Paired t-test

Data Analysis for Psychology in R 1

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Learning Objectives

- Understand when to use an paired sample *t*-test
- Understand the null hypothesis for an paired sample *t*-test
- Understand how to calculate the test statistic
- Know how to conduct the test in R
- Know how to calculate Cohen's *D* for each form of *t*-test

Topics for today

- Recording 1: Conceptual background and introduction to our example
- Recording 2: Calculations and R-functions
- Recording 3: Assumptions and effect size

Purpose & Data

- The paired sample *t*-test is used when we want to test the difference in mean scores for a sample measured at two points in time.
 - Thus this is a first example of a repeated measures design.
- Data Requirements
 - A continuously measured variable.
 - A binary variable denoting time.

Example

- I want to assess whether a time-management course helps reduce exam stress in students.
- I ask 50 students to take a self-report stress measure during their winter exams.
- At the beginning of semester 2 they take a time management course.
- I then assess their self-report stress in the summer exam block.
 - Let's assume for the sake of this example that I have been able to control the volume and difficulty of the exams the students take in each block.

Data

A tibble: 6 × 3 ## ID stress time <chr> <dbl> <fct> ## ## 1 ID1 14 t1 ## 2 ID2 7 t1 ## 3 ID3 8 t1 ## 4 ID4 8 t1 ## 5 ID5 7 t1 ## 6 ID6 7 t1

Calculating difference

• In the paired *t*-test, we specifically calculate and analyse the difference in scores at time 1 and time 2 per participant.

 $d_i = x_{i1} - x_{i2}$

Test statistic

• The resulting test statistic:

$$t=rac{ar{d}}{s_d/\sqrt{n}}$$

- where:
 - $\circ \ ar{d}$ = mean of the individual difference scores (d_i)
 - $\circ~~s_d$ = standard deviation of the difference scores (d_i)
 - \circ *n* = sample size
- The associated sampling distribution is a t-distributon with n-1 degrees of freedom.
 - Note, this is just essentially a one sample test on the difference scores.

Hypotheses

• Two-tailed:

 $egin{aligned} H_0: \mu_d = 0\ H_1: \mu_d
eq 0 \end{aligned}$

• One-tailed

 $egin{aligned} H_1:\mu_d < 0\ H_1:\mu_d > 0 \end{aligned}$

Time for a break

Welcome Back!

Let's calculate a paried-sample t-test!

Our Example

- I elect to use a two-tailed test with alpha of .01
- I want to be quite sure the intervention has worked and stress levels have changed.
- So my hypotheses are:

 $egin{aligned} H_0: \mu_d = 0\ H_1: \mu_d
eq 0 \end{aligned}$

Calculation

- Steps in my calculations:
 - Calculate the difference scores for individuals.
 - Calculate the mean of the difference scores.
 - Calculate the SD of the difference scores.
 - Check I know my N.
 - Calculate the standard error of the mean difference.
 - \circ Use all this to calculate t
 - Calculate my degrees of freedom

Data organisation

- Our data is currently in what is referred to as long format.
 - All the scores are in one column, with two entries per participant.
- To calcuate the d_i values, we will convert this to wide format.
 - Where there are two columns representing the score at time 1 and time 2
 - And a single row per person

Data organisation

```
## # A tibble: 6 × 3
##
    ID
             t1
                   t2
##
  <chr> <dbl> <dbl>
## 1 ID1
             14
                    7
                    7
## 2 ID2
              7
## 3 ID3
              8
                 9
## 4 ID4
              8
                 12
## 5 ID5
              7
                  10
## 6 ID6
              7
                    9
```

Calculation

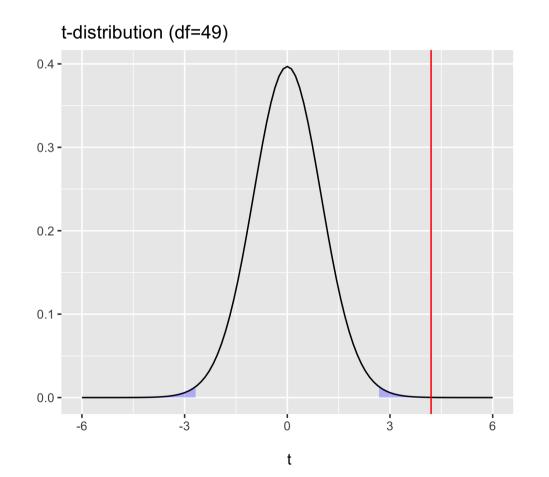
```
calc <- exam_wide %>%
  mutate(
    dif = t1 - t2) %>%
summarise(
    D = mean(dif),
    SDd = round(sd(dif),2),
    N = n()) %>%
mutate(
    SEd = round(SDd /sqrt(N),2),
    t = round(D/SEd,2)
)
```

A tibble: 1 × 5
D SDd N SEd t
<dbl> <dbl> <int> <dbl> <dbl> <dbl> <dbl> 4.2

Is my test significant?

- So we have all the pieces we need:
 - *t* = 4.2
 - $\circ df = n 1 = 49$
 - Hypothesis to test (two-tailed)
 - $\circ \ \alpha = 0.01$
- So now all we need is the critical value from the associated *t*-distribution in order to make our decision .

Is my test significant?



tibble(LowerCrit = round(qt(0.005, 49),2), UpperCrit = round(qt(0.995, 49),2), Exactp = round(2*(1-pt(calc[[5]], 49)),5)

##	#	A tibble:	1 × 3	
##		LowerCrit	UpperCrit	Exactp
##		<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
##	1	-2.68	2.68	0.00011

In R

```
res <- t.test(exam_wide$t1, exam_wide$t2,</pre>
        paired = TRUE,
        alternative = "two.sided")
##
##
       Paired t-test
##
## data: exam_wide$t1 and exam_wide$t2
## t = 4.1864, df = 49, p-value = 0.0001174
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 1.091937 3.108063
## sample estimates:
## mean of the differences
##
                       2.1
```

• Again, slight rounding differences.

Write-up

A paired-sample *t*-test was conducted in order to determine a if a statistically significant ($\alpha = .01$) mean difference in self-report stress was present, pre- and post-time management intervention in a sample of 50 undergraduate students. The pre-intervention mean score was higher (Mean=9.72) than the post intervention score (Mean = 7.62). The difference was statistically significant (t (49)= 4.19, p < .01, two-tailed). Thus, we reject the null hypothesis of no difference.

Time for a break

Welcome Back!

Now to check assumptions for a paired t-test and calculate Cohen's D

Assumption checks summary

	Description	One-Sample t- test	Independent Sample t- test	Paired Sample t-test
Normality	Continuous variable (and difference) is normally distributed.	Yes (Population)	Yes (Both groups/ Difference)	Yes (Both groups/ Difference)
Tests:	Descriptive Statistics; Shapiro-Wilks Test; QQ-plot			
Independence	Observations are sampled independently.	Yes	Yes (within and across groups)	Yes (within groups)
Tests:	None. Design issue.			
Homogeneity of variance	Population level standard deviation is the same in both groups.	NA	Yes	Yes
Tests:	F-test			
Matched Pairs in data	For paired sample, each observation must have matched pair.	NA	NA	Yes
Tests:	None. Data structure issue.			

Assumptions

- 1. Normality of the difference scores (d_i)
- 2. Independence of observations within group/time
- 3. Data are matched pairs (design)
- We will briefly show the normality assumptions again.
 O Hopefully these are becoming familiar.

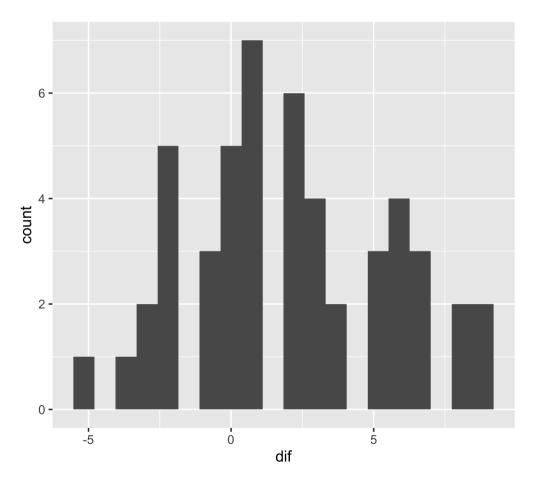
Adding the difference scores

- Our assumptions concern the difference scores.
- We showed these earlier in our calculations.
- Here we will add them to exam_wide for ease.

```
exam_wide <- exam_wide %>%
  mutate(
    dif = t1 - t2)
```

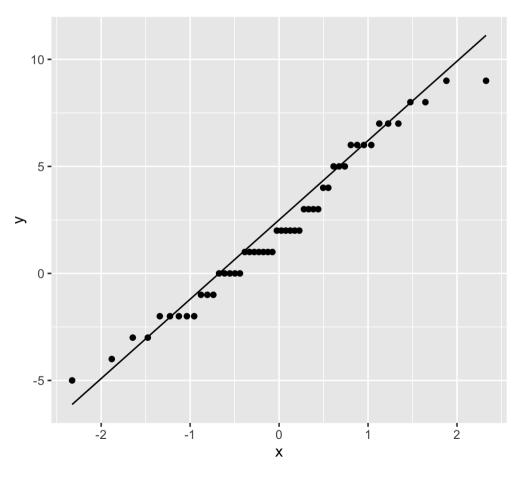
Histograms

exam_wide %>%
ggplot(., aes(x=dif)) +
geom_histogram(bins = 20)



QQ-plots

```
exam_wide %>%
ggplot(., aes(sample = dif)) +
stat_qq() +
stat_qq_line()
```



Shapiro-Wilks R

shapiro.test(exam_wide\$dif)

```
##
## Shapiro-Wilk normality test
##
## data: exam_wide$dif
## W = 0.97142, p-value = 0.264
```

- Fail to reject the null, *p* > .05
- Normality of the differences is met.

Cohen's D: Paired t

• Paired-sample t-test:

$$D=rac{ar{d}\,-0}{s_d}$$

- \bar{d} = mean of the difference scores (d_i)
- s_d = standard deviation of the difference scores (d_i)

Cohen's D in R

```
##
## Cohen's d
##
## d estimate: 0.8822584 (large)
## 95 percent confidence interval:
## lower upper
## 0.3893289 1.3751880
```

Summary: Three different t-tests

	One-sample	Independent Sample	Paired (Dependent) Sample
Outcome	Continuous Variable	Continuous Variable	Continuous Variable
Predictor	Single group vs population	Categorical: two groups	Categorical: two time points
Sample	One sample vs population value	Two independent groups	One group sampled at two time points
Measure of difference	Observed - known population value	Group 1 - Group 2	Time 1 - Time 2
Measure of Variability	Standard error of the mean	Pooled standard error of difference in means	Standard error of the difference in means