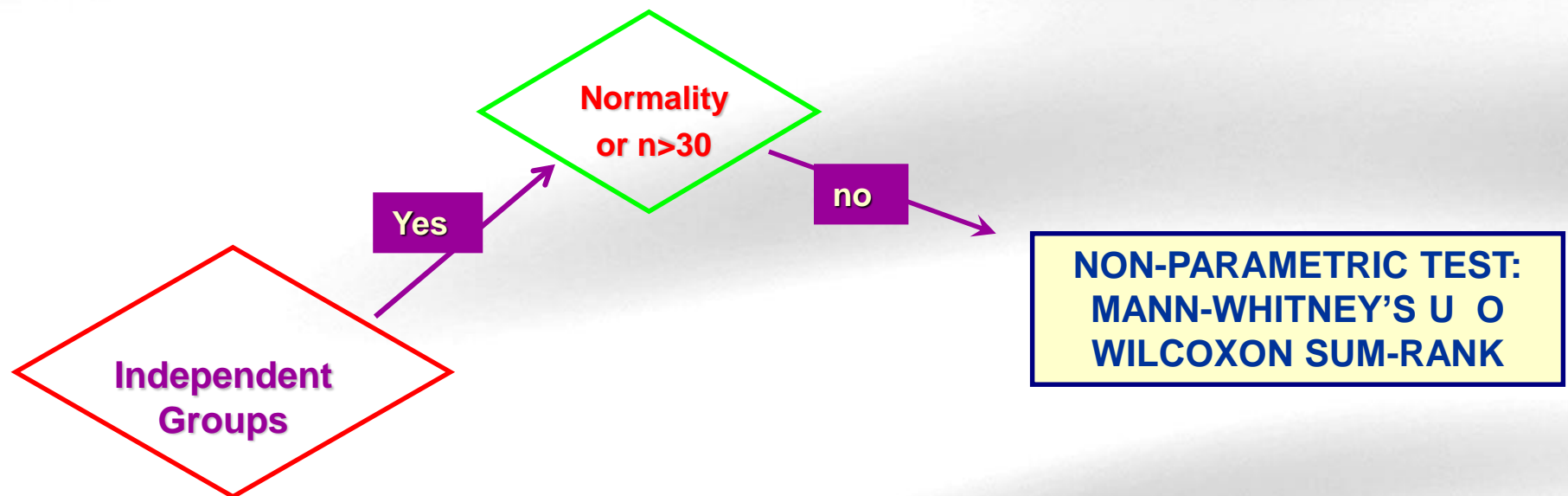
```
> t.test(tad1~sexo, alternative='two.sided', conf.level=.95, var.equal=FALSE,  
+ data=Dataset)
```

Welch Two Sample t-test

```
data: tad1 by sexo  
t = 0.3336, df = 38.144, p-value = 0.7405  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 -4.852834  6.768228  
sample estimates:  
mean in group MUJER mean in group VARON  
      88.78378      87.82609
```

Differences are not statistically significant

Two groups, data non normal



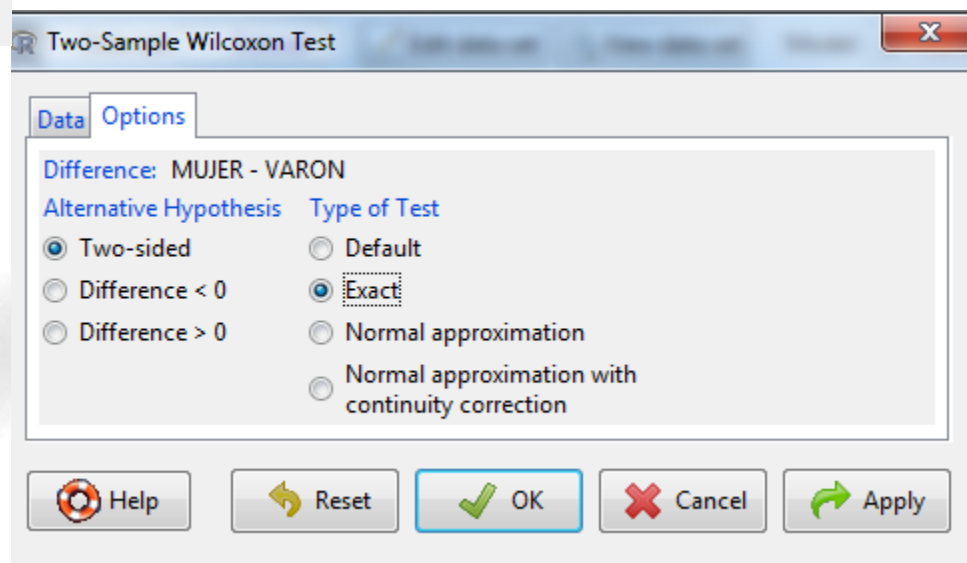
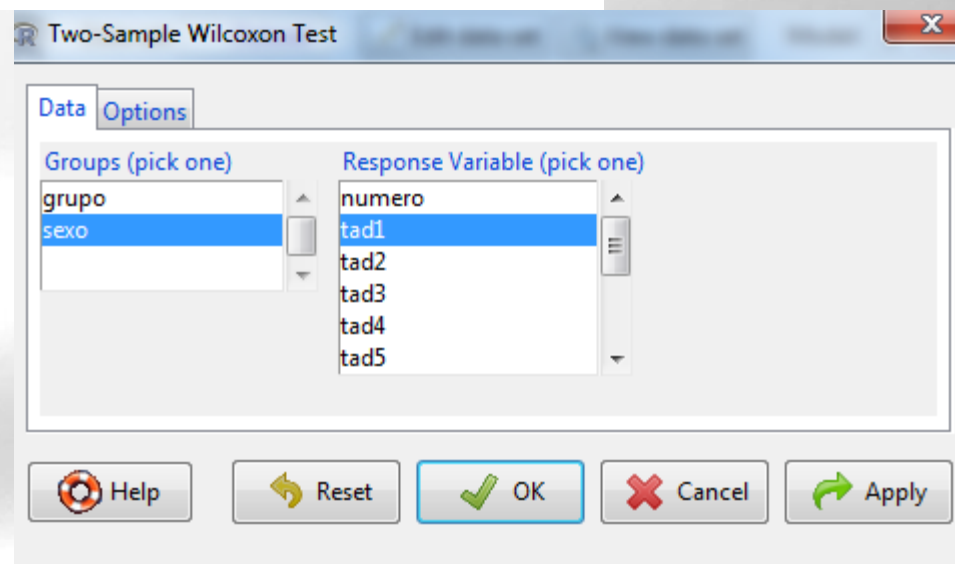
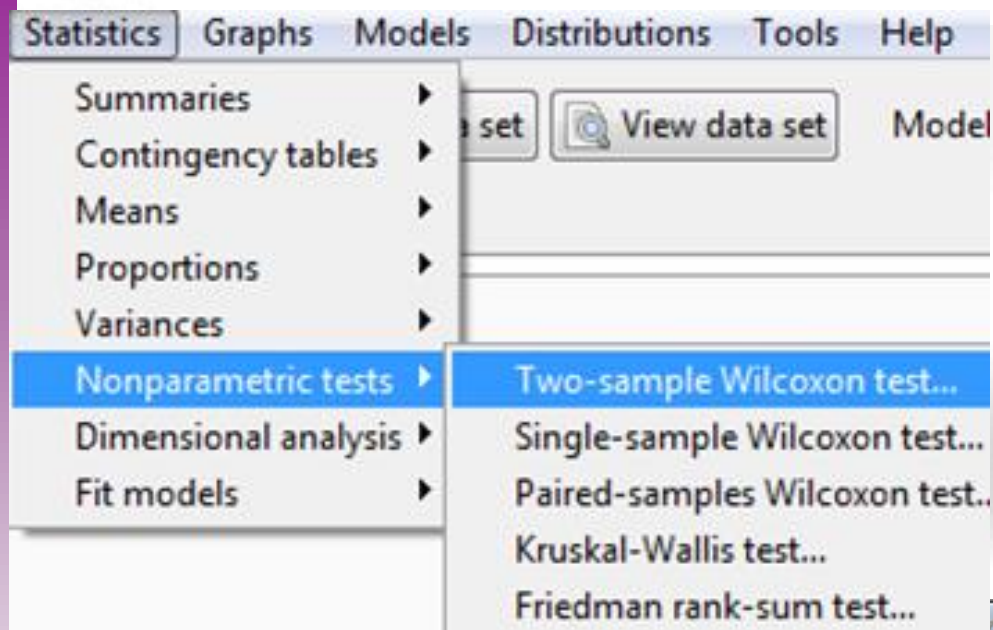
Non parametric tests

- If data distribution is unknown or mean is not the best way to summarize data ...
 - Non parametric test are not based on the usual parameters from a distribution, such as μ or σ^2 .
 - Instead they may be based ...
 - On order statistics, such as median or percentiles
 - They take into account the whole distribution.

Test based on ranks(Wilcoxon)

- Based on substituting original values by “ranks” in a joint sample
 - 12, 5, 14, 16, 3 → ranks are: 3, 2, 4, 5, 1
- Ranks only depend on the position of each value in the ordered sample.
 - 120, 95, 121, 130, 3 have the same ranks as values in the first sample
- ☺ NP test are more robust than parametric ones
- ☹ In the ideal situation where parametric tests are valid they are considered to be preferable.

Mann Whitney's U (Wilcoxon Rank-sum)



Mann Whitney's U (Wilcoxon Rank-sum)

```
> wilcox.test(tad1 ~ sexo, alternative='two.sided', exact=TRUE, correct=FALSE, data=Dataset)
```

```
Wilcoxon rank sum test
```

```
data: tad1 by sexo
```

```
W = 434, p-value = 0.8955
```

```
alternative hypothesis: true location shift is not equal to 0
```

```
> with(Dataset, tapply(tad1, sexo, median, na.rm=TRUE))
```

```
MUJER VARON
```

```
90      90
```

Differences are not significant

Questions to answer

- Are samples comparable at baseline time?
- Is blood pressure comparable between first and 12th measures?

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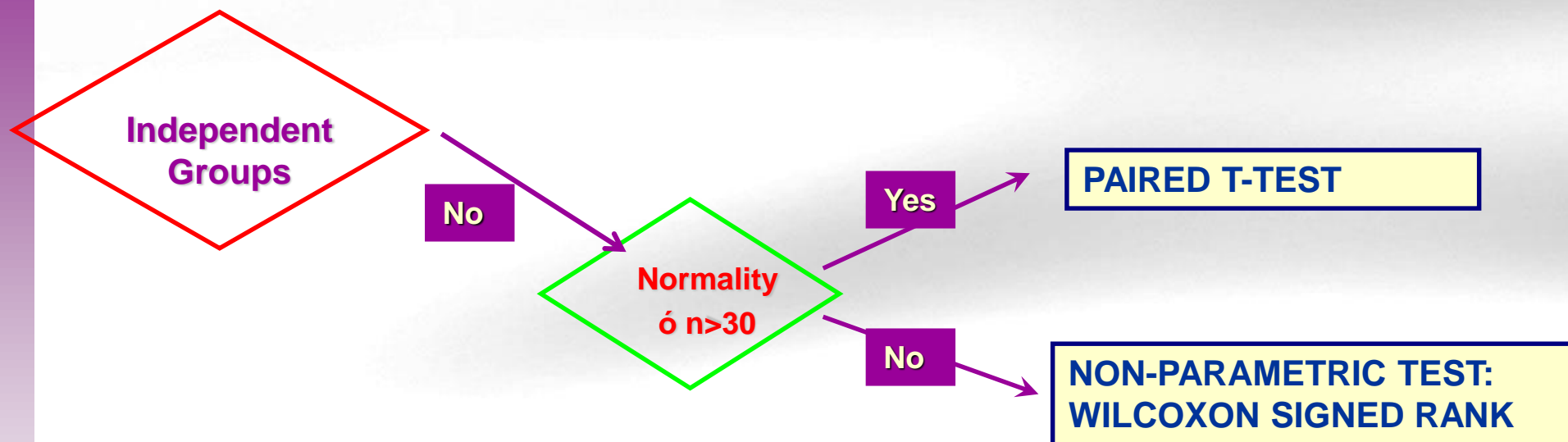
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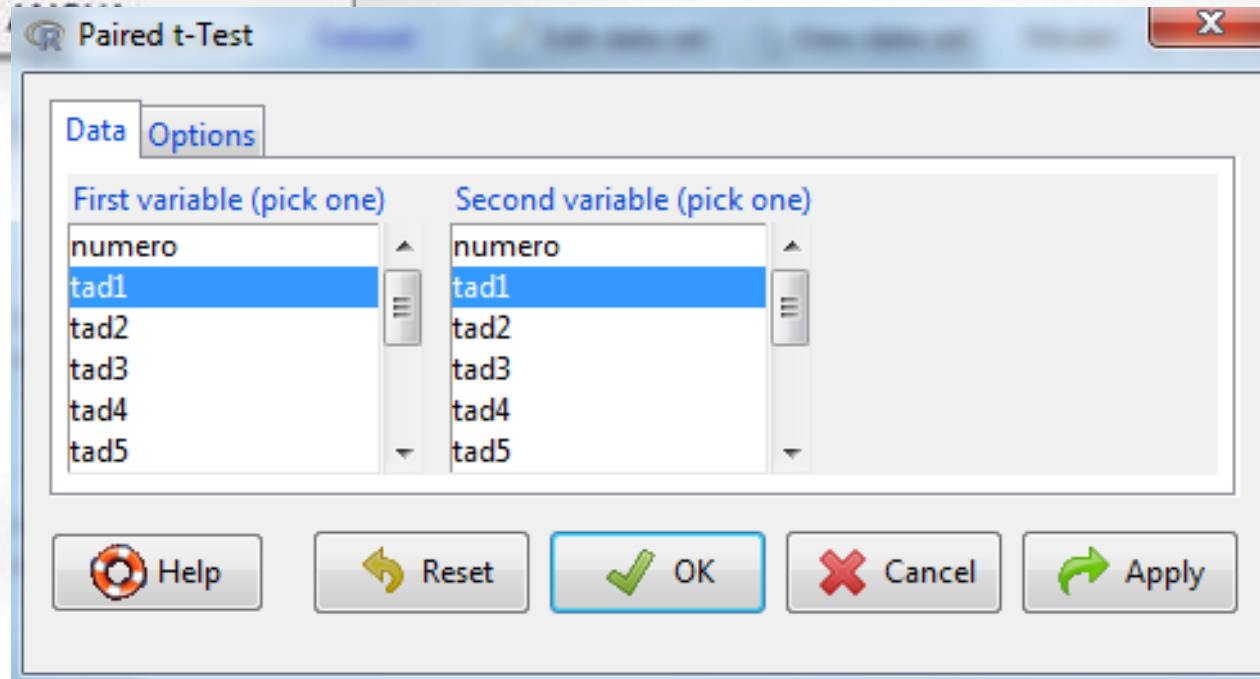
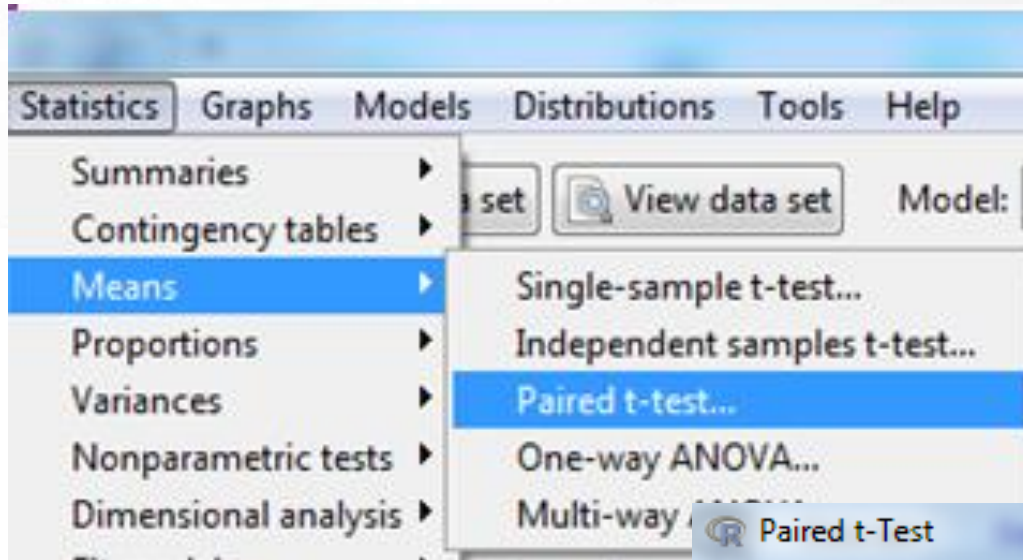
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Two dependent groups



Paired t-test



Paired t-test

```
> with(Dataset, (t.test(tad1, tad12, alternative='two.sided', conf.level=.95, paired=TRUE)))
```

```
Paired t-test
```

```
data: tad1 and tad12
```

```
t = 1.8507, df = 51, p-value = 0.07001
```

```
alternative hypothesis: true difference in means is not equal to 0
```

```
95 percent confidence interval:
```

```
-0.2364274  5.8133505
```

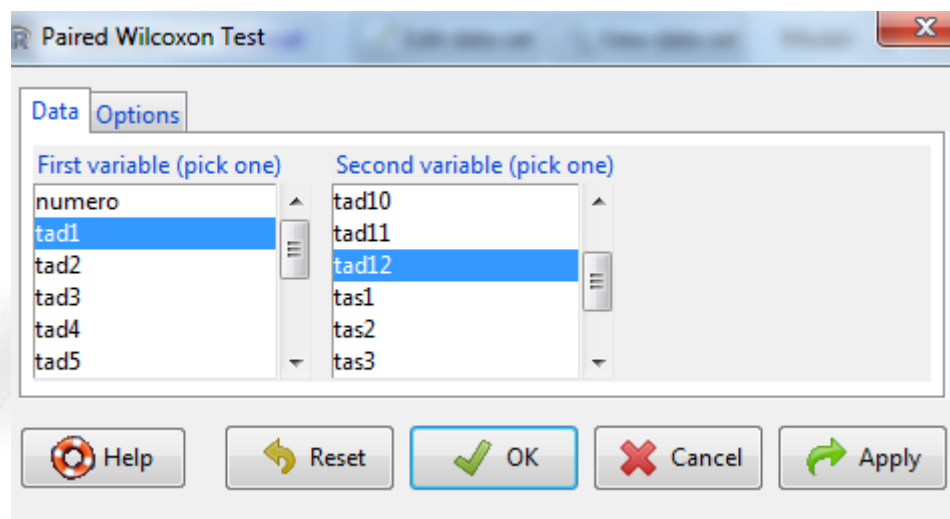
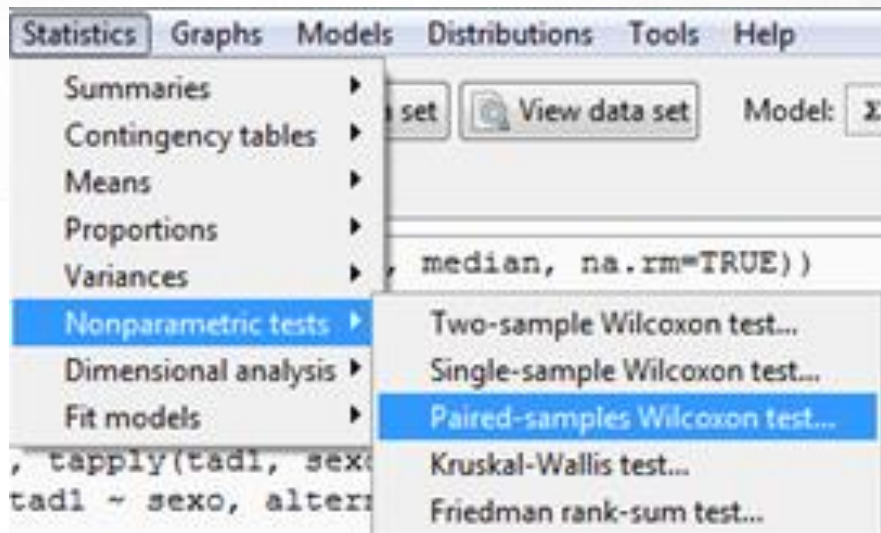
```
sample estimates:
```

```
mean of the differences
```

```
2.788462
```

Differences are not significant

Wilcoxon signed-rank test



Wilcoxon signed-rank test

```
> with(Dataset, wilcox.test(tad1, tad12, alternative='two.sided', exact=TRUE, paired=TRUE))
```

```
Wilcoxon signed rank test with continuity correction
```

```
data: tad1 and tad12
```

```
V = 478.5, p-value = 0.05333
```

```
alternative hypothesis: true location shift is not equal to 0
```

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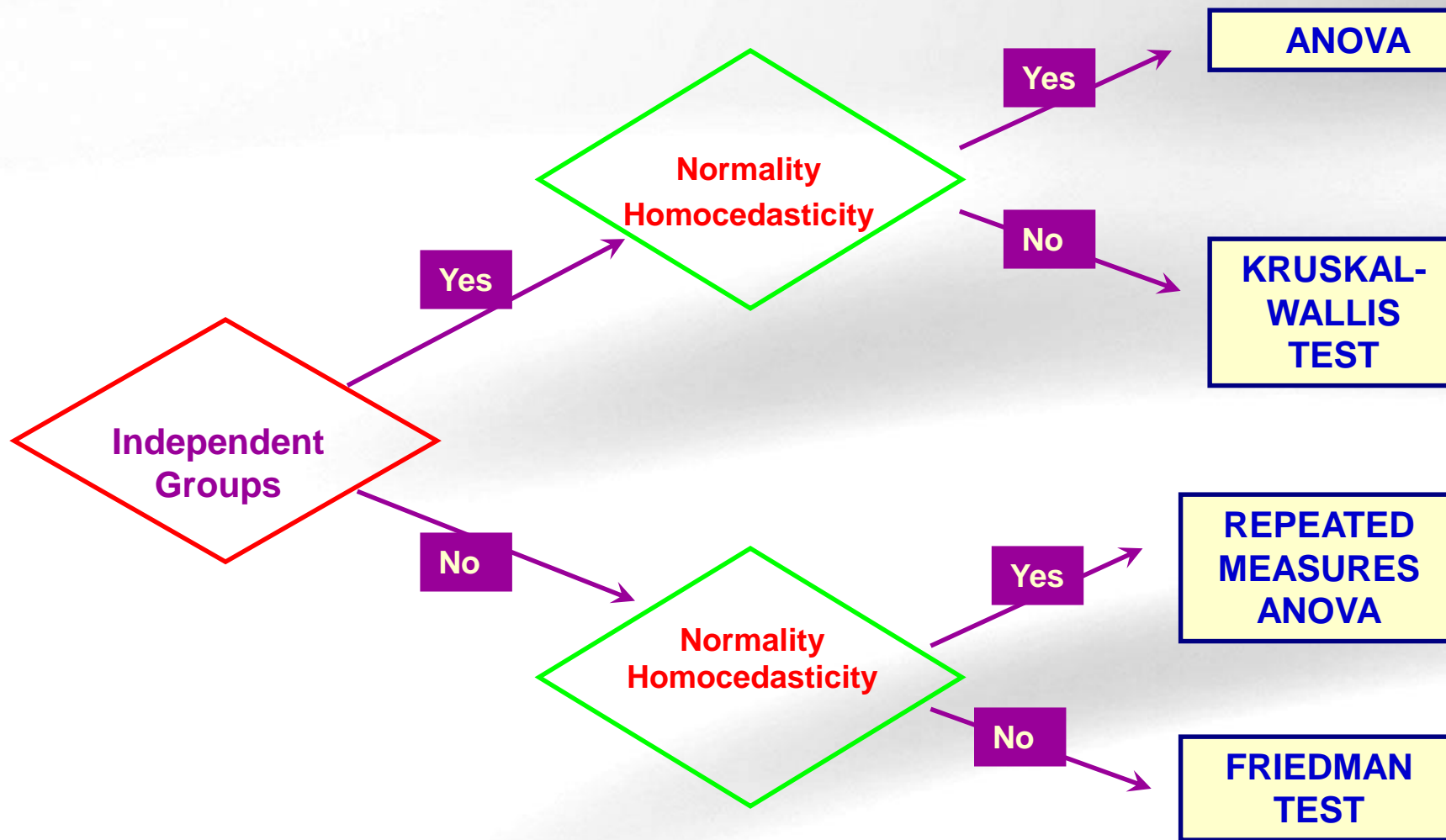
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Three or more groups to compare

Three or more groups



Analysis of the variance

Null Hypothesis

The means of all population are equal

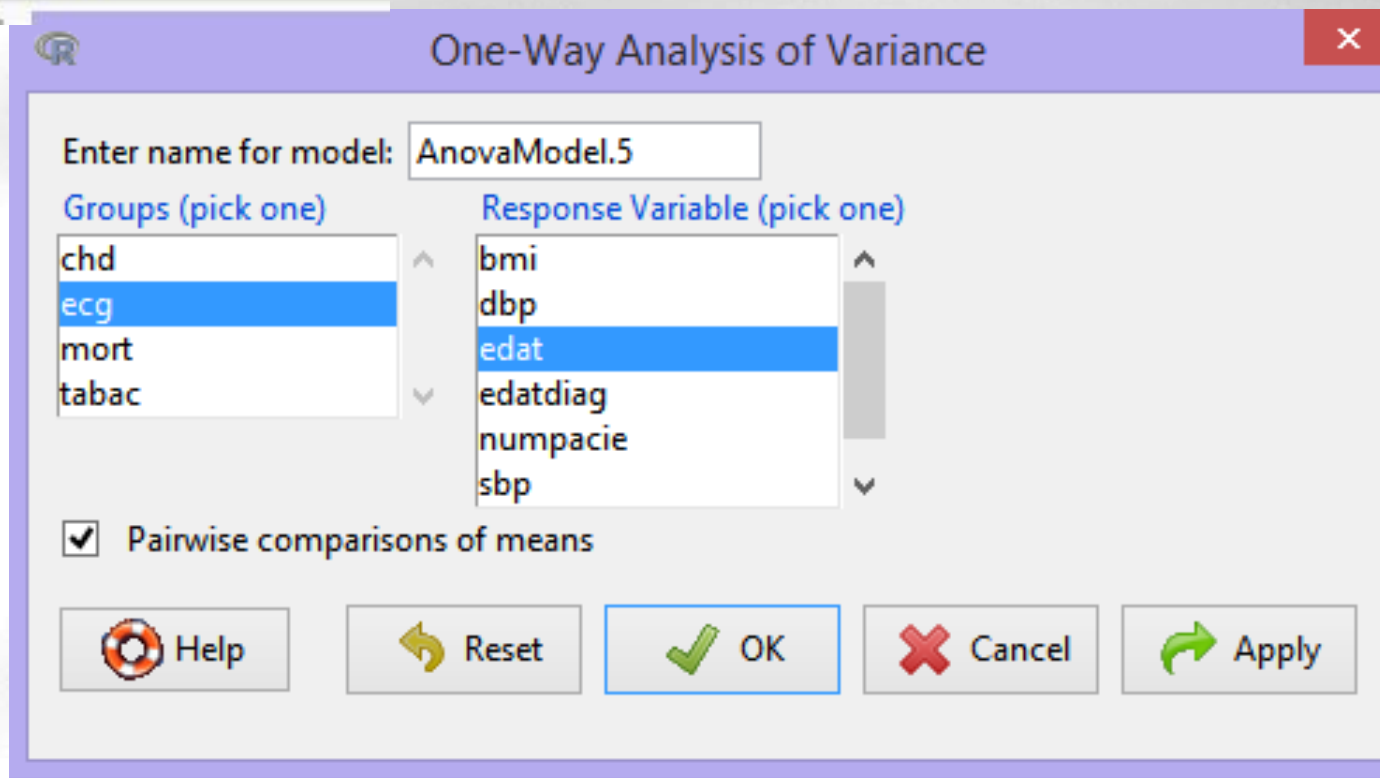
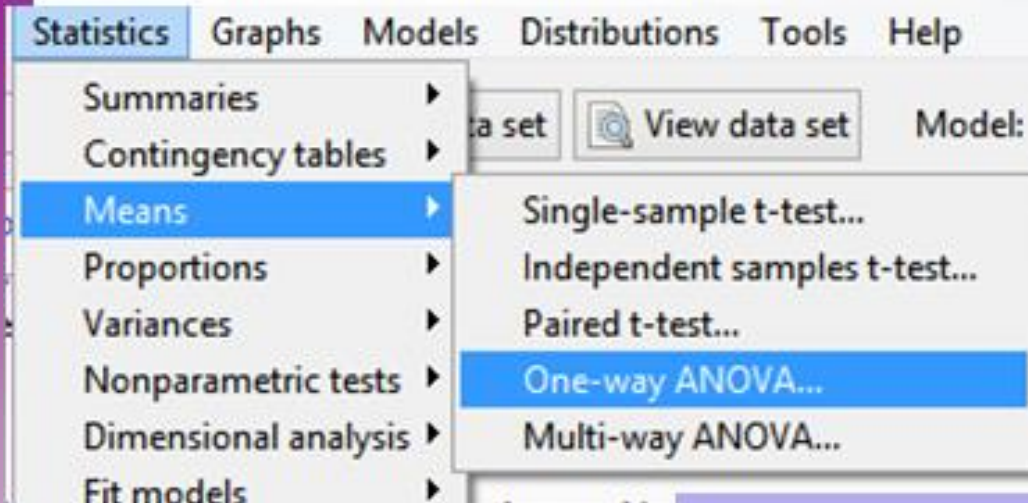
$$H_0 \quad \mu_1 = \mu_2 = \dots = \mu_k$$

Alternative Hypothesis

Not all the means are equal. At least there are two different means

$$H_a \quad \exists i, j \quad \mu_i \neq \mu_j$$

Anova in R Commander



```
> AnovaModel.4 <- aov(edat ~ ecg, data=diabetes)
> summary(AnovaModel.4)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
ecg	2	2166	1083.0	8.619	0.00029 ***
Residuals	146	18347	125.7		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```
> with(diabetes, numSummary(edat, groups=ecg, statistics=c("mean", "sd")))
```

	mean	sd	data:n
Normal	50.50450	11.492981	111
Frontera	53.81481	11.368097	27
Anormal	64.90909	6.759505	11

Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

```
Fit: aov(formula = edat ~ ecg, data = diabetes)
```

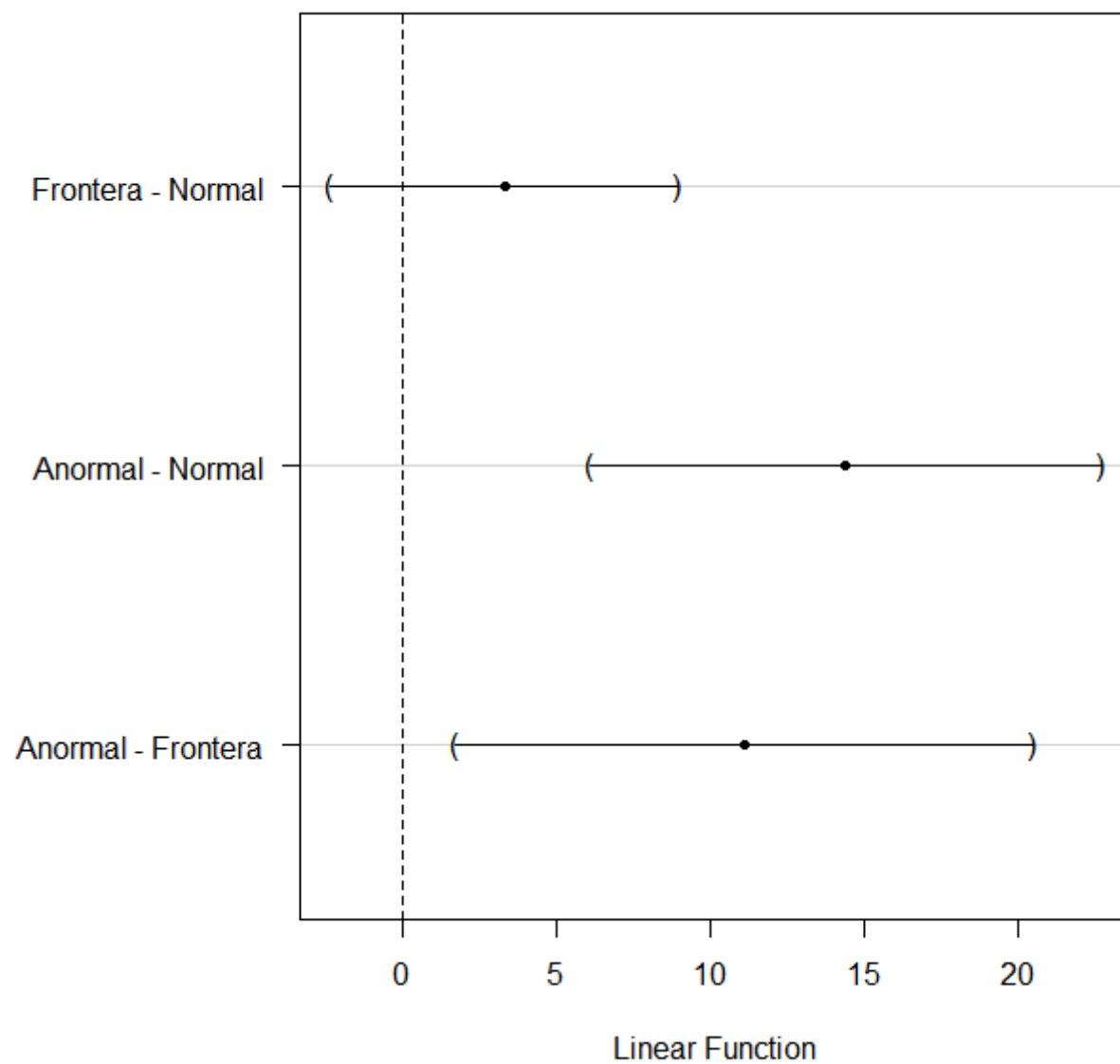
Linear Hypotheses:

	Estimate	Std. Error	t value	Pr(> t)
Frontera - Normal == 0	3.310	2.405	1.376	0.345713
Anormal - Normal == 0	14.405	3.543	4.065	0.000217 ***
Anormal - Frontera == 0	11.094	4.010	2.767	0.016472 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Adjusted p values reported -- single-step method)

95% family-wise confidence level



Multiple Comparisons



To cross or not to cross?



Imagine you are an adventurer that has the option of to cross a bridge in order to escape from danger, find a treasure...

and that there is a post in front of the bridge stating:

"This bridge has broken only one out of 100 times"



To cross or not to cross?



Imagine you are an adventurer that has the option of to cross a bridge in order to escape from danger, find a treasure...

and that there is a post in front of the bridge stating:

“This bridge has broken only one out of 100 times”

So, the p-value of our metaphor is 0.01

You could accept that **1%** is a **risk small** enough to **pass the bridge** and pursue your goal. OK



To cross or not to cross?



But... what do you decide if, in order to reach your goal, you have to cross hundreds of bridges of that kind?





To cross or not to cross?



But... what do you decide if, in order to reach your goal, you have to cross hundreds of bridges of that kind?

- In this case, the probability of falling while crossing one of the bridges is obviously too high ('cause we have just one life).





To cross or not to cross?



Therefore, in this case (multiple testing), the p-value by itself is not a good reference for accepting or not statistical significance.

We must apply some type of correction to the p-values (allowing us to be safe in crossing all the bridges).

- Bonferroni (α/k)
- Post-Hoc test ANOVA (Tukey, Scheffe, Dunn-test)
- False Discovery rate
- Benjamini-Hochberg correction



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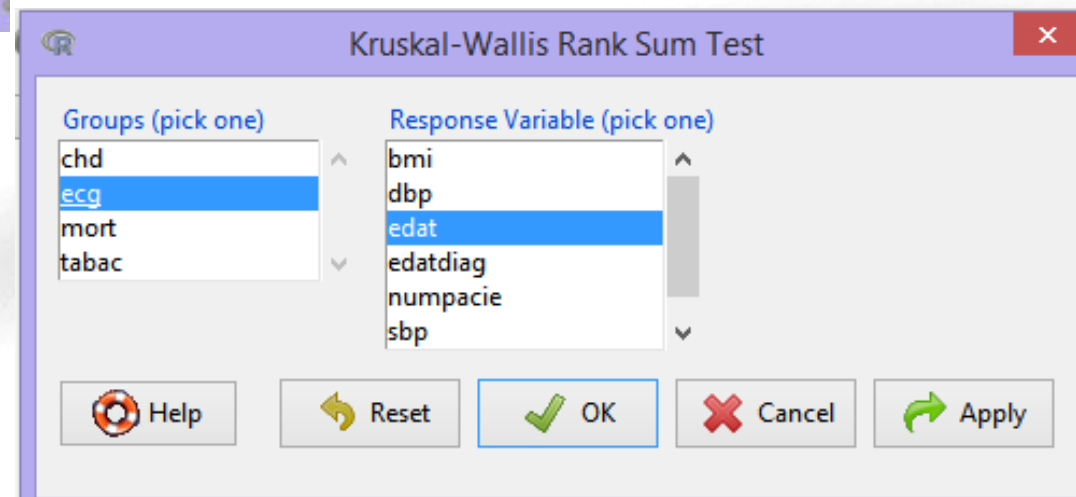
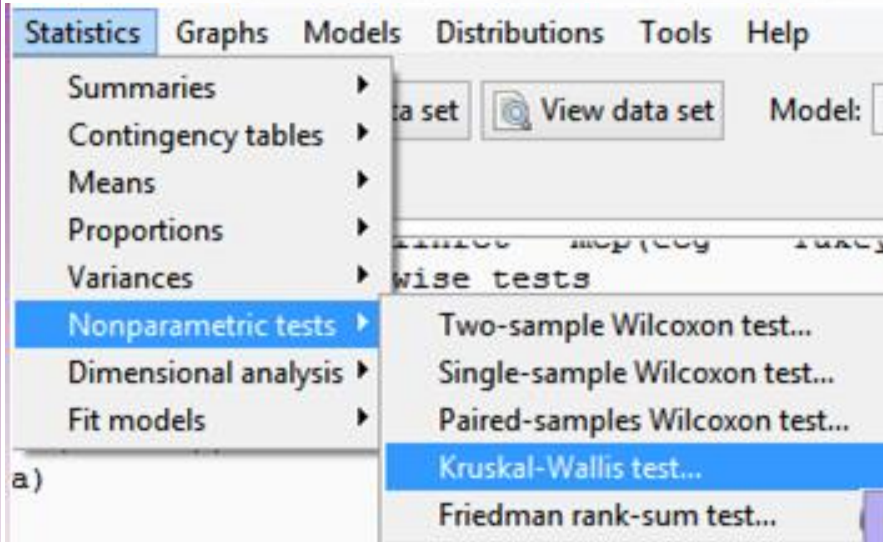
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Kruskal-Wallis test

- Is the non-parametric versió of ANOVA based on ranks



Kruskal-Wallis Test

```
> with(diabetes, tapply(edat, ecg, median, na.rm=TRUE))
```

Normal	Frontera	Anormal
49	53	64

```
> kruskal.test(edat ~ ecg, data=diabetes)
```

Kruskal-Wallis rank sum test

data: edat by ecg

Kruskal-Wallis chi-squared = 17.4826, df = 2, p-value = 0.0001598

Exercise

- Is there a relation between follow up time (tempsviu) and death (mort)?
- Is there a relation between systolic blood pressure (bdp) and ECG?



Common misunderstandings about the p-value

- The p-value is **not** the probability that the null hypothesis is true, nor it is the probability that the alternative hypothesis is false (it is not connected to either of these).
- The p-value **cannot** be used to figure out the probability of a hypothesis being true.
- The p-value is **not** the probability of wrongly rejecting the null hypothesis.
- The p-value is **not** the probability that replicating the experiment would yield the same conclusion.
- The p-value does **not** indicate the size or importance of the observed effect. The two do vary together however: the larger the effect (effect size), the smaller sample size will be required to get a significant p-value.