Reporting statistical result: t-test example

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Introduction

This vignette shows an example of reporting the results from a 2-sample t-test using data on the impact of invasive trout on salmon survival.

Outline of tasks

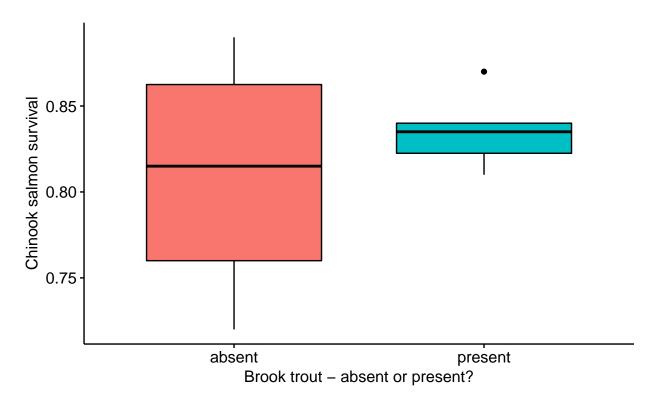
- Load the data into R
- Create a boxplot of the raw data
- State the relevant statistical null (Ho) and alternative (Ha) hypotheses
- Carry out an appropriate t-test
- Report the appropriate results in a full sentence as it would appear in a report or scientific paper

```
#load the data
## If I were loading the data by hand it would look something like this
## don't forget the commas!
salmon \leftarrow data.frame(survival = c(0.83,0.87,0.82,
                                   0.84,0.81,0.84,
                                   0.72, 0.84, 0.75,
                                   0.79, 0.89, 0.87),
                      brook.trout.PRES.ABS =
                        c("present","present","present",
                           "absent", "absent", "absent",
                           "absent", "absent", "absent"))
#Look at data
##I am using the salmon vs. brook trout data as an example
head(salmon)
     survival brook.trout.PRES.ABS
##
## 1
         0.83
                            present
## 2
         0.87
                            present
## 3
         0.82
                            present
## 4
         0.84
                            present
## 5
         0.81
                            present
## 6
         0.84
                            present
#Plot raw data
library(ggplot2)
library(ggpubr)
## Loading required package: magrittr
ggboxplot(data = salmon,
          y = "survival",
          x = "brook.trout.PRES.ABS",
          fill = "brook.trout.PRES.ABS",
```

"pre

```
xlab = "Brook trout - absent or present?",
ylab = "Chinook salmon survival")
```





State the hypotheses

##

mean in group absent mean in group present

0.810

Ho: The survival rates of Chinook salmon are the same whether brook trout are present or absent Ha: The persence of brook trout changes survival rates of salmon.

0.835

Report the results

For the real data, the results could be reported like this: "There was no evidence that the mean survival of salmon when brook trout are present (mean = 0.81) is different than when brook trout are absent (mean = 0.84; 2-sample t-test: p = 0.44, t = 0.82, n = 12 streams, df = 6)."

Normally I would also report the standard errors (SE) around the means, but for this exericise we will ignore it

Alternative results

What if the results really looked like this?

```
##
## Welch Two Sample t-test
##
## data: fake.surv by salmon$brook.trout.PRES.ABS
## t = 3.5938, df = 5.725, p-value = 0.01241
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.03173718 0.17227394
## sample estimates:
## mean in group absent mean in group present
## 0.8142305 0.7122249
```

The results could be reported like this: "Surival of chinook salmon in streams where brook were present (mean = 0.71) was significantly lower than when brook trout were absent (mean = 0.81) with a mean difference of 0.10 (95% CI: 0.03-0.17; 2-sample t-test p = 0.012, t = 3.6, n = 12 stream, df = 5.73)"