Statistical inference and linear models

February 20, 2018

If you get bored

- Go to the last slide for bonus exercises
- Work on code for your research and ask question during exercise time
- But try and keep an eye out, just in case



Disclaimer

- Assume you got lectures about statistics and
 - know why we need statistics
 - have heard of the general philosophy
- We may simplify to focus on practical aspects
- Correct us if we say something completely awful

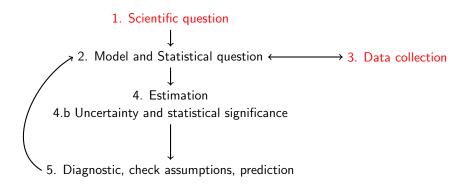
- Statistical inference
- 2 t-test, ANOVA, regression: all is one, one is all

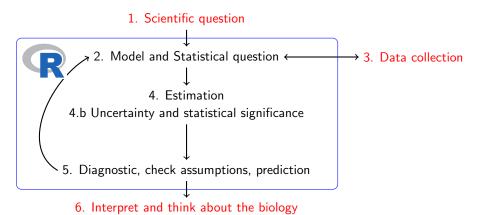
1. Scientific question

- 1. Scientific question
 - \downarrow
- 2. Model and Statistical question

- 1. Scientific question
- 2. Model and Statistical question ← → 3. Data collection

- 1. Scientific question
- 2. Model and Statistical question ← → 3. Data collection
 - 4. Estimation
- 4.b Uncertainty and statistical significance





6 / 19

```
data("iris")
```

```
str(iris)
plot(iris)
```

Scientific question: Are the taxa "setosa" and "versicolor" different species?

```
data("iris")
```

```
str(iris)
plot(iris)
```

- Scientific question: Are the taxa "setosa" and "versicolor" different species?
- Model and stat question:
 - ► Model:
 - * There is an intrinsic/expected sepal length value for a species; an individual value is the sum of this expectation and a random Gaussian deviation.
 - * $y_i = \mu_{species_i} + \epsilon_i$ with $\epsilon \sim N(0, \sigma^2)$
 - ★ t-test

```
data("iris")
```

```
str(iris)
plot(iris)
```

- Scientific question: Are the taxa "setosa" and "versicolor" different species?
- Model and stat question:
 - Model:
 - * There is an intrinsic/expected sepal length value for a species; an individual value is the sum of this expectation and a random Gaussian deviation.
 - * $y_i = \mu_{species} + \epsilon_i$ with $\epsilon \sim N(0, \sigma^2)$
 - ★ t-test
 - Statistical question:
 - Does sepal length differ significantly between the two taxa in our sample?
 - * Is the observed difference between taxa likely if both taxa have the same intrinsic/expected value?
- Data collection



One t-test for sepal length between setosa and versicolor:

```
t.test(x = iris$Sepal.Length[iris$Species == "setosa"],
        y = iris$Sepal.Length[iris$Species == "versicolor"])
Welch Two Sample t-test
data: iris$Sepal.Length[iris$Species == "setosa"] and iris$Sepal.Le
t = -10.521, df = 86.538, p-value < 2.2e-16
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-1.1057074 - 0.7542926
sample estimates:
mean of x mean of y
    5.006 5.936
```

- Statistical estimation
 - a Estimation
 - \star Cannot know true difference $\mu_{species_1} \mu_{species_2}$
 - ★ Estimated difference = Mean₁ Mean₂
 - ★ Difference contains random variation
 - b Quantify uncertainty / Statistical significance
 - * $t = \frac{\text{Mean}_1 \text{Mean}_2}{\text{Variation}} \frac{\sqrt{\text{Sample Size}}}{\sqrt{2}}$
 - * We know exactly how t is distributed when $\mu_{species_1} \mu_{species_2} = 0$
 - * Hence we know probability of $\geq t$ if $\mu_{species_1} \mu_{species_2} = 0$
 - Can derive confidence interval and standard error

- Statistical estimation
 - a Estimation
 - \star Cannot know true difference $\mu_{\mathit{species}_1} \mu_{\mathit{species}_2}$
 - ★ Estimated difference = Mean₁ Mean₂
 - * Difference contains random variation
 - b Quantify uncertainty / Statistical significance

```
\star t = \frac{\text{Mean}_1 - \text{Mean}_2}{\text{Variation}} \frac{\sqrt{\text{Sample Size}}}{\sqrt{2}}
```

- * We know exactly how t is distributed when $\mu_{species_1} \mu_{species_2} = 0$
- * Hence we know probability of $\geq t$ if $\mu_{species_1} \mu_{species_2} = 0$
- * Can derive confidence interval and standard error

Less uncertainty with

- Larger absolute difference
- Smaller variability
- Larger sample size

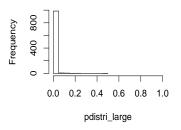


1. Larger absolute difference

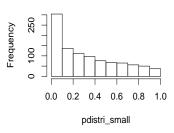
```
nbsim <- 1000
pdistri_large <- vector(length = nbsim)</pre>
pdistri_small <- vector(length = nbsim)</pre>
for (i in 1:nbsim)
  x1 \leftarrow rnorm(n = 10, mean = 2, sd = 1)
  x2 \leftarrow rnorm(n = 10, mean = 4, sd = 1) #large diff
  x3 \leftarrow rnorm(n = 10, mean = 2.5, sd = 1) #small diff
  out_large <- t.test(x1, x2)</pre>
  out_small <- t.test(x1, x3)</pre>
  pdistri_large[i] <-out_large$p.value
  pdistri_small[i] <-out_small$p.value
```

```
par(mfrow=c(1,2), cex=2)
hist(pdistri_large, xlim=c(0,1),
    main=paste("Prop signif=",mean(pdistri_large<0.05)))
hist(pdistri_small, xlim=c(0,1),
    main=paste("Prop signi=",mean(pdistri_small<0.05)))</pre>
```

Prop signif= 0.986



Prop signi= 0.184



When do we know it is different? Try it!

Exercise

Check the effect of smaller variability and/or larger sample size.

By the way, what are these p-values?

Blabla

Properties

- Reference to a null-model (H_0) with assumptions
- Uniform distribution under H_0 ...
- ... hence proportion(significance under H_0) = significance threshold

T-test exercise

```
t.test(x = ..., y=..., var.equal = TRUE)
t.test(x = ..., y=..., var.equal = FALSE)
```

What if variance are different by chance only?

```
set.seed(1234)
var(rnorm(20, mean = 0, sd = 1))
[1] 1.027806
var(rnorm(20, mean = 0, sd = 1))
[1] 0.6265501
```

Exercise

What option is more correct for var.equal?

- Statistical inference
- 2 t-test, ANOVA, regression: all is one, one is all

A small example

Animal behavior in response to weather Load data:

```
getwd()
setwd()
```

```
dat.behav <- read.csv(file = "datbehav.csv") # path to file</pre>
```

A small example

Animal behavior in response to weather Load data:

```
getwd()
setwd()
```

```
dat.behav <- read.csv(file = "datbehav.csv") # path to file</pre>
```

STEP 1: have a look at your data

```
str(dat.behav)
summary(dat.behav)
plot(dat.behav)
```

t-test

```
fitstudent <- t.test(x = dat.behav$activity[dat.behav$weather==
                                               "rainy"],
                     y = dat.behav$activity[dat.behav$weather==
                                               "sunny"],
                     var.equal = TRUE)
print(fitstudent)
Two Sample t-test
data: dat.behav$activity[dat.behav$weather == "rainy"] and dat.behav
t = 3.2752, df = 33, p-value = 0.002485
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.6138373 2.6270325
sample estimates:
mean of x mean of y
 6.781476 5.161041
```

Linear models

ANOVA

```
fitanova <- aov(data = dat.behav, formula = activity ~ weather)

summary(fitanova)

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

weather 1 11.25 11.253 10.73 0.00248 **

Residuals 33 34.62 1.049
---

Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Linear regression

```
fitlm <- lm(data = dat.behav, formula = activity ~ weather)
summary(fitlm)
Call:
lm(formula = activity ~ weather, data = dat.behav)
Residuals:
   Min 1Q Median 3Q Max
-2.3547 -0.6028 0.2346 0.6419 1.6534
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 6.7815 0.4581 14.805 3.94e-16 ***
weathersunny -1.6204 0.4948 -3.275 0.00248 **
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

Residual standard error: 1.024 on 33 degrees of freedom

Linear models

February 20, 2018

18 / 19

NB: aov() vs. anova()

```
aov(data = dat.behav, formula = activity ~ weather)
anova(fitlm)
```