Statistical Inference Project Part 2

bzg 12-1-2022

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Synopsis
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This is a project for the Coursera Statistical Inference Class. The project consists of two parts: - Simulation Exercise to explore inference - Basic inferential analysis using the ToothGrowth data in the R datasets package

Part 2 - Inferential Analysis using ToothGrowth data

Overview

Analyse the ToothGrowth data in the R datasets package.

Instructions

```
· Load the ToothGrowth data and perform some basic exploratory data analysis
• Provide a basic summary of the data
```

• Use confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose • State your conclusions and the assumptions needed for your conclusions Preparing the environment

Load all necessary libraries.

```
# Load packages
library(dplyr, warn.conflicts = F)
library(ggplot2)
library(knitr)
library(data.table)
```

Attaching package: 'data.table' ## The following objects are masked from 'package:dplyr':

```
between, first, last
```

Description

```
Load the Tooth Growth dataset
 # Load data
 toothGrowth <- data.table(ToothGrowth)</pre>
```

len supp dose

1: 4.2 VC 0.5 ## 2: 11.5 VC 0.5

4: 27.3 OJ 2

str(ToothGrowth)

[1] 0.5 1.0 2.0

geom_boxplot() +

30 -

30 -

level 2 there doesn't appear to be any significant difference.

Checking effectiveness dose levels

summarise(avg.length = mean(len))

26.1

Summary of initial data analysis

Extract the len and dose vectors from toothGrowth

(Test.a <- t.test(len_a~dose_a, paired = FALSE))

Test B, dose = 0.5 and dose = 2 (p-value = 4.398e-14)

len_b <- toothGrowth %>%

dose_b <- toothGrowth %>%

select(len) %>%

select(dose) %>%

unlist()

unlist()

Test B

filter(dose %in% c(0.5,2)) %>%

filter(dose %in% c(0.5, 2)) %>%

10.605

filter(dose %in% c(1,2)) %>%

len_c <- toothGrowth %>%

dose_c <- toothGrowth %>%

Welch Two Sample t-test

data: len_c by dose_c

-8.996481 -3.733519 ## sample estimates:

95 percent confidence interval:

mean in group 1 mean in group 2 19.735

select(len) %>%

unlist()

Test C, dose = 1 and dose = 2 (p-value = 1.906e-05)

#Extract the len and dose vectors from toothGrowth

26.100

Extract the len and dose vectors from toothGrowth

26.1

toothGrowth %>%

filter(dose == 2) %>% group_by(supp) %>%

A tibble: 2 x 2 ## supp avg.length ## <fct> <dbl>

len_a <- toothGrowth %>%

dose_a <- toothGrowth %>%

select(len) %>%

select(dose) %>%

unlist()

unlist()

Test A

filter(dose %**in**% c(0.5,1)) %>%

filter(dose %in% c(0.5,1)) %>%

1 OJ

2 VC

 $facet_grid(. \sim supp) +$

```
The response is the length of odontoblasts (cells responsible for tooth growth) in 60 guinea pigs. Each animal received one of three dose levels of
vitamin C (0.5, 1, and 2 mg/day) by one of two delivery methods, (orange juice or ascorbic acid (a form of vitamin C and coded as VC).
Format
```

A data frame with 60 observations on 3 variables.

Exploratory data analysis # Explore dataset head(toothGrowth); tail(toothGrowth)

```
## 3: 7.3 VC 0.5
## 4: 5.8 VC 0.5
## 5: 6.4 VC 0.5
## 6: 10.0 VC 0.5
   len supp dose
## 1: 24.8 OJ 2
## 2: 30.9 OJ 2
## 3: 26.4 OJ 2
```

```
## 5: 29.4 OJ 2
 ## 6: 23.0 OJ 2
The three variables are length, supplement, and dose
 summary(ToothGrowth)
              supp
        len
                              dose
 ## Min. : 4.20 OJ:30 Min. :0.500
 ## 1st Qu.:13.07 VC:30 1st Qu.:0.500
```

Mean :18.81 Mean :1.167 ## 3rd Qu.:25.27 3rd Qu.:2.000 ## Max. :33.90 Max. :2.000

Median :19.25 Median :1.000

```
## 'data.frame': 60 obs. of 3 variables:
 ## $ len : num 4.2 11.5 7.3 5.8 6.4 10 11.2 11.2 5.2 7 ...
 ## $ supp: Factor w/ 2 levels "OJ", "VC": 2 2 2 2 2 2 2 2 2 2 ...
 ## $ dose: num 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
There are two levels (delivery types) for supp (supplement): OJ (Orange Juice) and VC (Vitamin C).
Length and dose are both numeric values but we cannot determine how many values exist for dose. View the unique values of dose.
 unique(ToothGrowth$dose)
```

There are three discrete levels for dose: 0.5, 1.0, and 2. We can conveniently convert it to a factor variable with three levels

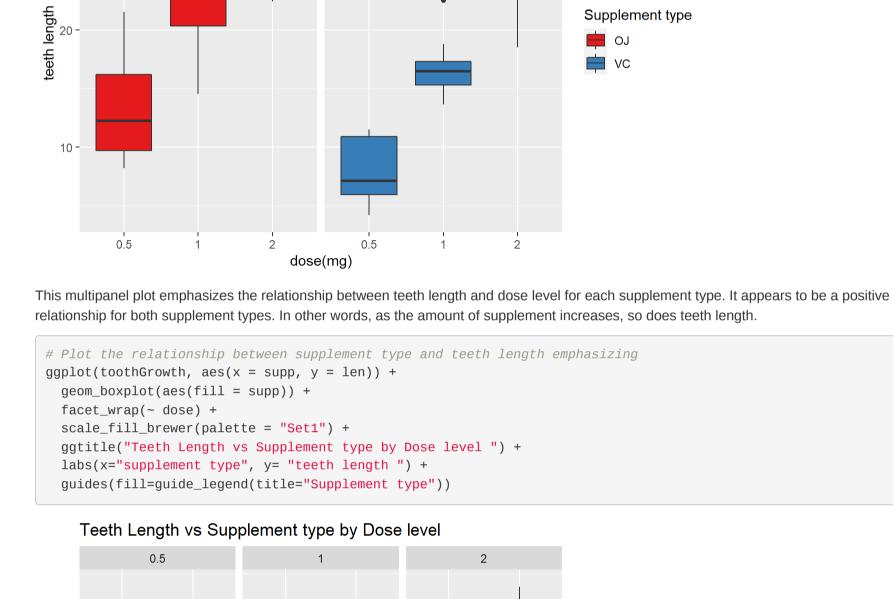
toothGrowth <- toothGrowth %>% mutate(dose = as.factor(dose))

ggplot(toothGrowth, aes(x=dose, y=len, fill = supp)) +

scale_fill_brewer(palette = "Set1") +

Visually examine the data by looking at the tooth length compared to dose by supplement. # Plot the relationship between teeth length and dose level for each supplement type

```
ggtitle("Teeth Length vs Dose level by Supplement type") +
labs(x="dose(mg)", y= "teeth length ") +
guides(fill=guide_legend(title="Supplement type"))
  Teeth Length vs Dose level by Supplement type
```



Supplement type

```
teeth length
                                                                                       OJ
            OJ
                                    supplement type
This second plot shows the relationship between supplement type and teeth length emphasizing direct comparison between supplement types.
```

Here the relationship is much less clear. Orange juice OJ appears to be more effective at dosage levels 0.5 and 1. On the other hand, at dosage

average length of 26.1 at that stage. Use confidence intervals and/or hypothesis tests to compare tooth

There appears to be an impact on tooth growth by increasing the dosage of Orange Juice, until a dosage of two. Both Supplement types score an

growth by supplement and dose Use t.test to determine if there is a difference in the performance of the treatments. First, we will run the test based on supplement. Looking to see if the p-value is smaller than 0.05 and if the confidence interval crosses 0. Testing by dose levels Test A, dose = 0.5 and dose = 1 (p-value = 1.268e-07)

```
## Welch Two Sample t-test
## data: len_a by dose_a
## t = -6.4766, df = 37.986, p-value = 1.268e-07
## alternative hypothesis: true difference in means between group 0.5 and group 1 is not equal to 0
## 95 percent confidence interval:
## -11.983781 -6.276219
## sample estimates:
## mean in group 0.5 mean in group 1
         10.605 19.735
```

(Test.b <- t.test(len_b~dose_b, paired = FALSE))</pre> ## Welch Two Sample t-test ## data: len_b by dose_b ## t = -11.799, df = 36.883, p-value = 4.398e-14 ## alternative hypothesis: true difference in means between group 0.5 and group 2 is not equal to 0 ## 95 percent confidence interval: ## -18.15617 -12.83383 ## sample estimates: ## mean in group 0.5 mean in group 2

filter(dose %in% c(1,2)) %>% select(dose) %>% unlist() #Test C (Test.c <- t.test(len_c~dose_c, paired = FALSE))</pre>

t = -4.9005, df = 37.101, p-value = 1.906e-05

26.100

```
We went through all possible combinations of levels from the factor variable dose and in all cases the p-value is lower than the default significance
level 0.05. We can safely reject the null. In other words there appears to be a positive relationship between dose level and teeth length.
Testing by Supplement
 #Extract the len and supp vectors from toothGrowth
 len <- toothGrowth %>%
   select(len) %>%
   unlist()
 supp <- toothGrowth %>%
   select(supp) %>%
   unlist()
 #Test T
 t.test(len~supp, paired=F)
```

alternative hypothesis: true difference in means between group 1 and group 2 is not equal to 0

```
# Test T 1
t.test(toothGrowth$len[toothGrowth$dose==2],
       toothGrowth$len[toothGrowth$dose==1],
       paired = FALSE, var.equal = TRUE)
##
## Two Sample t-test
## data: toothGrowth$len[toothGrowth$dose == 2] and toothGrowth$len[toothGrowth$dose == 1]
## t = 4.9005, df = 38, p-value = 1.811e-05
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 3.735613 8.994387
## sample estimates:
## mean of x mean of y
    26.100 19.735
```

Two Sample t-test ## data: toothGrowth\$len[toothGrowth\$dose == 1] and toothGrowth\$len[toothGrowth\$dose == 0.5]

```
19.735 10.605
Again, we see the p-value is still small although slighty larger than the previous test, therefore, it is significant. We can again reject the null
hypothesis and assume the dosage increase from .5mg to 1mg creates an positive effect on tooth growth.
There is no need for more testing of dosages given the previous tests.
State conclusions and the assumptions needed for conclusions
In this experiment, we assume there is a common variance in the population and that the guinea pigs were chosen at random. The delivery type
does not show a significant increase in tooth growth even though it does have a confidence level that crosses 0 at the 95% confidence.
However, there does appear to be a difference with an increase in tooth growth when the dosage is increased. The tests comparing the dosage
show confidence intervals of differences never crossing zero.
```

Increasing the dosage leads to an increase in tooth growth in guinea pigs.

95 percent confidence interval:

6.276252 11.983748 ## sample estimates: ## mean of x mean of y

Conclusion

```
## Welch Two Sample t-test
 ## data: len by supp
 ## t = 1.9153, df = 55.309, p-value = 0.06063
 ## alternative hypothesis: true difference in means between group OJ and group VC is not equal to 0
 ## 95 percent confidence interval:
 ## -0.1710156 7.5710156
 ## sample estimates:
 ## mean in group OJ mean in group VC
             20.66333
                               16.96333
We can see that the p-value of the test is 0.06. Since the p-value is greater than 0.05 and the confidence interval of the test contains zero, there is
not enough evidence to reject the null hypothesis. We cannot assume the delivery type has a significant effect on tooth growth.
Test the tooth length comparing the dosage of 1mg to 2mg to determine the effects of an increased dosage.
We see the p-value (1.811e-05) is very small, and is significant. Therefore, we can reject the null hypothesis and assume the dosage increase from
1mg to 2mg creates an positive effect on tooth growth.
Next, perform the test comparing the dosage of 0.5mg to 1mg.
 t.test(toothGrowth$len[toothGrowth$dose==1],
        toothGrowth$len[toothGrowth$dose==0.5],
         paired = FALSE, var.equal = TRUE)
 ## t = 6.4766, df = 38, p-value = 1.266e-07
 \#\# alternative hypothesis: true difference in means is not equal to 0
```