# Statistical Inference Project Part 2

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Now in the second portion of the project, we're going to analyze the ToothGrowth data in the R datasets package.

#### Instructions

- 1. Load the ToothGrowth data and perform some basic exploratory data analyses
- 2. Provide a basic summary of the data.
- 3. Use confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose. (Only use the techniques from class, even if there's other approaches worth considering)
- 4. State your conclusions and the assumptions needed for your conclusions.

#### Loading Libraries

```
library("ggplot2")
```

## Warning: package 'ggplot2' was built under R version 3.6.3

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

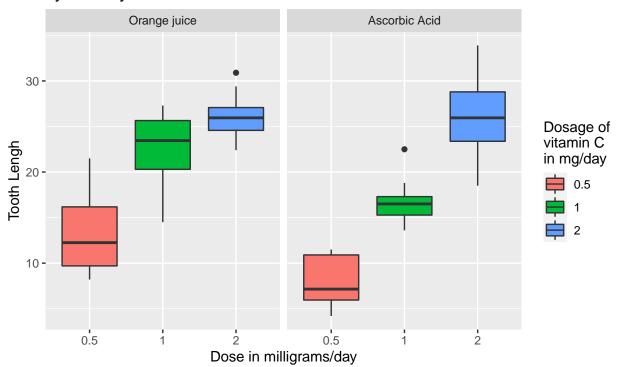
```
# Load the data ToothGrowth
data(ToothGrowth)
# Look at the structure of the data
str(ToothGrowth)
## 'data.frame':
                   60 obs. of 3 variables:
## $ len : num 4.2 11.5 7.3 5.8 6.4 10 11.2 11.2 5.2 7 ...
## $ supp: Factor w/ 2 levels "OJ", "VC": 2 2 2 2 2 2 2 2 2 2 ...
## $ dose: num 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
# Look at the first 5 rows of the data
head(ToothGrowth, 5)
##
     len supp dose
## 1
    4.2
           VC 0.5
## 2 11.5
           VC 0.5
## 3 7.3
           VC 0.5
     5.8
           VC
              0.5
## 5 6.4
           VC 0.5
```

2. Provide a basic summary of the data.

```
# Look at the summary of the data
summary(ToothGrowth)
```

```
##
         len
                    supp
                                 dose
##
           : 4.20
                    OJ:30
                                   :0.500
   Min.
                            Min.
                    VC:30
   1st Qu.:13.07
                            1st Qu.:0.500
  Median :19.25
                            Median :1.000
##
##
   Mean
           :18.81
                            Mean
                                   :1.167
##
   3rd Qu.:25.27
                            3rd Qu.:2.000
   Max.
           :33.90
                            Max.
                                   :2.000
# Compare means of the different delivery methods
tapply(ToothGrowth$len,ToothGrowth$supp, mean)
##
         OJ
## 20.66333 16.96333
# Make a plot to look at data graphically
ggplot(ToothGrowth, aes(factor(dose), len, fill = factor(dose))) +
      geom_boxplot() +
      # facet_grid(.~supp)+
      facet_grid(.~supp, labeller = as_labeller(
            c("OJ" = "Orange juice",
              "VC" = "Ascorbic Acid"))) +
      labs(title = "Tooth growth of 60 guinea pigs
           by dosage and nby delivery method of vitamin C",
           x = "Dose in milligrams/day",
           y = "Tooth Lengh") +
      scale_fill_discrete(name = "Dosage of\nvitamin C\nin mg/day")
```

## Tooth growth of 60 guinea pigs by dosage and by delivery method of vitamin C



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

```
# Comparison by delivery method for the same dosage
t05 <- t.test(len ~ supp,
       data = rbind(ToothGrowth[(ToothGrowth$dose == 0.5) &
                                       (ToothGrowth$supp == "OJ"),],
                    ToothGrowth[(ToothGrowth$dose == 0.5) &
                                       (ToothGrowth$supp == "VC"),]),
       var.equal = FALSE)
t1 <- t.test(len ~ supp,
       data = rbind(ToothGrowth[(ToothGrowth$dose == 1) &
                                       (ToothGrowth$supp == "OJ"),],
                    ToothGrowth[(ToothGrowth$dose == 1) &
                                       (ToothGrowth$supp == "VC"),]),
       var.equal = FALSE)
t2 <- t.test(len ~ supp,
       data = rbind(ToothGrowth[(ToothGrowth$dose == 2) &
                                       (ToothGrowth$supp == "OJ"),],
                    ToothGrowth[(ToothGrowth$dose == 2) &
                                       (ToothGrowth$supp == "VC"),]),
       var.equal = FALSE)
# Make summary of the conducted t.tests, which compare the delivery methods by dosage
# take p-values and CI
summaryBYsupp <- data.frame(</pre>
      "p-value" = c(t05\$p.value, t1\$p.value, t2\$p.value),
      "Conf.Low" = c(t05\$conf.int[1], t1\$conf.int[1], t2\$conf.int[1]),
      "Conf.High" = c(t05$conf.int[2],t1$conf.int[2], t2$conf.int[2]),
      row.names = c("Dosage .05", "Dosage 1", "Dosage 2"))
# Show the data table
summaryBYsupp
                  p.value Conf.Low Conf.High
## Dosage .05 0.006358607
                           1.719057 8.780943
              0.001038376 2.802148
                                     9.057852
```

## Dosage 1 0.963851589 -3.798070 3.638070 ## Dosage 2

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

With 95% confidence we reject the null hypothesis, stating that there is no difference in the tooth growth by the delivery method for .5 and 1 milligrams/day. We observe p-values less than the treshold of .05 and the confidence levels don't include 0. So, for dosage of .5 milligrams/day and 1 milligrams/day does matter the delivery method. With 95% confidence we fail to reject the null hypothesis, stating that there is no difference in the tooth growth by the delivery method for 2 milligrams/day. We observe p-values more than the treshold of .05 and the confidence levels include 0. So, for dosage of 2 milligrams/day the delivery method doesn't matter.