# **Chapter 4: Basic Analyses**

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Introduction

T-tests

**ANOVA** 

Linear Regression

# Introduction

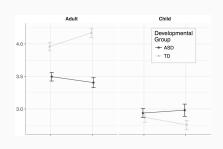
#### **Basic Analyses**

Basic Analyses: The analyses taught in the first stats course

#### These include:

- 1. T-tests
- 2. ANOVA
- 3. Linear Regression

These allow us to assess relationships like that in the figure.



Maybe surprising:\ These all are doing essentially the same thing!

First, **T-TESTS!** 

# **T**-tests

### Three Types

- 1. Simple
- 2. Independent Samples
- 3. Paired Samples

### Three Types

# Each will be demonstrated using:

```
Α
    0 -0.888158433 0.294452230
    1 0.384032654 -2.022480886
3
    0 -0.978200548  0.363196635
    0 0.597665769 0.306536631
4
5
    0 0.849400438 -0.444227641
6
    0 -0.890268979 -1.254064551
    0 -0.854688613 0.938866598
8
    0 -0.148777057 -2.283803888
9
      1.046238407 -0.149862421
```

#### Simple

t.test(df\$B, mu = 0)

Comparing a mean of a variable with  $\mu$ .

```
One Sample t-test
data: df$B
t = 0.29332, df = 99, p-value = 0.7699
alternative hypothesis: true mean is not equal to 0
95 percent confidence interval:
-0.1803399 0.2429081
sample estimates:
mean of x
0.03128405
```

#### **Independent Samples**

Comparing the means of two groups (dfA is the grouping variable).

```
t.test(df$B ~ df$A)
```

Welch Two Sample t-test

mean in group 0 mean in group 1 0.08514805 -0.04309956

```
data: df$B by df$A
t = 0.59367, df = 89.632, p-value = 0.5542
alternative hypothesis: true difference in means is not equal to
95 percent confidence interval:
  -0.3009458   0.5574410
sample estimates:
```

#### **Paired Samples**

Comparing repeated measures (e.g., Pretest vs. Posttest).

```
t.test(df$B, df$C, paired = TRUE)
```

Paired t-test

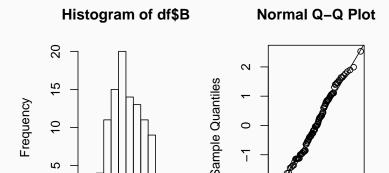
0.2401172

```
data: df$B and df$C
t = 1.7086, df = 99, p-value = 0.09066
alternative hypothesis: true difference in means is not equal to
95 percent confidence interval:
  -0.03873655   0.51897089
sample estimates:
mean of the differences
```

### **Testing Assumptions of T-Tests**

T-tests require that the data be normally distributed with approximately the same variance.

```
## Normality
par(mfrow = c(1,2))
hist(df$B)
qqnorm(df$B)
abline(a=0, b=1)
```



# **ANOVA**

#### **Analysis of Variance**

The Analysis of Variance (ANOVA) is highly related to t-tests but can handle 2+ groups.

- 1. Provides the same p-value as t-tests
- 2.  $t^2 = F$

For example:

```
fit_ano = aov(df$B ~ df$A)
summary(fit_ano)
```

```
t.test(df$B ~ df$A)$p.value
```

[1] 0.5542261

### **Analysis of Variance**

```
fit_ano = aov(df$B ~ df$A)
summary(fit_ano)
t.test(df$B ~ df$A)$p.value
```

#### Notice in the code:

- We assigned the aov() the name fit\_ano (which we could have called anything)
- We used the summary() function to see the F and p values.
- We pulled the p-value right out of the t.test() function.

- 1. One-Way
- 2. Two-Way (Factorial)
- 3. Repeated Measures
- 4. A combination of Factorial and Repeated Measures

#### **Types**

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We will use the following data set for the examples:

1 -0.825585271 -1.068904014 4

```
library(tidyverse)
df <- data.frame("A"=sample(c(0,1), 100, replace = TRUE) %>% fac
              "B"=rnorm(100),
              "C"=rnorm(100),
              "D"=sample(c(1:4), 100, replace = TRUE) %>% fac
df
   Α
                         C D
   1 0.393203143 0.166533149 3
   1 0.325532077 0.121702597 4
   3
   4
5
   0 -1.040692589 -0.353863739 1
6
   1 1.086275696 -0.249269105 2
   0 -0.304352002 -1.226131107 2
```

#### One-Way

A One-Way ANOVA can be run using aov().

```
fit1 = aov(B ~ D, data = df)
summary(fit1)
```

```
Df Sum Sq Mean Sq F value Pr(>F)
D 3 1.55 0.5152 0.499 0.684
Residuals 96 99.12 1.0325
```

#### Two-Way

A Two-Way ANOVA uses essentially the exact same code with a minor change—including the other variable in an interaction.

```
fit2 = aov(B ~ D * A, data = df)
summary(fit2)
```

	Df	Sum Sc	Mean	Sq F	value	Pr(>F)
D	3	1.55	0.51	52	0.502	0.682
A	1	1.28	1.28	18	1.250	0.266
D:A	3	3.50	1.16	79	1.139	0.338
Residuals	92	94.34	1.02	54		

The D:A line highlights the interaction term whereas the others show the main effects.

#### **Repeated Measures**

To show this, we will add a fake ID variable to our already fake data set df.

```
df$ID = 1:100
```

And change our data to long (Can you remember how to do it?)

```
library(tidyverse)
df_long = gather(df, "var", "value", 2:3)
df_long
```

```
A D
      ID var
             value
   1 3 1
           B 0.393203143
 1 4 2 B 0.325532077
3
   0 3 B -0.716154628
   0 2 4 B -0.544302202
4
5
   0 1 5 B -1.040692589
6
   1 2
        6
           B 1.086275696
           B -0.304352002
```

### Repeated Measures

The repeated measures, besides using a long-form of the data, is very similar in code. In addition to our usual formula (e.g., something ~ other + stuff), we have the Error() function. This function tells R how the repeated measures are clustered. In general, you'll provide the subject ID. The next slide highlights this.

#### **Repeated Measures**

```
fit3 = aov(value ~ var + Error(ID), data = df_long)
summary(fit3)
```

```
Error: ID

Df Sum Sq Mean Sq F value Pr(>F)

Residuals 1 2.222 2.222

Error: Within

Df Sum Sq Mean Sq F value Pr(>F)

var 1 1.95 1.9474 2.101 0.149

Residuals 197 182.57 0.9267
```

Here, value was the value of the repeated measures where var is the time. That means our oucome is testing if there were any differences from pre-test to post-test across all the groups.

#### **Combination**

To take the repeated measures a step further, we can do a Three-Way Repeated Measures ANOVA.

```
fit4 = aov(value ~ var * D * A + Error(ID), data = df_long)
summary(fit4)
```

The output is on the next slide. . .

#### **Combination**

Error: ID

var:A

var:D:A

D:A

```
Df Sum Sq Mean Sq
D 1 2.222 2.222

Error: Within

Df Sum Sq Mean Sq F value Pr(>F)

var 1 1.95 1.9474 2.088 0.150
D 3 1.43 0.4780 0.512 0.674
A 1 0.76 0.7593 0.814 0.368

var:D 3 1.26 0.4202 0.450 0.717
```

3 2.72 0.9055

Residuals 183 170.70 0.9328

1 0.66 0.6576 0.705 0.402

3 5.04 1.6794 1.800 0.149

0.971 0.408

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### **Checking Assumptions**

Of course, as with any statistical analysis, there are assumptions.

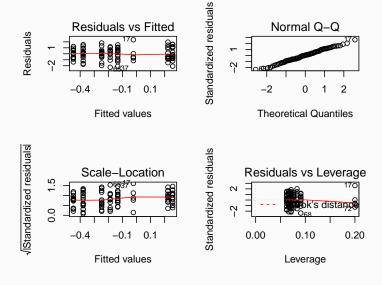
Many of these we can test.

Using our fitX objects from our ANOVAs above, we can look at our assumptions:

```
par(mfrow = c(2,2))
plot(fit2)
```

Again, the output is on the next slide. . .

## **Checking Assumptions**



### **Checking Assumptions**

They don't fit great on the slides but trust me that normality looks good. The assumption of homogeneity of variance looks good as well.

But, if you wanted to test it, you could.

```
library(car)
leveneTest(fit2)
```

```
Levene's Test for Homogeneity of Variance (center = median)

Df F value Pr(>F)
group 7 0.2113 0.9821
92
```

#### Large p-value here is a good thing: emo::ji("smile") 1

 $<sup>^{1}\</sup>mathrm{This}$  shows a smiley in 'R', just not on these slides—from the 'emo' package on GitHub.

# **Linear Regression**

## **Linear Regression**

Once again, linear regression is essentially the more flexible twin of ANOVA and t-tests.<sup>2</sup>

#### It can:

- 1. Handle continuous and categorical predictors (i.e., independent variables)
- 2. Less stringent assumption of equality of variances
- Is what many other methods are built on (Chapter 5 and 6 will talk about some of these)

 $<sup>^2</sup>$ It mainly only differs from ANOVA in the way it takes a dummy code rather than an effect code of the categorical variables.

### Linear Regression

We will use lm() (Linear Model) to fit these models.

```
fit5 = lm(B ~ A, data = df)
summary(fit5)
```

```
Call:
```

lm(formula = B ~ A, data = df)

#### Residuals:

Min 1Q Median 3Q Max -2.2232 -0.5694 -0.0839 0.6275 2.5542

#### Coefficients:

Residual standard error: 1.01 on 98 degrees of freedom

