

Lab 5 Tutorial

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Dependent and Independent Samples t-test

Intro

This week we will extend our exploration of t-tests by conducting dependent and independent samples t-tests. Using a hypothetical dataset and research hypothesis, we will walk through how to conduct t-tests by using the `oneSampleTTest()` function.

Note: **All material is taken from Dr. Zopluoglu's 2021 EDUC 614 materials and adapted for this year's course.**

We have adapted it to slide format using R.

Want to learn how we created R slide presentations? This linked tutorial is helpful.

Independent samples t-test

Importing Data

Our first step is to import the data we need. We'll use the Add dataset for this walkthrough, which you can access via Canvas. We'll also use the `head()` function to familiarize ourselves with the structure of the data.

```
Add <- read.csv(file = "/Users/aguha/Downloads/Add.csv",  
                 header = TRUE)
```

```
head(Add, 3) #this pulls up the first 3 lines of data
```

	##	CaseNum	ADDSC	Sex	Repeat	IQ	GPA	SocProb	Dropout
##	1	1	45	1	0	111	2.60	0	0
##	2	2	50	1	0	102	2.75	0	0
##	3	3	49	1	0	108	4.00	0	0

Research Hypothesis

The column ADDSC is a score for each observation related to the level of Attention Deficit Disorder (ADD). Higher scores indicate more severity. The column Repeat is a binary variable with 0 indicating that a student did not repeat a grade and 1 indicating that a student did repeat a grade.

Research Hypothesis: Students who do not repeat a grade have lower scores of ADD than the students who repeat a grade.

Using a verbal language, we can also write the null hypothesis and alternative hypothesis. . .

Null and Alternative Hypothesis

- ▶ **Null hypothesis:** The average ADD scores for the students who repeat a grade are equal to the average ADD scores for the students who does not repeat a grade.
- ▶ **Alternative hypothesis:** The average ADD scores for the students who do not repeat a grade are lower than the ADD scores for the students who repeat a grade.

You can write same null and alternative hypothesis by using symbolic notation. Suppose that μ_2 is the population mean for students who repeated a grade, and μ_1 is the population mean for students who did not repeat a grade.

$$H_0 : \mu_1 = \mu_2$$

$$H_a : \mu_1 < \mu_2$$

Note that we will be conducting a one-tailed t-test because the alternative hypothesis implies a specific direction for the difference.

One-Sided Independent samples t-test using the `independentSamplesTTest()` function

Also from the `{lsr}` package, we can use the `independentSamplesTTest()` function to conduct an independent samples t-test using the following arguments:

- ▶ `formula`: `outcome variable(left) ~ independent variable(right)`
 - ▶ you are looking at the outcome variable for different levels of the independent variable
 - ▶ variable names must match those in your dataset
- ▶ `data`: name of your dataset
- ▶ `one.sided`: specify the name of the group that you're expecting to have the higher score
- ▶ `var.equal`: whether or not you are making the assumption that both populations have the same variance
 - ▶ if you can reasonably assume that this assumption holds, set to `TRUE`
 - ▶ otherwise set to `FALSE` to conduct Welch's test

Data Checks

Let's run the independent samples t-test for the same null and alternative hypothesis specified earlier. But, first we will do some checks on the variables of interest in this dataset.

```
# First, make sure to convert your independent variable  
# to a factor variable if it isn't already.  
typeof(Add$Repeat)
```

```
## [1] "integer"
```

```
Add$Repeat <- as.factor(Add$Repeat)
```

```
# Second, make sure your independent variable has  
## only two distinct values  
table(Add$Repeat)
```

```
##  
##  0  1  
## 76 12
```

```
# Third, make sure your dependent variable is numeric  
typeof(Add$ADDSC)
```

```
## [1] "integer"
```

Descriptives

It is also a good idea to check the descriptive statistics for the variable to make sure the numbers look as expected.

```
require(psych)
# Overall descriptives for the dependent variable
describe(Add$ADDSC)
```

```
##      vars   n mean    sd median trimmed   mad min max range
## X1      1  88 52.6 12.42    50   52.18 10.38  26  85    59
```

```
# Conditional descriptive statistics by Repeat  
describeBy(Add$ADDSC,Add$Repeat)
```

```
##
```

```
## Descriptive statistics by group
```

```
## group: 0
```

```
##      vars  n  mean      sd median trimmed  mad min max range
```

```
## X1      1 76 50.62 11.59      50   50.08 8.9  26  85    59
```

```
## -----
```

```
## group: 1
```

```
##      vars  n  mean      sd median trimmed  mad min max range
```

```
## X1      1 12 65.17 10.26     66.5    64.8 6.67  49  85    36
```

Conducting the t-test assuming variances are equal

Now, we can run the t-test.

```
require(lsr)
independentSamplesTTest(formula    = ADDSC ~ Repeat,
                        data       = Add,
                        var.equal  = TRUE,
                        one.sided = 0)
```

##

Student's independent samples t-test

##

Outcome variable: ADDSC

Grouping variable: Repeat

##

Descriptive statistics:

0 1

mean 50.618 65.167

std dev. 11.590 10.259

##

Hypotheses:

null: population means equal for both groups

alternative: different population means in each group

##

Test results:

t-statistic: -4.098

degrees of freedom: 86

p-value: <.001

##

As you see, this function returns an output that includes all the relevant numbers.

- ▶ t-statistic is reported as -4.098
- ▶ df is reported as 86
- ▶ p-value is reported as $<.001$
- ▶ sample means for each group, 50.62 and 65.17
- ▶ Cohen's d is 1.273
- ▶ CI is [-21.605, -7.491]

Summarizing Results

Finally, you can summarize the results using a paragraphs as the following:

The average ADD scores was 50.62 with a standard deviation of 11.59 for 76 students who did not repeat a grade and was 65.17 with a standard deviation of 10.26 for 12 students who repeated a grade. An independent samples t-test was conducted to test whether or not the average ADD score for students who did not repeat a grade was significantly lower than the students who repeated a grade. The findings indicates that the average ADD score for students who did not repeat a grade was 14.55 points lower, and the difference was statistically significant, $t(86) = -4.098$, $p < 0.001$, Cohen's $d = 1.27$.

Conducting the t-test assuming variances are NOT equal

If you think the equal variance assumption is not plausible, then you can also run the t-test with Welch approximation by specifying `var.equal = FALSE`.

```
independentSamplesTTest(formula = ADDSC ~ Repeat,  
                          data     = Add,  
                          var.equal = FALSE,  
                          one.sided = 0)
```

##

Welch's independent samples t-test

##

Outcome variable: ADDSC

Grouping variable: Repeat

##

Descriptive statistics:

0 1

mean 50.618 65.167

std dev. 11.590 10.259

##

Hypotheses:

null: population means equal for both groups

alternative: different population means in each group

##

Test results:

t-statistic: -4.482

degrees of freedom: 15.786

p-value: <.001

##

Summarizing Results

If you do run the t-test with Welch approximation, you can report the numbers accordingly. Notice that everything in the following write up is same except the numbers related to the t-statistic.

The average ADD scores was 50.62 with a standard deviation of 11.59 for 76 students who did not repeat a grade and was 65.17 with a standard deviation of 10.26 for 12 students who repeated a grade. An independent samples t-test was conducted to test whether or not the average ADD score for students who did not repeat a grade was significantly lower than the students who repeated a grade. The findings indicates that the average ADD score for students who did not repeat a grade was 14.55 points lower, and the difference was statistically significant, $t(15.79) = -4.482$, $p < 0.001$, Cohen's $d = 1.33$.

Dependent Samples t-test

Import the data

```
anorexia <- read.csv(file = "/Users/aguha/Downloads/anorexia.csv",  
                     header = TRUE)
```

```
str(anorexia) #this shows an overview of the data
```

```
## 'data.frame':    17 obs. of  3 variables:  
## $ id           : int  1 2 3 4 5 6 7 8 9 10 ...  
## $ Weight_before: num  83.8 83.3 86 82.5 86.7 79.6 76.9 ...  
## $ Weight_after : num  95.2 94.3 91.5 91.9 100.3 ...
```

- ▶ 17 patients diagnosed with anorexia participated in a study.
- ▶ The weight was measured before the family therapy (Weight_before)
- ▶ Following a 4 weeks of therapy, the weight was measured again for the same 17 subjects(Weight_after)

Note the data structure: Each individual is represented in a separate row. Dependent measures for the same individual are in columns.

Research Hypothesis

Research Hypothesis: Participants in the study will gain weight after being exposed to the 4-week family therapy.

- ▶ **Null hypothesis:** The average weight for patients before the therapy is equal to the average weight after the therapy.
- ▶ **Alternative hypothesis:** The average weight for patients before the therapy is smaller than the average weight after the therapy.

You can write same null and alternative hypothesis by using symbolic notation. Suppose that μ_1 is the population mean weight for patients before exposed to the therapy, and μ_2 is the population mean weight for patients after being exposed to the therapy.

$$H_0 : \mu_1 = \mu_2$$

$$H_a : \mu_1 < \mu_2$$

Note that we will be conducting a one-tailed t-test because the alternative hypothesis implies a specific direction for the difference.

Describe Data

```
describe(anorexia$Weight_before)
```

```
##      vars   n  mean    sd median trimmed  mad   min   max range  
## X1      1 17 83.23 5.02   83.3   83.15 4.15 73.4 94.2  20.8
```

```
describe(anorexia$Weight_after)
```

```
##      vars   n  mean    sd median trimmed  mad   min   max range  
## X1      1 17 90.49 8.48   92.5   90.77  4 75.2 101.6  26.4
```

Dependent samples t-test using the `oneSampleTTest()` function

Because of the data structure of dependent samples data (the same person is being used at two time points, for example), we can create another variable in the data set that computes the differences at each data point.

```
anorexia$Weight_difference <-  
  anorexia$Weight_after - anorexia$Weight_before
```

Using this new variable, we can conduct the `oneSampleTTest()` like before.

To conduct a one-sample t-test we need the following arguments, adopted for dependent samples:

- ▶ `x`: the column in your data for the variable you just created (difference variable)
- ▶ `one.sided`: this argument can take one of two forms depending on the direction of your alternative hypothesis:
 - ▶ If your alternative hypothesis is non-directional (i.e., two-sided) you'll use `FALSE`
 - ▶ If your alternative hypothesis implies a certain direction you'll use either `greater` or `less` depending on the direction
- ▶ `mu`: the value in your null hypothesis that you are testing against
 - ▶ will usually be 0 since you are seeing whether the groups are different

Dependent samples t-test

```
oneSampleTTest(x = anorexia$Weight_difference,  
               mu = 0,  
               one.sided = "greater") #since you are seeing
```

```
##
##      One sample t-test
##
## Data variable:   anorexia$Weight_difference
##
## Descriptive statistics:
##           Weight_difference
##      mean                7.265
##      std dev.            7.157
##
## Hypotheses:
##      null:           population mean less than or equal to 0
##      alternative: population mean greater than 0
##
## Test results:
##      t-statistic:   4.185
##      degrees of freedom: 16
##      p-value:   <.001
##
## Other information:
```

Summarizing Results

Summarize the results using a paragraphs as the following:

The average weight for 17 patients diagnoses with anorexia was 83.23 with a standard deviation of 5.02 before the family therapy, and the average weight increased to 90.49 with a standard deviation of 8.48 after four weeks of therapy. A dependent samples t-test was conducted to test whether or not the average weight gain after the family therapy was significant. The findings indicated that the patients increased their weight by 7.26 lbs, and the increase was statistically significant, $t(16) = 4.185$, $p < .001$, Cohen's $d = 1.01$.