# Statistical Inference Course Project - Part II - Basic Inferential Data Analysis

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#### Overview

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

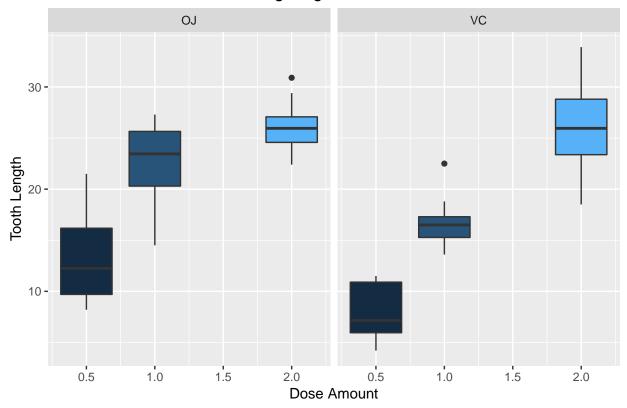
- 1. Load the ToothGrowth data and perform some basic exploratory data analyses
- 2. Provide a basic summary of the data.
- 3. Use confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose.
- 4. Conclusions

# Load the ToothGrowth data and perform some basic exploratory data analyses & Provide a basic summary of the data.

```
Load ToothGrowth data
data("ToothGrowth")
Summary of data
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
           :18.81
                             Mean
                                    :1.167
   3rd Qu.:25.27
                             3rd Qu.:2.000
## Max.
           :33.90
                             Max.
                                    :2.000
First few rows
head (ToothGrowth)
##
      len supp dose
## 1 4.2
            VC 0.5
## 2 11.5
            VC 0.5
## 3 7.3
            VC 0.5
## 4 5.8
            VC 0.5
## 5 6.4
            VC 0.5
## 6 10.0
            VC 0.5
Exploratory Plot
library(ggplot2)
ggplot(data = ToothGrowth, aes(x = dose, y = len)) +
```

geom\_boxplot(aes(fill = dose, group = dose)) +

## Tooth Length against Dose Amount



#### Run t-tests for different variables

```
ToothGrowth$dose <- as.factor(ToothGrowth$dose)

t1 <- t.test(data = ToothGrowth, len ~ supp)

ToothGrowth_0.5_1.0 <- subset(ToothGrowth, ToothGrowth$dose %in% c(0.5, 1.0))

t2 <- t.test(len ~ dose,data = ToothGrowth_0.5_1.0)

ToothGrowth_0.5_2.0 <- subset(ToothGrowth, ToothGrowth$dose %in% c(0.5, 2.0))

t3 <- t.test(len ~ dose,data = ToothGrowth_0.5_2.0)

ToothGrowth_1.0_2.0 <- subset(ToothGrowth, ToothGrowth$dose %in% c(1.0, 2.0))

t4 <- t.test(len ~ dose,data = ToothGrowth_1.0_2.0)
```

## Conclusions

Look at the results

```
t1$estimate
## mean in group OJ mean in group VC
           20.66333
                             16.96333
t1$p.value
## [1] 0.06063451
t1$conf.int
## [1] -0.1710156 7.5710156
## attr(,"conf.level")
## [1] 0.95
The p-value of this test is 0.06. Since the p-value is greater than 0.05 and the confidence interval of the test
contains zero we can say that supplement types seems to have no impact on Tooth growth based on this test.
t2$estimate
## mean in group 0.5
                        mean in group 1
##
              10.605
                                  19.735
t2$p.value
## [1] 1.268301e-07
t2$conf.int
## [1] -11.983781 -6.276219
## attr(,"conf.level")
## [1] 0.95
t3$estimate
                        mean in group 2
## mean in group 0.5
##
              10.605
                                  26.100
t3$p.value
## [1] 4.397525e-14
t3$conf.int
## [1] -18.15617 -12.83383
## attr(,"conf.level")
## [1] 0.95
t4$estimate
## mean in group 1 mean in group 2
##
            19.735
                             26.100
t4$p.value
## [1] 1.90643e-05
t4$conf.int
## [1] -8.996481 -3.733519
## attr(,"conf.level")
## [1] 0.95
```

As can be seen, the p-value of each test was essentially zero and the confidence interval of each test does not cross over zero (0).

Based on this result we can assume that the average tooth length increases with an inceasing dose, and therefore the null hypothesis can be rejected.

- 1. The sample is representative of the population
- 2. The distribution of the sample means follows the Central Limit Theorem
- 3. The 95% confidence interval is adapted as the standard

With the results of t-tests above, we can conclude that supplement delivery method has no effect on tooth length with 95% confidence interval. However increased dosages result in increased tooth length.