Statistical Inference Course Project (Part 2)

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Overview

In this second part of a two-part project assignment, we are being asked to investigate the effect of vitamin C on tooth growth in guinea pigs.

This analysis will compare the effects of different doses of vitamin C using two delivery methods to study the tooth growth in guinea pigs. The sample will consist of 60 guinea pigs. Each guinea pig will receive 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 supplements.

The study will be performed using the ToothGrowth dataset which is included in the R datasets package. The dataset is a data frame that contains 60 observations and 3 variables:

- len A numeric vector indicating the measurement of tooth length after vitamin C delivery
- supp A factor vector describing the delivery method used: Orange Juice (OJ) or Ascorbic Acid (VC)
- dose A numeric vector indicating the dosage level in milligrams per day (0.5, 1, or 2mg)

Further information on the ToothGrowth dataset can be found in the R documentation using ?ToothGrowth.

Environment Setup

Load packages used in this analysis.

```
if (!require(ggplot2)) {
    install.packages("ggplot2", repos = "http://cran.us.r-project.org")
    library(ggplot2)
}

## Loading required package: ggplot2

if (!require(dplyr, warn.conflicts = FALSE)) {
    install.packages("dplyr", repos = "http://cran.us.r-project.org")
    library(dplyr, warn.conflicts = FALSE)
}
```

Loading required package: dplyr

Load ToothGrowth dataset.

```
library(datasets)
data(ToothGrowth)
```

Basic Data Summary

```
After loading the ToothGrowth dataset, provide a basic summary of the data.
str(ToothGrowth)
                  60 obs. of 3 variables:
## 'data.frame':
## $ 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 ...
summary(ToothGrowth)
##
        len
                  supp
                              dose
## Min. : 4.20 OJ:30 Min. :0.500
## 1st Qu.:13.07 VC:30 1st Qu.:0.500
## Median :19.25
                         Median :1.000
                         Mean :1.167
## Mean :18.81
## 3rd Qu.:25.27
                         3rd Qu.:2.000
## Max. :33.90
                         Max. :2.000
# tabulate delivery method and dosage level values
table(ToothGrowth$supp, ToothGrowth$dose)
##
##
       0.5 1 2
    OJ 10 10 10
##
##
   VC 10 10 10
# summary of tooth length data grouped by delivery method and dosage level
by(data = ToothGrowth$len, INDICES = list(ToothGrowth$supp, ToothGrowth$dose), summary)
## : OJ
## : 0.5
   Min. 1st Qu. Median
##
                          Mean 3rd Qu.
     8.20 9.70 12.25 13.23 16.18 21.50
## : VC
## : 0.5
##
   Min. 1st Qu. Median Mean 3rd Qu.
                                          Max.
   4.20 5.95 7.15 7.98 10.90 11.50
## -----
## : OJ
## : 1
     Min. 1st Qu. Median Mean 3rd Qu.
                                          Max.
## 14.50 20.30 23.45
                          22.70 25.65
                                         27.30
## : VC
##
    Min. 1st Qu. Median
                          Mean 3rd Qu.
    13.60 15.28 16.50 16.77 17.30 22.50
## : OJ
```

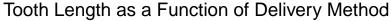
Exploratory Data Analysis

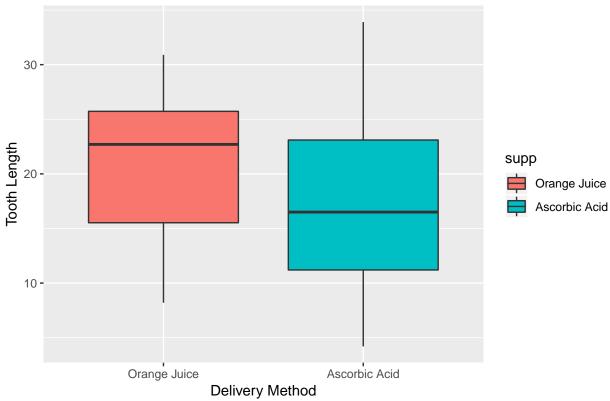
Perform some basic exploratory data analyses of the data. The analyses will explore the following relations:

- 1. Tooth Length (len) as a function of Delivery Method (supp)
- 2. Tooth Length (len) as a function of Dosage Level (dose)
- 3. Tooth Length (len) as a function of Delivery Method (supp) and Dosage Level (dose)

Tooth Length to Delivery Method

```
tg <- ToothGrowth
levels(tg$supp) <- c("Orange Juice", "Ascorbic Acid")
gLenSupp <- ggplot(data = tg, aes(x = supp, y = len)) +
    geom_boxplot(aes(fill = supp)) +
    xlab("Delivery Method") +
    ylab("Tooth Length") +
    theme(plot.title = element_text(size = 14, hjust = 0.5)) +
    ggtitle("Tooth Length as a Function of Delivery Method")
print(gLenSupp)</pre>
```





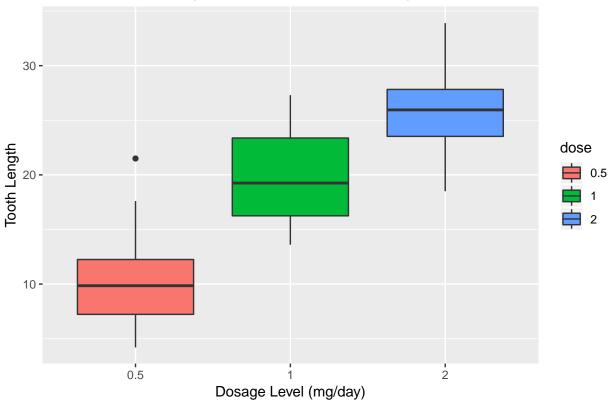
Observation

The above chart shows that using orange juice as the delivery method, independent of dosage level, had a more favorable effect on tooth growth than ascorbic acid.

Tooth Length to Dosage Level

```
gLenDose <- ggplot(data = ToothGrowth, aes(x = factor(dose), y = len)) +
    geom_boxplot(aes(fill = factor(dose))) +
    xlab("Dosage Level (mg/day)") +
    ylab("Tooth Length") +
    guides(fill=guide_legend(title="dose")) +
    theme(plot.title = element_text(size = 14, hjust = 0.5)) +
    ggtitle("Tooth Length as a Function of Dosage Level")
print(gLenDose)</pre>
```



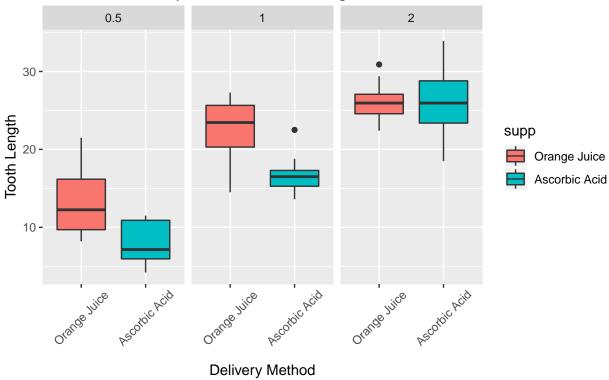


Observation

The above chart shows a positive relationship that higher dosage levels of vitamin C, independent of delivery method, had a more favorable effect on tooth growth than lower dosages of vitamin C. The 2 mg/day dosage level had the best effect on tooth growth followed by 1 mg/day and then 0.5 mg/day.

Tooth Length to Delivery Method and Dosage Level

Tooth Length as a Function of Delivery Method and Dosage Level



Observation

Looking at the above chart which shows tooth growth as a function of delivery method and dosage level, it appears that orange juice is more effective than ascorbic acid as the delivery method when the dosage level is 0.5 to 1 mg/day. The higher dosage level of 2 mg/day is more effective than the lower dosages; however, both delivery methods are equally as effective.

Inferential Statistics

Exploratory data analysis is helpful to study and visually interrogate data; however, the use of inferential statistics allows us to use techniques such as the confidence interval and/or hypothesis testing to more precisely draw conclusions about a population from which representative samples were taken.

In this section, hypothesis testing will be employed to study the impact of delivery method (orange juice and ascorbic acid supplements) and dosage level on tooth growth. A p-value less than or greater than a significance level of 5% will be used as the threshold to reject or accept the null hypothesis.

Hypothesis 1

Hypothesis testing will be conducted to study the impact of delivery method on tooth growth, independent of dosage level. A t-test will be performed on the null hypothesis that the two delivery methods have no effect on tooth growth.

```
t1 <- t.test(len ~ supp, data = ToothGrowth, conf.level = 0.95)
paste0("p-value = ", round(t1$p.value, 4))</pre>
```

[1] "p-value = 0.0606"

```
paste0("confidence interval = (", round(t1$conf.int[1], 4) , ", ", round(t1$conf.int[2], 4), ")")
## [1] "confidence interval = (-0.171, 7.571)"
```

The observed p-value 0.0606 is greater than 0.05 and the 95% confidence interval contains zero. This indicates weak evidence against the null hypothesis so we fail to reject the null hypothesis.

Hypothesis 2

Hypothesis testing will be conducted to study the impact of delivery method on tooth growth for a single dosage level. A t-test will be performed on the null hypothesis that the two delivery methods at a dosage level of 0.5 mg/day have no effect on tooth growth.

```
t2 <- t.test(len ~ supp, data = subset(ToothGrowth, dose == 0.5), conf.level = 0.95)
paste0("p-value = ", round(t2$p.value, 4))

## [1] "p-value = 0.0064"

paste0("confidence interval = (", round(t2$conf.int[1], 4), ", ", round(t2$conf.int[2], 4), ")")

## [1] "confidence interval = (1.7191, 8.7809)"

The observed p-value 0.0064 is less than 0.05 and the 95% confidence interval does not contain zero. This
```

The observed p-value 0.0064 is less than 0.05 and the 95% confidence interval does not contain zero. This indicates strong evidence against the null hypothesis so the null hypothesis can be rejected.

Hypothesis 3

Hypothesis testing will be conducted to study the impact of delivery method on tooth growth for a single dosage level. A t-test will be performed on the null hypothesis that the two delivery methods at a dosage level of 1 mg/day have no effect on tooth growth.

```
t3 <- t.test(len ~ supp, data = subset(ToothGrowth, dose == 1), conf.level = 0.95)
paste0("p-value = ", round(t3$p.value, 4))

## [1] "p-value = 0.001"

paste0("confidence interval = (", round(t3$conf.int[1], 4) , ", ", round(t3$conf.int[2], 4), ")")

## [1] "confidence interval = (2.8021, 9.0579)"</pre>
```

The observed p-value 0.001 is less than 0.05 and the 95% confidence interval does not contain zero. This indicates strong evidence against the null hypothesis so the null hypothesis can be rejected.

Hypothesis 4

Hypothesis testing will be conducted to study the impact of delivery method on tooth growth for a single dosage level. A t-test will be performed on the null hypothesis that the two delivery methods at a dosage level of 2 mg/day have no effect on tooth growth.

```
t4 <- t.test(len ~ supp, data = subset(ToothGrowth, dose == 2), conf.level = 0.95)
paste0("p-value = ", round(t4$p.value, 4))

## [1] "p-value = 0.9639"
paste0("confidence interval = (", round(t4$conf.int[1], 4) , ", ", round(t4$conf.int[2], 4), ")")

## [1] "confidence interval = (-3.7981, 3.6381)"</pre>
```

The observed p-value 0.9639 is greater than 0.05 and the 95% confidence interval contains zero. This indicates weak evidence against the null hypothesis so we fail to reject the null hypothesis.

Conclusion