### "It's hard....."

#### **Previous Learning Experience Bloom's Taxonomy** Decreasing Produce new or original work create Design, assemble, construct, conjecture, develop, formulate, author, investigate level of Justify a stand or decision comfort evaluate appraise, argue, defend, judge, select, support, value, critique, weigh Draw connections among ideas differentiate, organize, relate, compare, contrast, distinguish, examine, analyze experiment, question, test **Data Analytics** Use information in new situations execute, implement, solve, use, demonstrate, interpret, operate, apply schedule, sketch Stage 1 Explain ideas or concepts understand Increasing classify, describe, discuss, explain, identify, locate, recognize, report, select, translate number of **Pre-university** Recall facts and basic concepts remember mistakes made define, duplicate, list, memorize, repeat, state

Vanderbilt University Center for Teaching

**17**C

# Laboratory & Professional Skills: Data Analysis

# Emma Rand Data Analysis in R

Week 5 Introduction to one- and two-sample tests. One sample tests

### Last week

- The normal distribution because it is the basis of many tests (parametric tests such as t-test, regression and ANOVA)
  - Properties of normal distributions
  - Sampling distribution of the mean and the standard error
  - Confidence intervals
- In RStudio
  - Calculate probabilities and quantiles from normal distributions
  - Calculate confidence intervals

## Summary of this week

#### We will cover

- General intro to one- and twosample tests
- one sample-tests
  - The one-sample t-test
  - The paired sample t-test
  - The one sample Wilcoxon

### R practice

- No workshop
- Activities to consolidate previous skills: Workflow basics and projects

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  - The one-sample t-test
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'Parametric' tests – Based on the normal distribution

### Summary of this week

- We will cover one sample-tests
  - The one-sample t-test
  - The paired sample t-test
  - The one sample Wilcoxon
     Non-parametric
- R practice
  - No workshop
  - Activities to consolidate previous skills: Workflow basics and projects

### Learning objectives

By actively following the material and carrying out the independent study the successful student will be able to:

- Appreciate that t-tests are based on the normal distribution and have assumptions relating to it
- Understand the principles of *t*-tests
- Select, appropriately, one-sample t-tests and their nonparametric equivalent (MLO 2)
- Recognise when two samples are not independent (MLO 2)
- Know what functions are used in R to run these tests and how to interpret them(MLO 3 and 4)
- Know how to state the results of these tests scientifically (MLO 3 and 4)

# Introduction to one- and twosample tests

T-tests and their non-parametric equivalents

# Reminder: The choice of test depends on ....

### 1. Type of data

The type of values a variable can take: <u>Discrete</u> or continuous?

### 2. Their role in the analysis

Which is the response and which is/are explanatory?

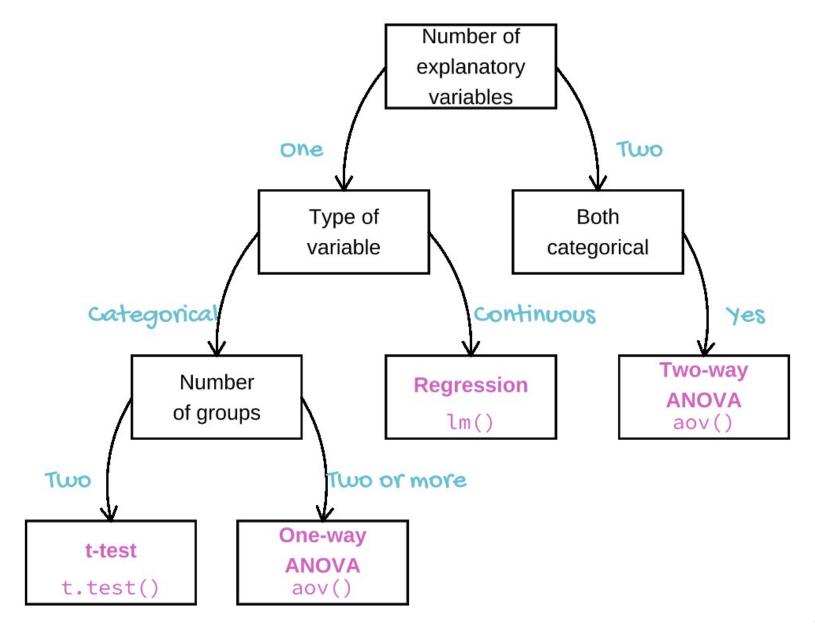
(week 3 Hypothesis testing, data types, reading data in to R and saving figures

# Choosing tests: 3 steps

- 1. What is a one sentence description of what you want to know?
- 2. What are your explanatory variables?
  - Categories: *t*-tests, ANOVA, Wilcoxon, Mann-Whitney
  - Continuous: Regression, correlation
- 3. What is your response variable?
  - Normally distributed: *t*-tests, ANOVA, regression
  - Counts: Chi-squared or stage 2 ©

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## Types of *t*-test

### 1. One-sample

- Compares the mean of sample to a particular value (compares the response to a reference)
  - Includes paired-sample test compares the mean difference to zero (i.e., compares dependent means)

### 2. Two-sample

Compares two (independent) means to each other

### t-tests in general

### Assumptions

All *t*-tests assume the "residuals" are normally distributed and have homogeneity of variance

A residual is the difference between the predicted and observed value

Predicted value is the mean / group mean

### *t*-tests in general: assumptions

## **Checking Assumptions**

- Common sense
  - Data should be continuous
  - No/few repeats
- Plot the residuals
- Using a test in R

### *t*-tests in general: assumptions

# When residuals are not normally distributed

- Transform (not really covered)
  - E.g. Log to remove skew
- Use a non-parametric test (covered)
  - Fewer assumptions
  - Generally less powerful

# The one-sample *t*-test

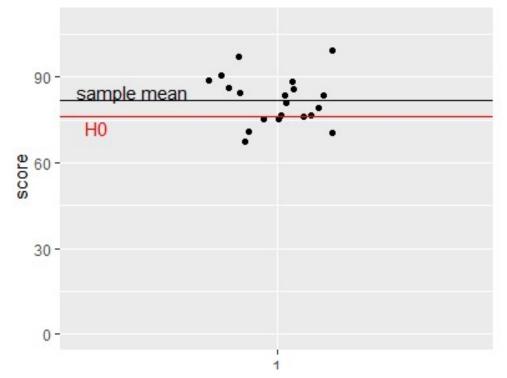
A parametric test

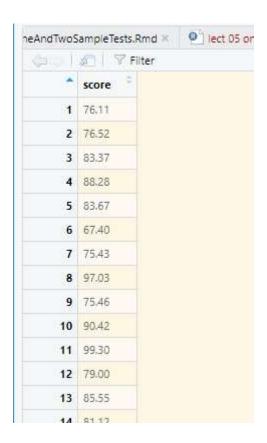
## One-sample *t*-tests

Tests whether the mean of a single sample differs from an expected value (i.e.,  $H_0$ )

- Example: Fields are sprayed if crop plants have a disease score\* of 76.
- 20 plants in a field are measured
- Is their mean significantly different from the reference of 76?

# One-sample t-tests - example





## One-sample t-tests - example

•  $H_0$ : mean = 76

• Standard formula for all t-tests  $t = \frac{statistic - hypothesised\ value}{s.e.\ of\ statistic}$ 

• d.f.= n - 1

## One-sample t-tests - example

$$t = \frac{\text{statistic} - \text{hypothesised value}}{\text{s.e. of statistic}}$$

$$\bar{x} = 81.8$$

$$\mu = 76.00$$

Is the difference between the obtained value and the expected value big relative to the variability?

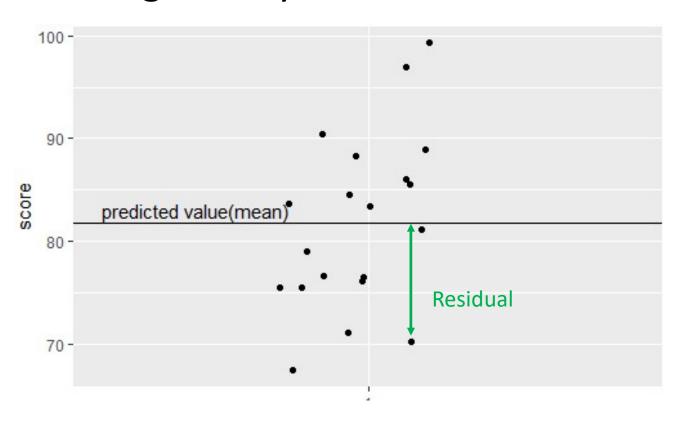
### One-sample *t*-tests - example

Note: you can also use

```
t.test(data = plants, score ~ 1, mu = 76)
```

## One-sample *t*-tests - example

Checking the assumptions: normally and homogenously distributed residuals



### One-sample *t*-tests - example

Checking the assumptions: normally and homogenously distributed residuals

```
residuals <- plants$score - mean(plants$score)
hist(residuals)
shapiro.test(residuals)

Shapiro-Wilk normality test

data: residuals
W = 0.9725, p-value = 0.8065

Histogram of residuals

**The state of the state o
```

### One-sample *t*-tests - example

Reporting the result: "significance of effect, direction of effect, magnitude of effect"

The disease score for plants in this field

$$(\bar{x} = 81.8)$$
 is significantly higher than

76 (
$$t = 2.52$$
;  $d.f. = 19$ ;  $p = 0.021$ ).

### One-sample *t*-tests - summary

- Parametric
- To test whether the mean of a single sample differs from an expected value
- t is size of difference relative to the s.e.
- Function in R: t.test(df\$response, mu = Exp)
- If p < 0.05 the test is significant
- assumptions: normally and homogenously distributed residuals
- Significance, direction, magnitude
- Figure: probably not needed

# The paired sample t-test

A parametric test

### Paired-sample *t*-tests

 Two samples but values are not independent (could not reorder)

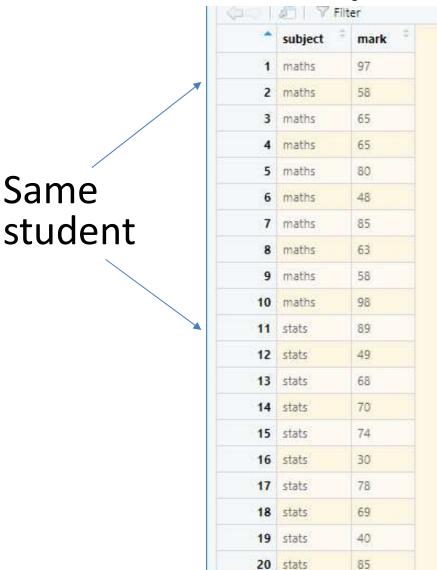
Actually a one-sample test

# Paired-sample *t*-tests example

Is there a difference between the maths and stats marks of 10 students?

The one sample is the difference between the pairs of values

n.b. tidy data



### Paired-sample t-tests - example

- $H_0$ : mean difference = 0
- Standard formula for all t-tests

$$t = \frac{statistic - hypothesised\ value}{s.e.\ of\ statistic}$$

• 
$$t_{[d.f]} = \frac{\bar{d}-0}{s.e. of \ \bar{d}}$$

• d.f.= n - 1 (where n is the number of pairs)

## Paired-sample *t*-tests

### Run paired sample *t*-test

```
t.test(data = marks, mark ~ subject, paired = TRUE)
        Paired t-test
data: mark by subject
t = 2.3399, df = 9, p-value = 0.04403
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.2159788 12.7840212
sample estimates:
mean of the differences
                   6.5
```

## Paired-sample t-tests - example

# Checking the assumptions: normally and homogenously distributed residuals

```
diffs <- marks$mark[marks$subject == "maths"] -</pre>
marks$mark[marks$subject == "stats"]
                                                  Histogram of residuals
residuals <- diffs - mean(diffs)</pre>
hist(residuals)
                                              Frequency
shapiro.test(residuals)
                                                 0
                                                 N
                                                 0.
                                                 0.0
Shapiro-Wilk normality test
                                                    -15
                                                        -5 0 5
                                                                 15
data:
       residuals
                                                        residuals
W = 0.91246, p-value = 0.2983
```

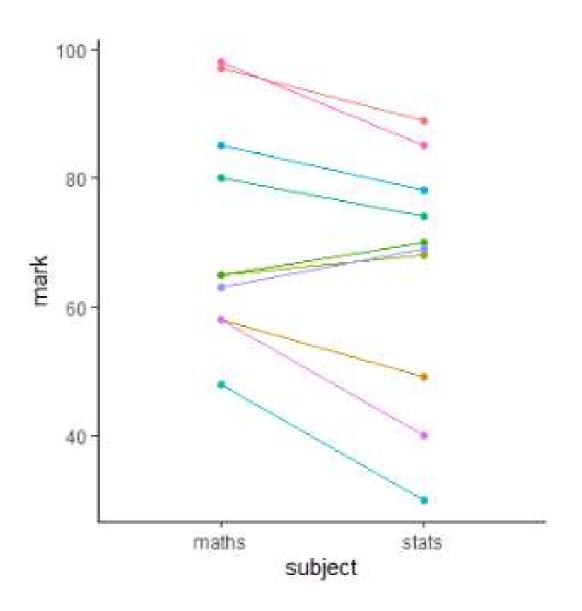
### Paired-sample *t*-tests

Reporting the result: "significance of effect, direction of effect, magnitude of effect"

Individual students score significantly higher in maths than in statistics (t = 2.34; d.f. = 9; p = 0.044) with an average difference of 6.5%.

*t*-tests

Paired-sample *t*-tests: figure



### Paired-sample *t*-tests - summary

- Parametric
- To test whether the mean difference between pairs of values is zero
- t is size of mean difference relative to the s.e.
- Function in R: t.test(data = df, response ~ explanatory, paired = TRUE)
- If p < 0.05 the test is significant
- assumptions: normally and homogenously distributed residuals
- Significance, direction, magnitude
- Figure: none or spaghetti plot

## The one sample Wilcoxon

Non-parametric equivalent of the paired-sample *t*-test

# When the *t*-test assumptions are not met: non- parametric tests

Non-parametric tests make fewer assumptions

Based on the ranks rather than the actual data

Null hypotheses are about the mean rank (not the mean)

### Non-parametric tests

### t-test equivalents

i,.e., the type of question is the same but the response variable is not normally distributed or it is impossible to tell (small samples)

- one sample *t*-test and paired-sample *t*-test:
   the one-sample Wilcoxon
- Two-sample t-test (next week): two-sample
   Wilcoxon aka Mann-Whitney

### Non-parametric tests

# one/paired-sample Wilcoxon

Marks – small sample.

### Wilcoxon might be more appropriate

```
wilcox.test(data = marks, mark ~ subject, paired = TRUE)

Wilcoxon signed rank test with continuity correction

data: mark by subject
V = 48.5, p-value = 0.03641
alternative hypothesis: true location shift is not equal to 0

Warning message:
In wilcox.test.default(x = c(97L, 58L, 65L, 65L, 80L, 48L, 85L, : cannot compute exact p-value with ties
```

# Non-parametric tests one/paired-sample Wilcoxon

Reporting the result: "significance of effect, direction of effect, magnitude of effect"

Individual students score significantly higher in maths than in statistics (Wilcoxon: V = 48.5; n = 10; p = 0.036) with a median difference of 7.5%.

### Wilcoxon- summary

- Non-parametric
- when assumptions for t-test not met
- To test whether the mean rank difference between pairs of values is zero
- Function in R:
  wilcox.test(data = df, response ~ explanatory, paired =
  TRUE)
- If p < 0.05 the test is significant
- Few assumptions
- Figure: none or spaghetti plot