

Hypothesis Testing. Quantitative Variables

Basic Statistics with R
UEB-VHIR

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Syllabus

1. INTRODUCTION
2. TYPE OF TEST
3. NORMALITY TEST
4. **ONE GROUP** COMPARISON
5. **TWO GROUPS** COMPARISON IN INDEPENDENT SAMPLES
6. **TWO GROUPS** COMPARISON IN DEPENDENT SAMPLES
7. **MORE THAN TWO GROUPS** COMPARISON IN INDEPENDENT SAMPLES
8. MULTIPLE COMPARISONS AND MULTIPLE TESTING

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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**4. TWO GROUPS COMPARISON IN INDEPENDENT
SAMPLES**

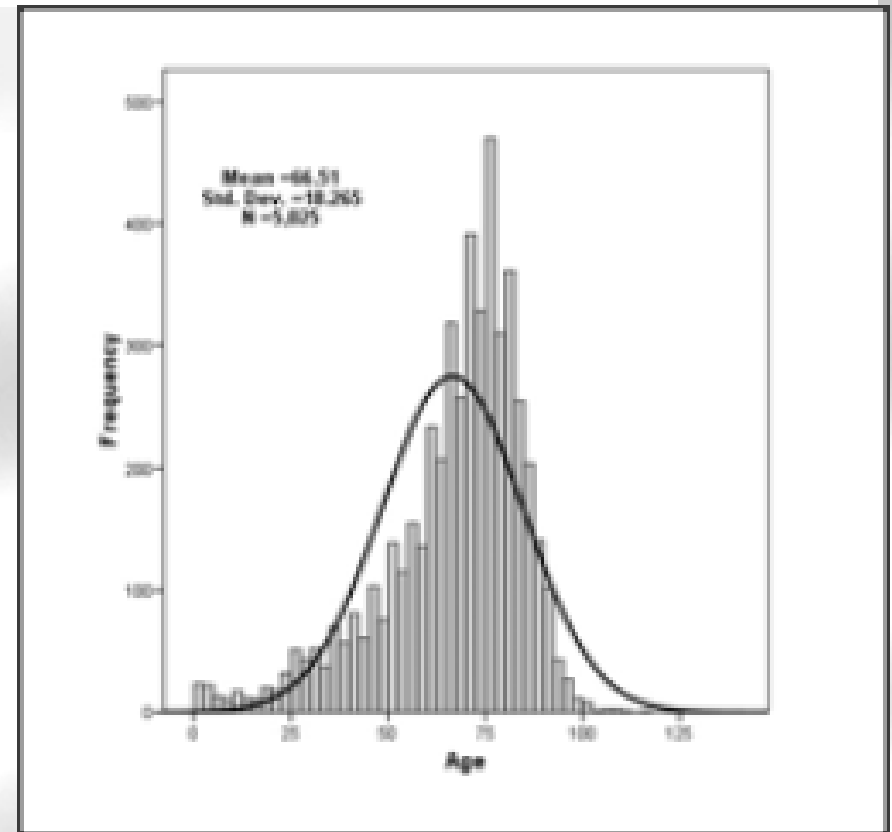
**5. TWO GROUPS COMPARISON IN DEPENDENT
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Normality test

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

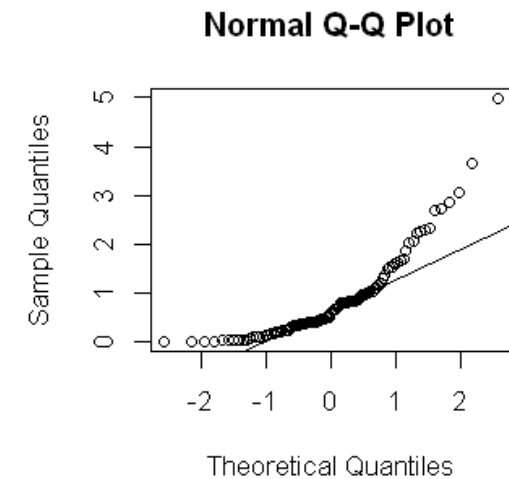
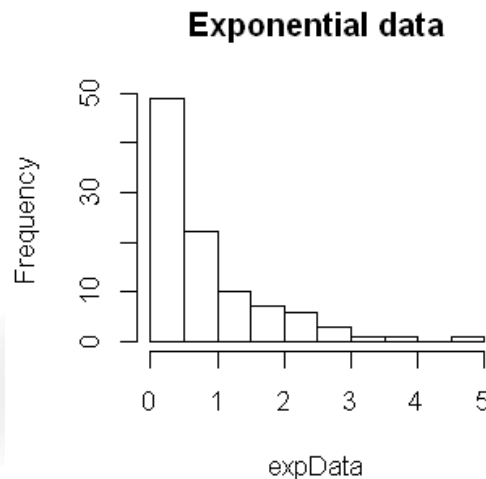
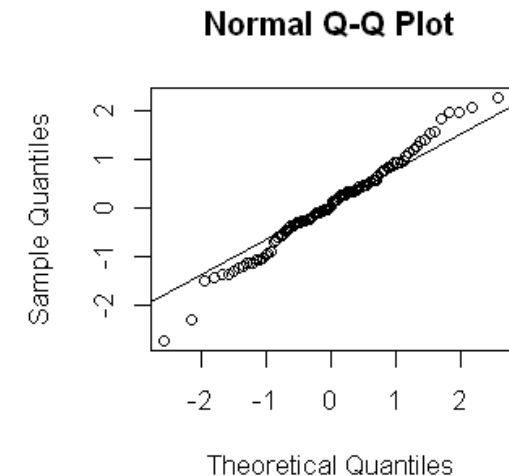
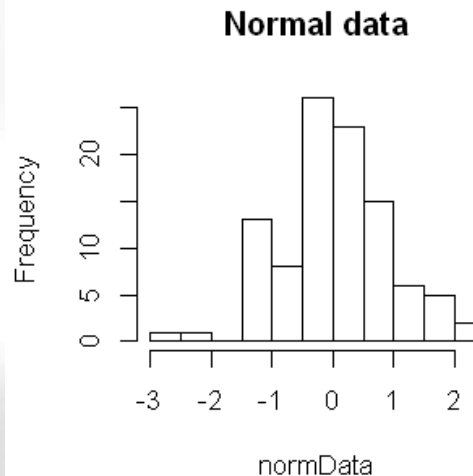


Testing normality

- 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

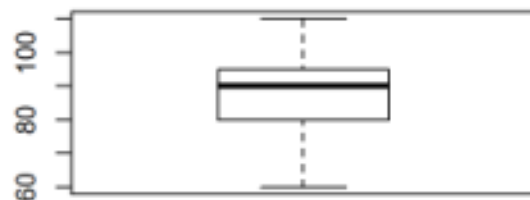
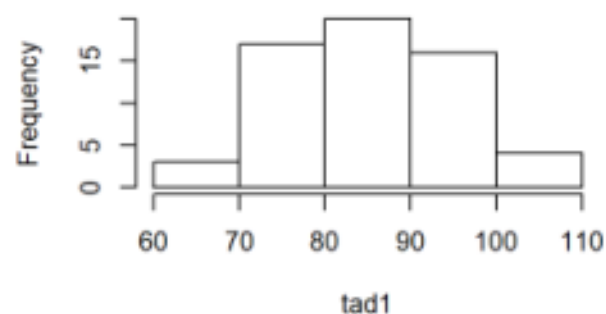
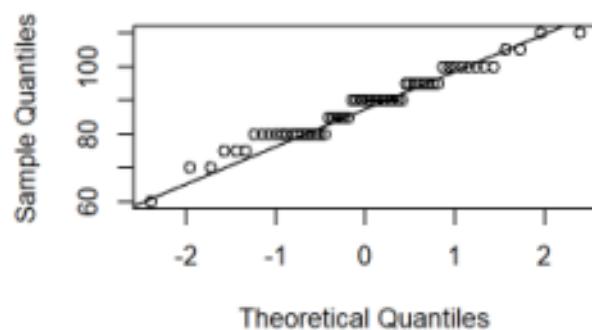



```
oldpar<-par(mfrow=c(1,1)) # Guarda los parámetros para el dibujo
par(mfrow=c(2,2)) # Dibuja cuatro gráficos por grafico
with(hta, boxplot(tad1, main="Box-plot") )

with(hta, hist(tad1) )

with(hta, qqnorm(tad1, main="Normal QQplot") )
;with(hta, qqline(tad1) )

par(oldpar) # Vuelve a los parámetros de dibujo originales
```

Box-plot**Histogram of tad1****Normal QQplot**

- 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

Normality Test

```
```{r normtest , mysize=TRUE, size='\\small'}  
with(hta,shapiro.test(tad1)) # Shapiro wilk test
...`
```

Shapiro-Wilk normality test

data: tad1  
W = 0.96622, p-value = 0.09512

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# 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 Hipertensive patients?*

# One sample Test

```
with(hta,t.test(tad1,mu=90)) # One sample T.test
```

```

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
```

# Exercise 1

---

- a) Check the normality of *tas1* variable in **hta** dataset
- b) Can it be accepted that the initial TAS is 120 in Hipertensive patients?
- c) Find the 95% confidence interval for the mean of *tas1* variable
- d) **Extra:** Can it be accepted that the initial TAS is *higher than* 120 in Hipertensive *women*?

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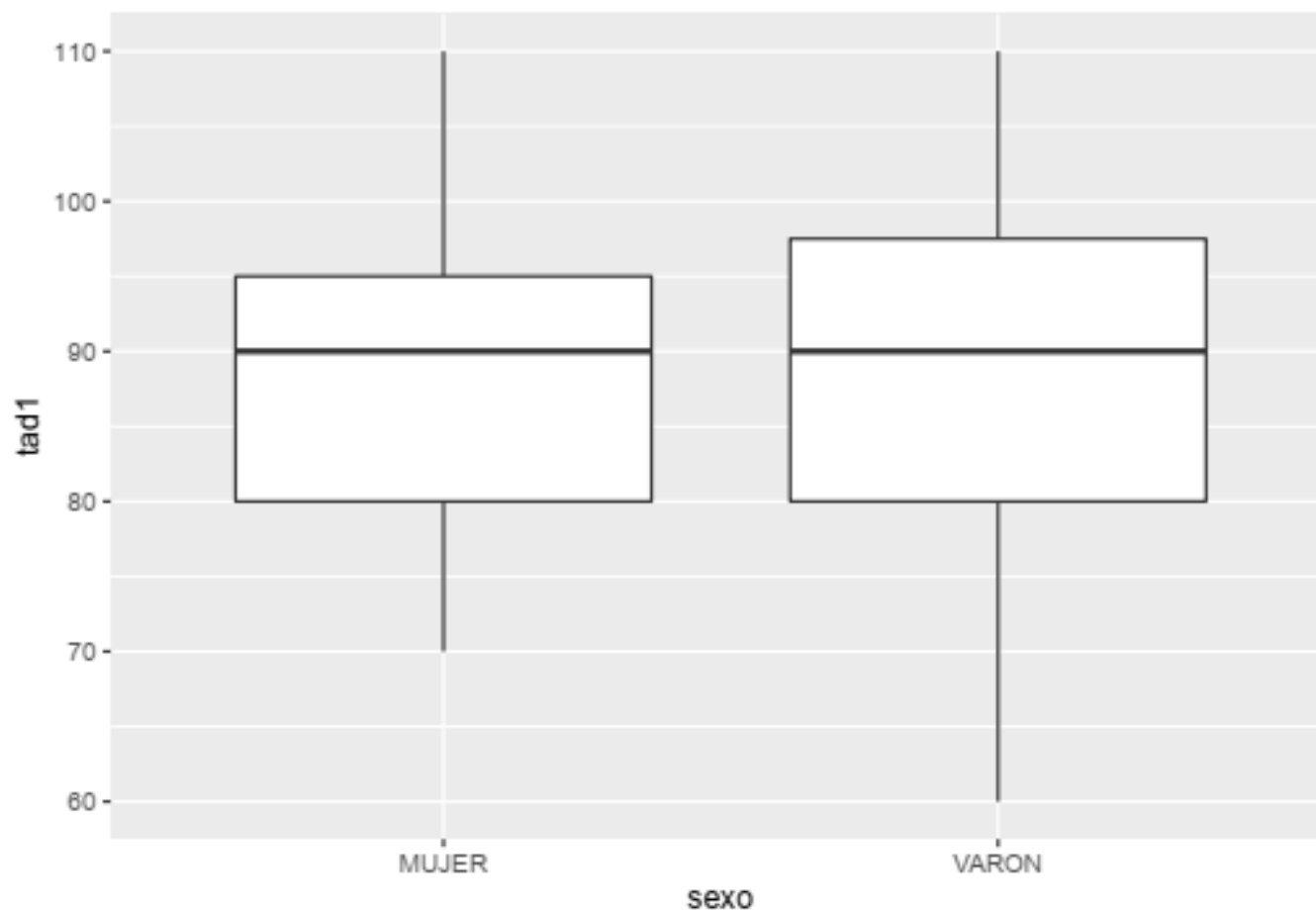
# Questions to answer

---

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

# Boxplot *tad1*, by *sexo*

```
ggplot(hta, aes(x=sexo, y=tad1)) +
 geom_boxplot()
```



# 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



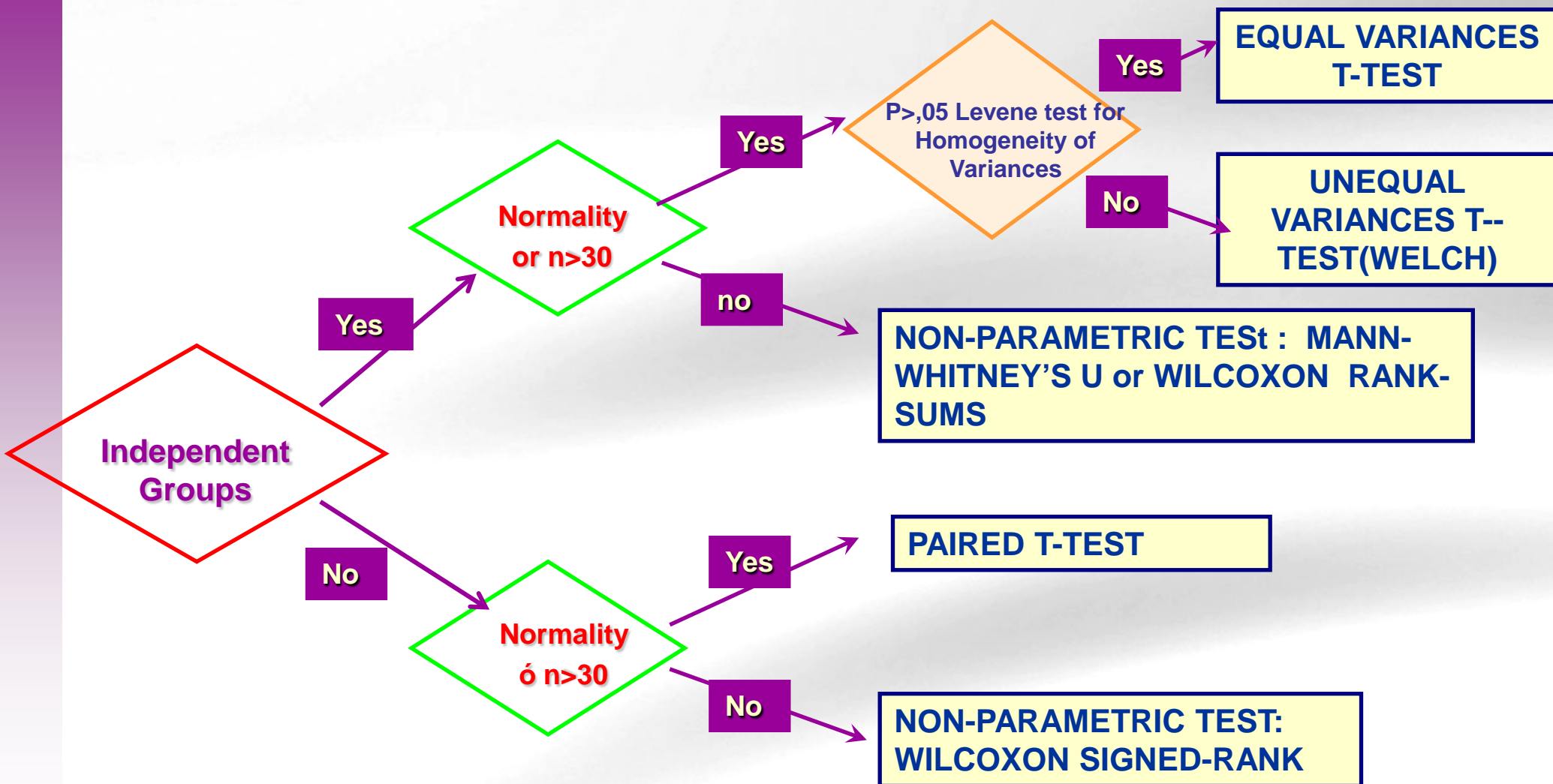
## INDEPENDENT

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

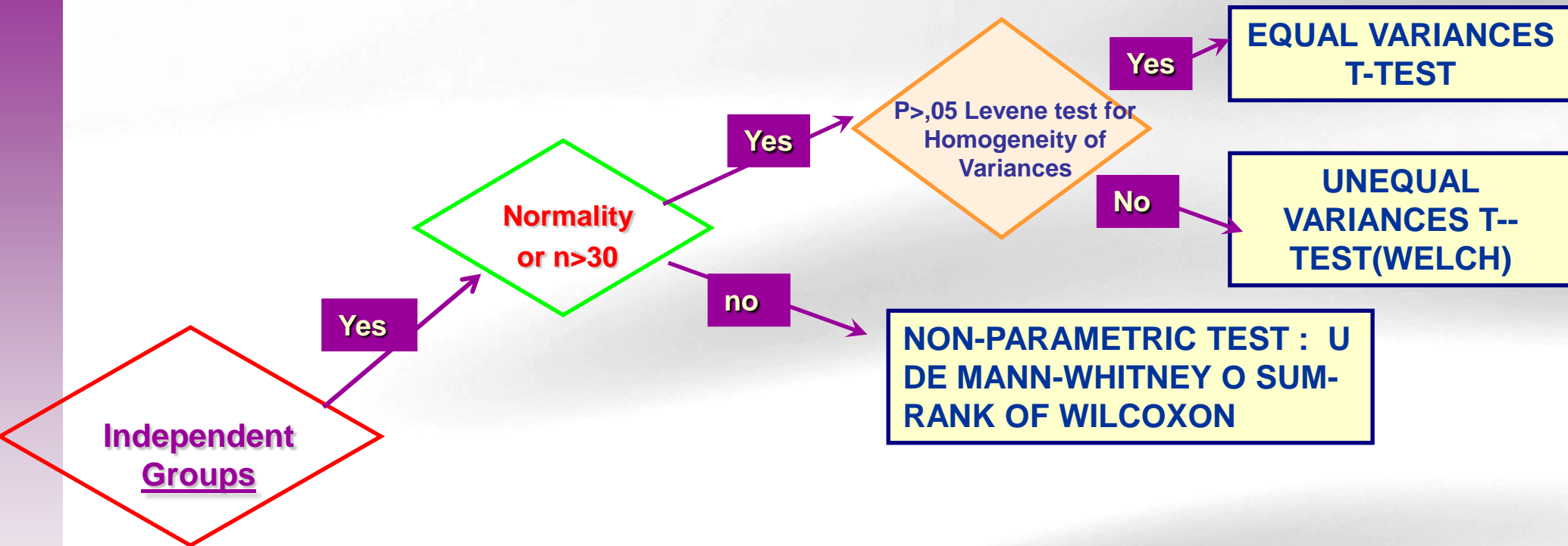
## DEPENDENT

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

# Homogeneity variance Test

```
library(car)
hta%>%
 group_by(sexo) %>%
 summarize(var = sd(tad1))
```

```
A tibble: 2 x 2
sexo var
<chr> <dbl>
1 MUJER 9.08
2 VARON 11.8
```

```
with(hta, leveneTest(tad1~factor(sexo), 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

```
with(hta,t.test(tad1~factor(sexo),var.equal=TRUE))

##
Two Sample t-test
##
data: tad1 by factor(sexo)
t = 0.35427, df = 58, p-value = 0.7244
alternative hypothesis: true difference in means is not equal
95 percent confidence interval:
-4.453505 6.368899
sample estimates:
mean in group MUJER mean in group VARON
88.78378 87.82609
```

- Type I Error is over than 0.05
- We cannot reject mean equality

# T test when variances are unequal

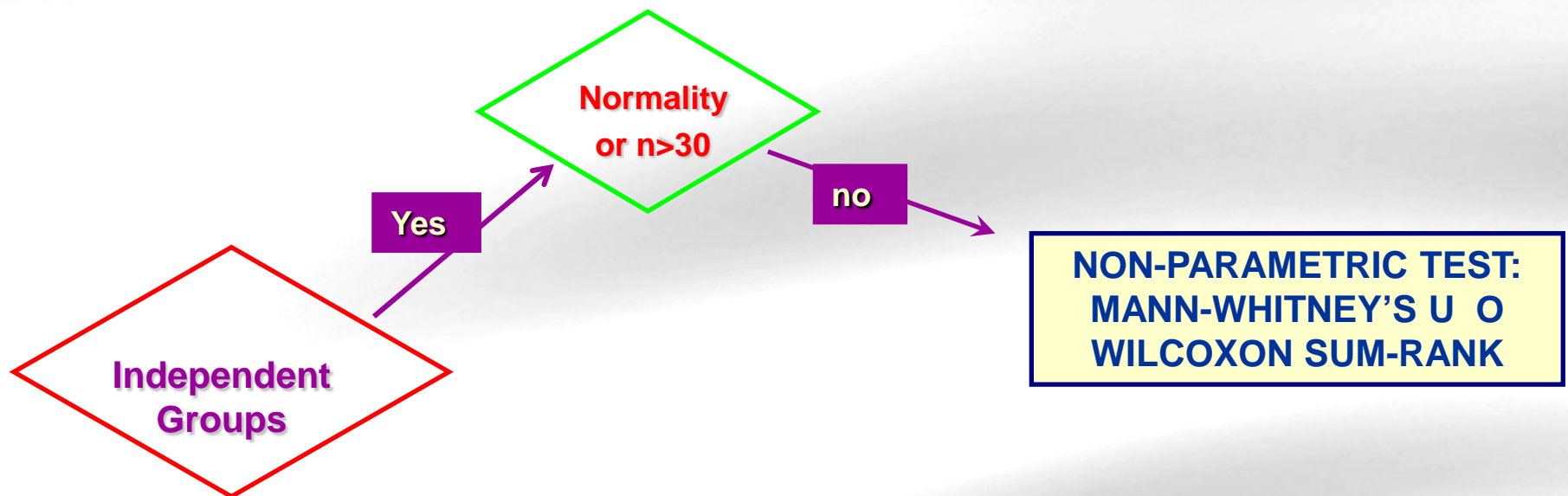
```
with(hta,t.test(tad1~factor(sexo),var.equal=FALSE))

##
Welch Two Sample t-test
##
data: tad1 by factor(sexo)
t = 0.33362, df = 38.144, p-value = 0.7405
alternative hypothesis: true difference in means is not equal
95 percent confidence interval:
-4.852834 6.768228
sample estimates:
mean in group MUJER mean in group VARON
88.78378 87.82609
```

- Same conclusions as before
- Test is also known as Welch test



# 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  $\mu$  or  $\sigma^2$ .
  - 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.

# U Mann-Whitney or Sum Rank non parametric test

```
with(hta,wilcox.test(tad1~factor(sexo)
,alternative='two.sided',exact=TRUE, correct=FALSE))
```

```
##
Wilcoxon rank sum test
##
data: tad1 by factor(sexo)
W = 434, p-value = 0.8955
alternative hypothesis: true location shift is not equal to 0
```

```
hta%>%
 group_by(sexo) %>%
 summarize(median = median(tad1))
```

```
A tibble: 2 x 2
sexo median
<chr> <dbl>
1 MUJER 90
2 VARON 90
```

- Null Hypothesis cannot be rejected

# Exercise 2

---

- Is TAD comparable at baseline time between groups?
  - a) What is the Hypothesis that we want to test? Describe the null hypothesis and the alternative hypothesis.
  - b) What test would be appropriate to perform to answer the question?
  - c) Answer the question
  - d) Apply a non-parametric test and compare the results

# Questions to answer

---

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

# 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	

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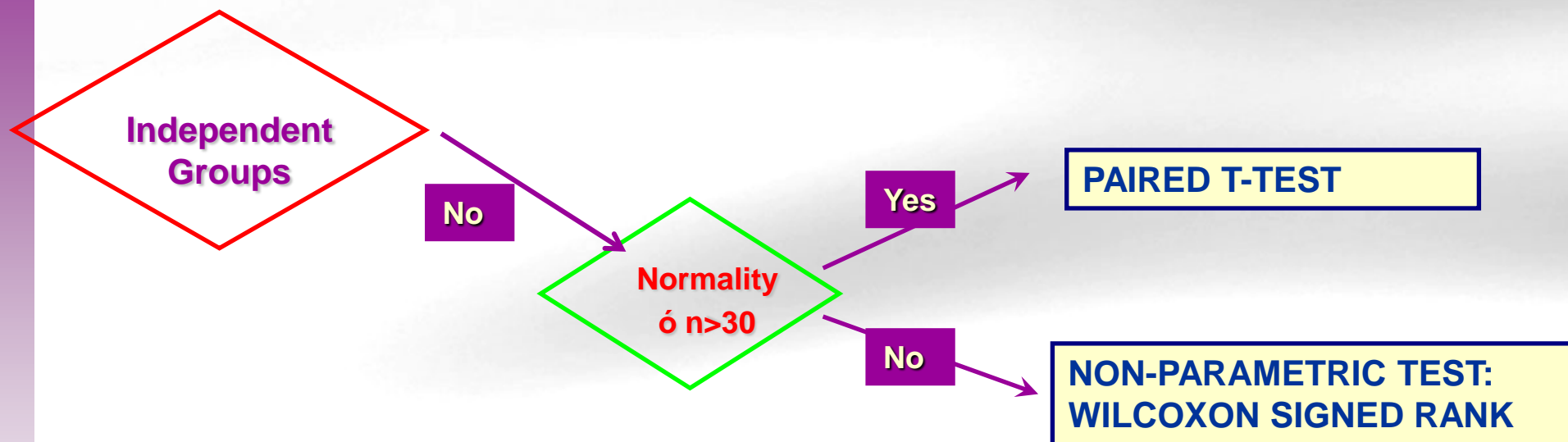
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# Two dependent groups



# Questions to answer

---

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

# Paired T-test

```
with(hta,t.test(tad1,tad12,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
95 percent confidence interval:
-0.2364274 5.8133505
sample estimates:
mean of the differences
2.788462
```

- P value is over 0.05

# Paired Sign-Rank Wilcoxon Test

```
with(hta,wilcox.test(tad1,tad12,
 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
```

## Exercise 3

---

- Is systolic blood pressure (TAS) comparable between first and 12th measures?



# Exercise 4

---

Indica que tipo de análisis o que pruebas estadísticas utilizarías y si fuera necesario algún tipo de prueba adicional para llevar a cabo el análisis. Formula la hipótesis a contrastar de acuerdo con las hipótesis seleccionadas

- a. Se efectúa un estudio de seguimiento a 1018 sujetos atendidos en una clínica de obesidad. Se mide el Índice de Masa Corporal(IMC) y el perfil lipídico. Al cabo de 12 meses se evalúa de nuevo el IMC y el colesterol estando interesados en cuantificar la disminución de ambos parámetros
- b. Se analizan un grupo de variables inmunológica(leucocitos totales, linfocitos B, natural Killer, etc) en una muestra de 102 hombres y 147 mujeres mayores de 65 años. Se está interesado en ver la existencia de diferencias por sexo.

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# Question to answer

---

A pharmaceutical laboratory wants to test which of three drugs are better:



drug 1



drug 2



drug 3



# Question to answer

Could we use Student's t test? We will see with an example:

A pharmaceutical laboratory wants to test which of three drugs are better:



drug 1



drug 2









drug 3

To know which of the drugs is the best one, one could think to perform the following comparison using a t test:

# Question to answer

Could we use Student's t test? We will see with an example:

Comparisons with separate t test would be:

			<u>Chance of Type I error</u>	<u>Chance of Accept H0</u>
	<u>vs</u>		$\alpha = 5\%$	$1 - \alpha = 95\%$
	<u>vs</u>		$\alpha = 5\%$	$1 - \alpha = 95\%$
	<u>vs</u>		$\alpha = 5\%$	$1 - \alpha = 95\%$

# Question to answer

Could we use Student's t test? We will see with an example:

Comparisons with t test would be:



vs



Chance of Type I error



$$\alpha = 5\%$$

Chance of Accept H0

$$1 - \alpha = 95\%$$



vs



$$\alpha \approx 10\%$$

$$1 - \alpha = 95\% * 95\%$$



vs



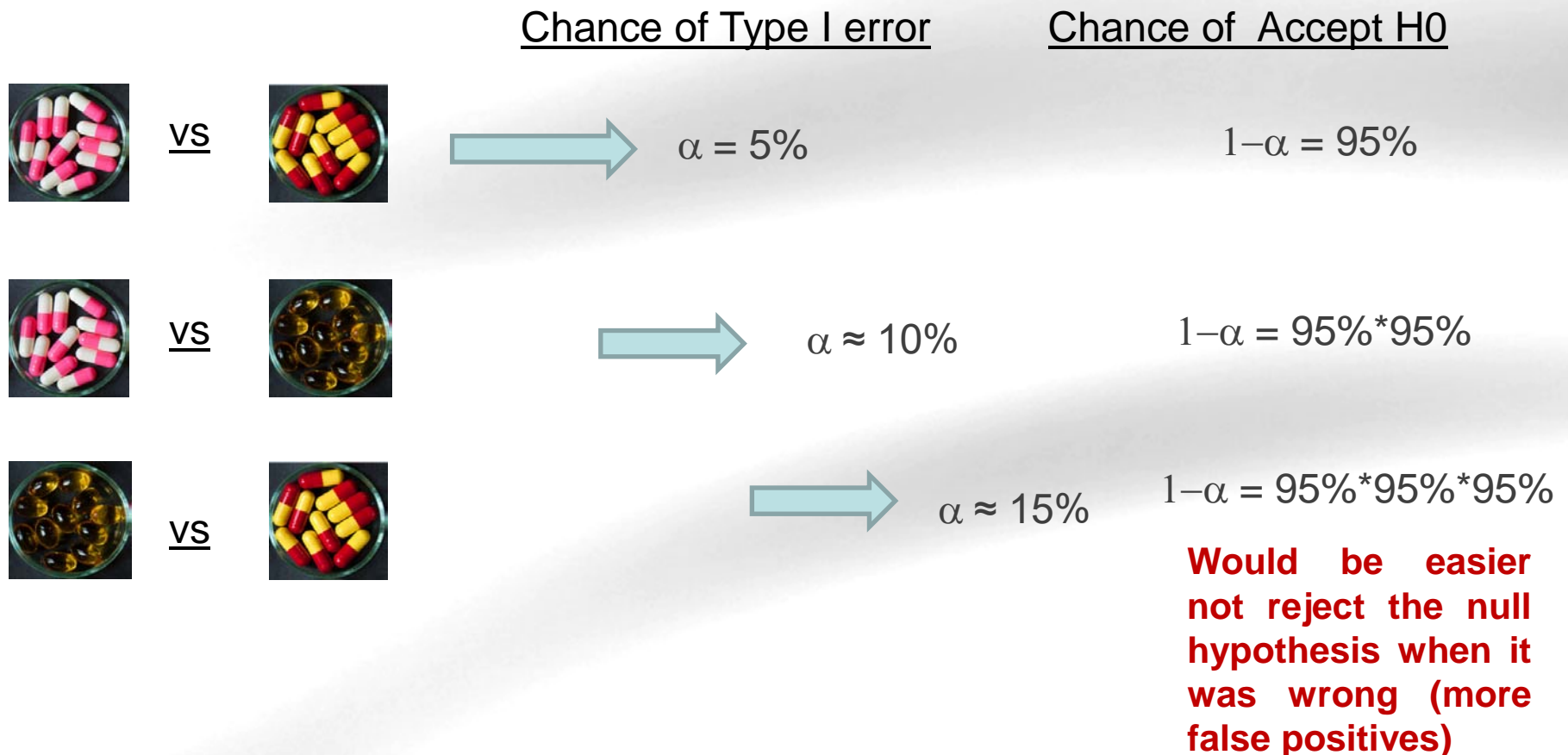
$$\alpha \approx 15\%$$

$$1 - \alpha = 95\% * 95\% * 95\%$$

# Question to answer

Could we use Student's t test? We will see with an example:

Comparisons with t test would be:





# 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

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

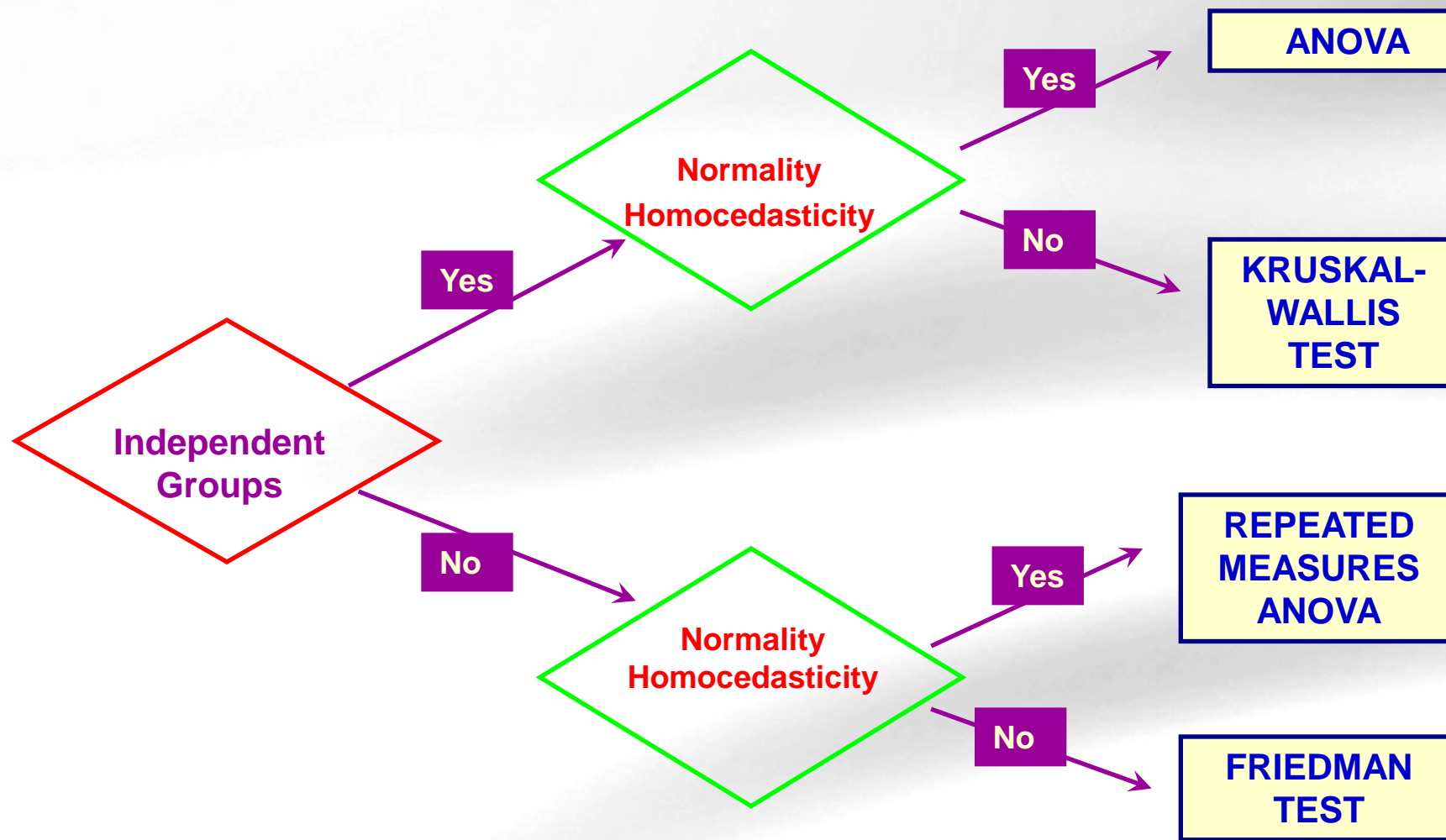


## Post-hoc ANOVA tests

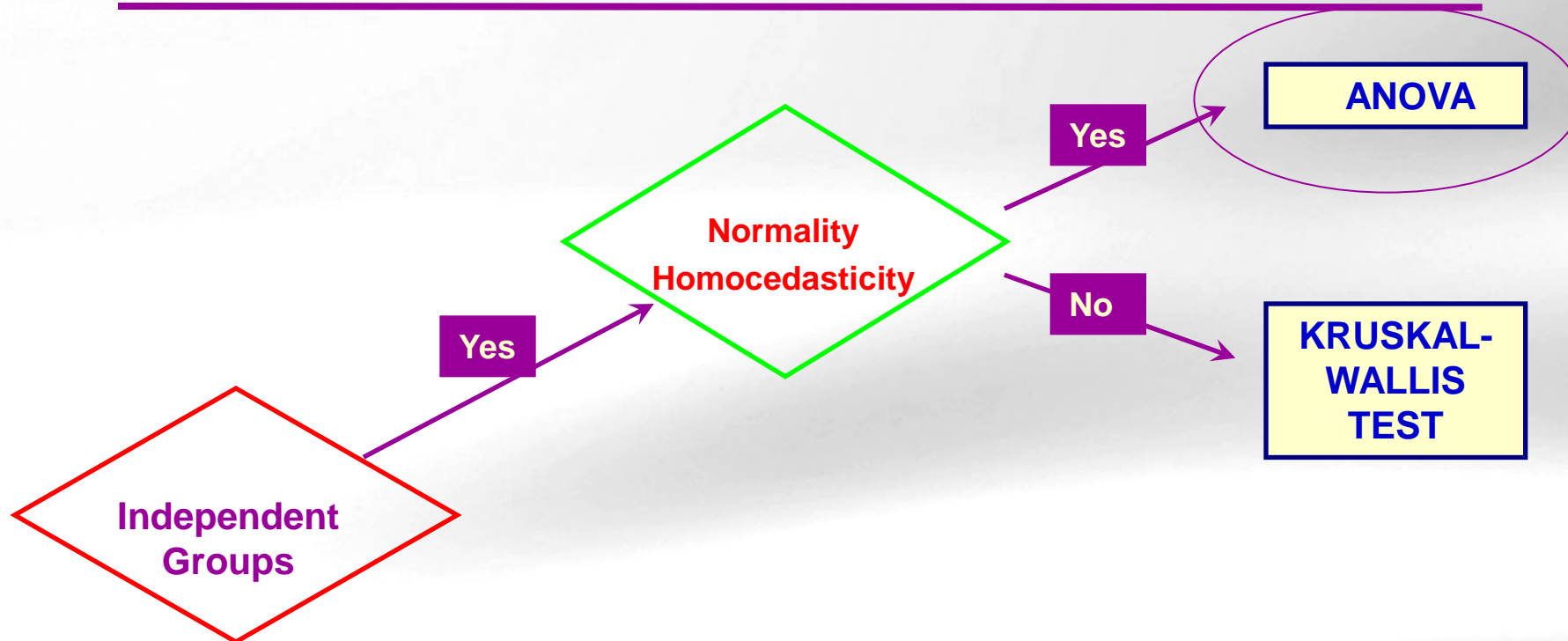
---

- If we wish to compare all means against all means the number of tests increases quickly (to compare all pairs of means if there are  $k$  groups  $(k*k-1)/2$  tests are required).
- This is usually called **multiple comparisons** and common methods of adjustment are Tukey, Fisher HSD or Bonferroni.

# Three or more groups



# Three or more groups





# Read diabetes data

```
library(readxl)
library(dplyr)
library(magrittr)
diabetes <- read_excel("datasets/diabetes.xls")
sapply(diabetes, class)
```

```
numpacie mort tempsviu edat bmi edatdiag
"numeric" "character" "numeric" "numeric" "numeric" "numeric"
tabac sbp dbp ecg chd
"character" "numeric" "numeric" "character" "character"
```

```
diabetes_factor <- diabetes %>%
 mutate_if(sapply(diabetes, is.character), as.factor) %>%
 select (-numpacie)
```

```
diabetes_factor %>%
 group_by(ecg) %>%
 summarise(n=n(),
 mean = mean(edat),
 sd=sd(edat))
```

```
A tibble: 3 x 4
ecg n mean sd
<chr> <int> <dbl> <dbl>
1 Anormal 11 64.9 6.76
2 Frontera 27 53.8 11.4
3 Normal 111 50.5 11.5
```

```
anova<-aov(edat~ecg,data=diabetes)
summary(anova)
```

```
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.
```

# Multicomparison

```
library(multcomp)
tuk <- glht(anova, linfct = mcp(ecg = "Tukey"))

print(summary(tuk)) # pairwise tests

##
Simultaneous Tests for General Linear Hypotheses
##
Multiple Comparisons of Means: Tukey Contrasts
##
##
Fit: aov(formula = edat ~ ecg, data = diabetes_factor)
##
Linear Hypotheses:
Estimate Std. Error t value Pr(>|t|)
Frontera - Anormal == 0 -11.094 4.010 -2.767 0.016496 *
Normal - Anormal == 0 -14.405 3.543 -4.065 0.000217 ***
Normal - Frontera == 0 -3.310 2.405 -1.376 0.345732

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Adjusted p values reported -- single-step method)
```



```
print(confint(tuk, level=0.95)) # confidence intervals
```

```

Simultaneous Confidence Intervals

Multiple Comparisons of Means: Tukey Contrasts

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

Quantile = 2.3459
95% family-wise confidence level

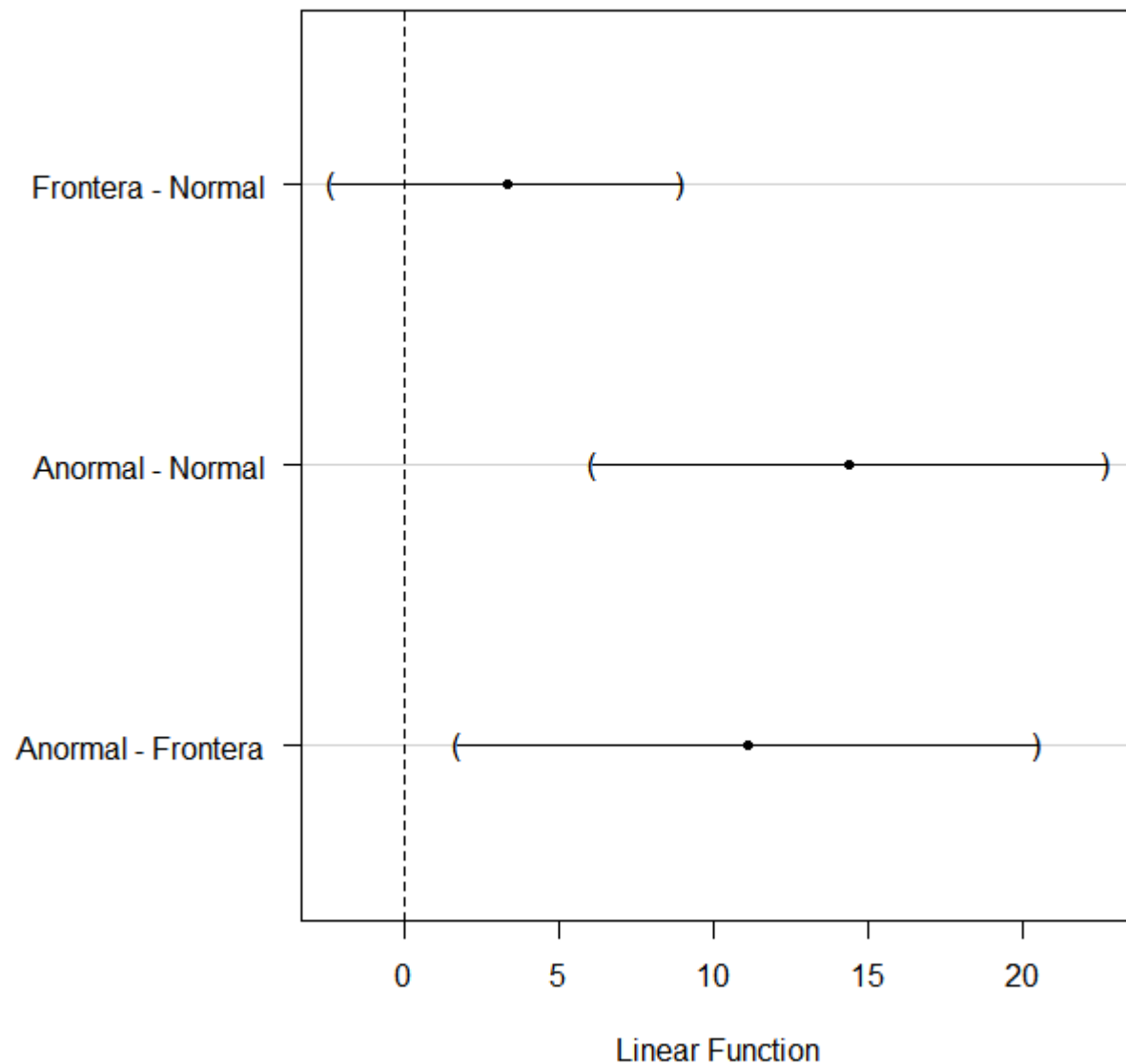
Linear Hypotheses:

Estimate lwr upr
Frontera - Anormal == 0 -11.0943 -20.5009 -1.6876
Normal - Anormal == 0 -14.4046 -22.7173 -6.0919
Normal - Frontera == 0 3.103 -8.9534 2.3328
```

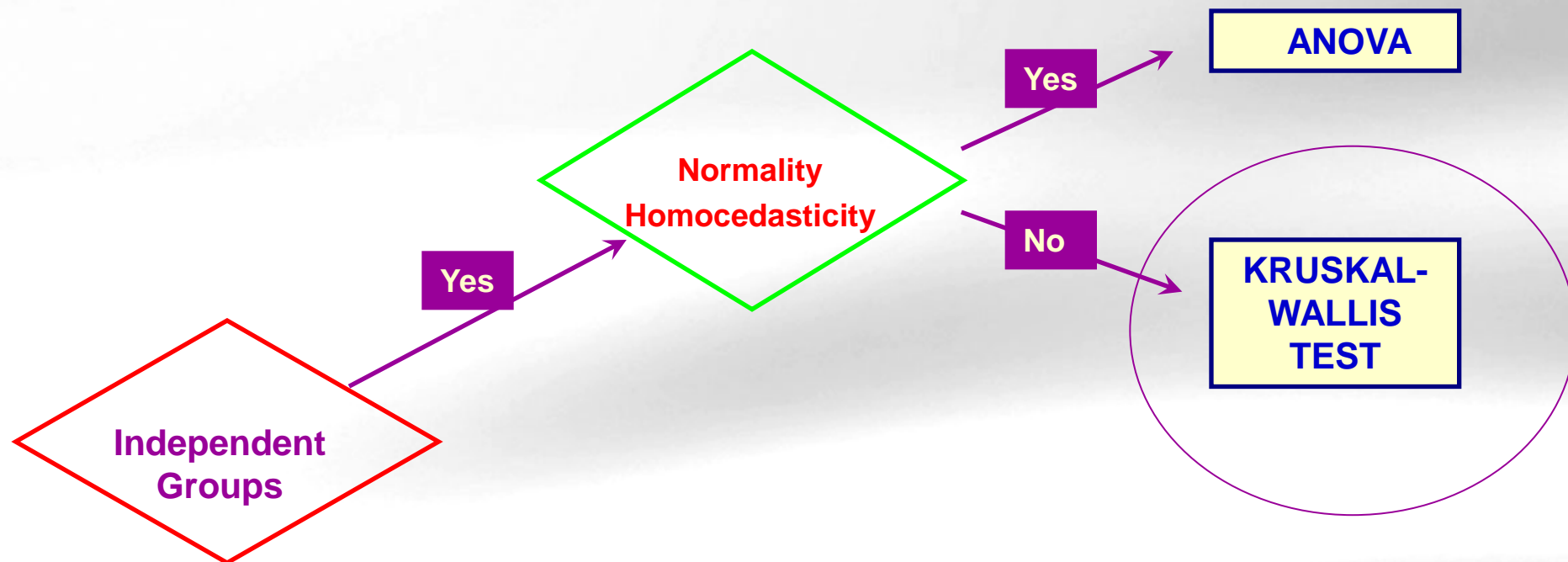
# Multicomparison

```
plot(confint(tuk))
```

95% family-wise confidence level



# Three or more groups



# Kruskal-Wallis Test

```
diabetes_factor%>%
 group_by(ecg) %>%
 summarise(median = median(edat))
```

```
A tibble: 3 x 2
ecg median
<fct> <dbl>
1 Anormal 64
2 Frontera 53
3 Normal 49
```

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

```
##
Kruskal-Wallis rank sum test
##
data: edat by ecg
Kruskal-Wallis chi-squared = 17.483, df = 2, p-value = 0.0001
```

## Dunn Test for multiple comparison

```
library(dunn.test)
with(diabetes_factor,dunn.test(edat,ecg,method="bonferroni"))
```

```
Kruskal-Wallis rank sum test
##
data: edat and ecg
Kruskal-Wallis chi-squared = 17.4826, df = 2, p-value = 0
```

```
##
##
Comparison of edat by ecg
(Bonferroni)
```

```
Col Mean-|
Row Mean | Anormal Frontera
-----+-----
```

Frontera	2.721182	
	0.0098*	
Normal	4.075469	1.467464
	0.0001*	0.2134

```
##
```



# Exercise 5

---

- Are there differences between systolic pressure (*sbp*) and *ECG* in diabetic patients?
  - a) For each variable, perform some descriptives and check normality assumptions
  - b) Set the hypothesis contrast and perform the test
  - c) If needed, apply a post-hoc test for multiple comparisons to test which groups are different.
  - d) **Extra:** Analyze the relation between the follow-up time (*tempsviu*) and the *ECG*.



# Exercise 6

---

*Indica que tipo de análisis o que pruebas estadísticas utilizarías en cada uno de los casos e indica también si es necesario hacer algún tipo de prueba adicional para llevar a cabo el análisis. Formula la hipótesis de trabajo de acuerdo con las hipótesis seleccionadas.*

**e) Se estudia un grupo de 33 pacientes afectados de Carcinoma hepatocelular, un grupo de 22 afectados únicamente de cirrosis y un grupo control de 31 donantes de sangre. Se determina la actividad celular NK y el número de células CK en los tres grupos. (Nota: la media de actividad celular es de 39 unidades líticas / $10^7$  de linfocitos y la mediana es de 28 y la media del número de células es de 178 y la mediana de 163)**