## Statistical Inference Course Project Part 2

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2022-10-30

### Part 2: Basic Inferential Data Analysis Instructions

### Overview

In this second part, we perform basic inferential analyses using the ToothGrowth data in the R datasets package.

As per the help file of this dataset, the response is the length of odontoblasts (teeth) in each of 10 guinea pigs at each of three dose levels of Vitamin C (0.5, 1, and 2 mg) with each of two delivery methods (orange juice or ascorbic acid).

### Load Necessary Libraries

```
library(ggplot2)
```

### Load the ToothGrowth data and perform some basic exploratory data analyses.

```
# load the dataset
library(datasets)
data(ToothGrowth)
# look at the dataset variables
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 ...
   # convert variable dose from numeric to factor
ToothGrowth$dose <- as.factor(ToothGrowth$dose)</pre>
# look at the dataset variables after conversion
str(ToothGrowth)
## '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: Factor w/ 3 levels "0.5", "1", "2": 1 1 1 1 1 1 1 1 1 1 ...
```

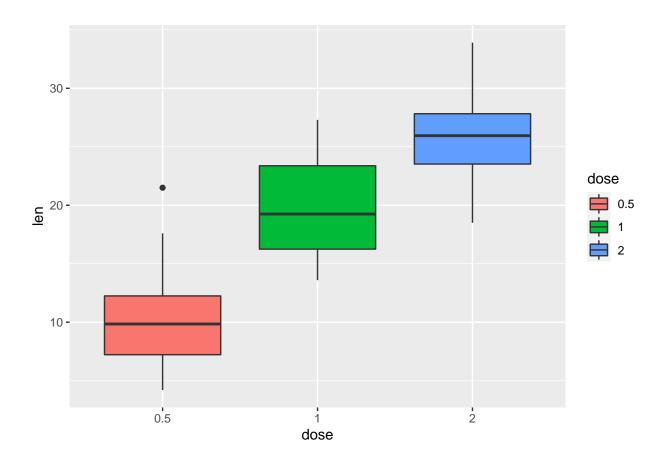
## Provide a basic summary of the data.

# # summary statistics for all variables summary(ToothGrowth)

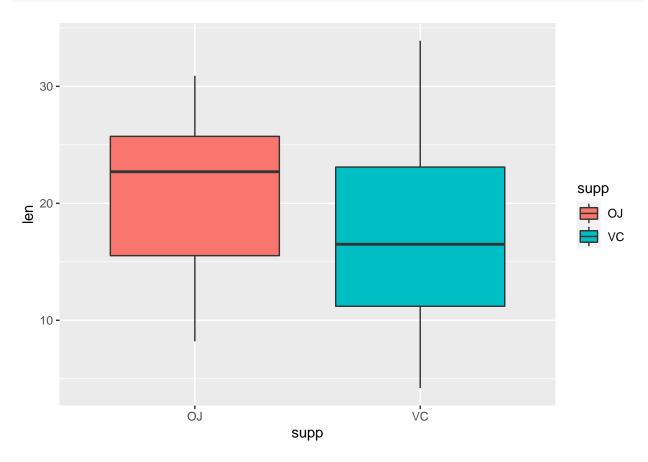
```
##
        len
                  supp
                           dose
         : 4.20
                  OJ:30
                          0.5:20
##
  Min.
  1st Qu.:13.07
                  VC:30
                         1 :20
## Median :19.25
                          2 :20
## Mean
         :18.81
## 3rd Qu.:25.27
## Max.
         :33.90
```

## # split of cases between different dose levels and delivery methods table(ToothGrowth\$dose, ToothGrowth\$supp)

### ggplot(aes(x=dose, y=len), data=ToothGrowth) + geom\_boxplot(aes(fill=dose))



```
# visualization of tooth growth as function of supplement type
ggplot(aes(x=supp, y=len), data=ToothGrowth) + geom_boxplot(aes(fill=supp))
```



Use confidence intervals and hypothesis tests to compare tooth growth by supp and dose.

```
# check for group differences due to different supplement type
# assuming unequal variances between the two groups
t.test(len ~ supp, data = ToothGrowth)
##
   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 O
## 95 percent confidence interval:
## -0.1710156 7.5710156
## sample estimates:
## mean in group OJ mean in group VC
##
           20.66333
                            16.96333
```

The p-value is 0.06, and the confidence interval contains zero. This indicates that we can not reject the null hypothesis that the different supplement types have no effect on tooth length.

```
# first create three sub-groups as per dose level pairs
ToothGrowth.doses_0.5_1.0 <- subset (ToothGrowth, dose %in% c(0.5, 1.0))
ToothGrowth.doses_0.5_2.0 <- subset (ToothGrowth, dose %in% c(0.5, 2.0))
ToothGrowth.doses_1.0_2.0 <- subset (ToothGrowth, dose %in% c(1.0, 2.0))
# Check for group differences due to different dose levels (0.5, 1.0)
# assuming unequal variances between the two groups
t.test(len ~ dose, data = ToothGrowth.doses_0.5_1.0)
##
   Welch Two Sample t-test
##
## data: len by dose
## 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
# Check for group differences due to different dose levels (0.5, 2.0)
# assuming unequal variances between the two groups
t.test(len ~ dose, data = ToothGrowth.doses_0.5_2.0)
##
##
   Welch Two Sample t-test
## data: len by dose
## 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
##
              10.605
                                26.100
# Check for group differences due to different dose levels (1.0, 2.0)
# assuming unequal variances between the two groups
t.test(len ~ dose, data = ToothGrowth.doses_1.0_2.0)
##
## Welch Two Sample t-test
##
## data: len by dose
## t = -4.9005, df = 37.101, p-value = 1.906e-05
## alternative hypothesis: true difference in means between group 1 and group 2 is not equal to 0
## 95 percent confidence interval:
## -8.996481 -3.733519
## sample estimates:
## mean in group 1 mean in group 2
                            26.100
##
            19.735
```

For all three dose level pairs, the p-value is less than 0.05, and the confidence interval does not contain zero. The mean tooth length increases on raising the dose level. This indicates that we can reject the null hypothesis, and establish that increasing the dose level leads to an increase in tooth length.

### State your conclusions and the assumptions needed for your conclusions.

#### Conclusions

- 1. Supplement type has no effect on tooth growth.
- 2. In reasing the dose level leads to increased tooth growth.

### Assumptions

- 1. The experiment was done with random assignment of guinea pigs to different dose level categories and supplement type to control for confounders that might affect the outcome.
- 2. Members of the sample population, i.e. the 60 guinea pigs, are representative of the entire population of guinea pigs. This assumption allows us to generalize the results.
- 3. For the t-tests, the variances are assumed to be different for the two groups being compared. This assumption is less stronger than the case in which the variances are assumed to be equal.