

# Statistical Inference Course Project - Part2

## Overview

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

```
library(ggplot2)
library(dplyr)
```

## Part 2: Basic Inferential Data Analysis Instructions

Load the ToothGrowth data and perform some basic exploratory data analyses

```
data(ToothGrowth)
```

Provide a basic summary of the data.

```
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
```

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

```
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 0.5 ...
```

```
# plot len & supp
ggplot(data=ToothGrowth) +
  aes(x=supp, y=len)+
  geom_boxplot(aes(fill=supp)) +
  facet_grid(cols = vars(dose)) +
  ggtitle("Length-Supplement Relation split by dose")
```



From Figure above, mean seems to be equal for both supp only for dose=2.

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)

```
# create t test
```

```
# perform t test between supp types where dose = 2
```

```
t.test(ToothGrowth$len[ToothGrowth$supp=="OJ" & ToothGrowth$dose==2], ToothGrowth$len[ToothGrowth$supp=="VC" & ToothGrowth$dose==2], paired = TRUE)
```

```
##
```

```
## Paired t-test
```

```
##
```

```
## data: ToothGrowth$len[ToothGrowth$supp == "OJ" & ToothGrowth$dose == 2] and ToothGrowth$len[ToothGrowth$supp == "VC" & ToothGrowth$dose == 2]
```

```
## t = -0.042592, df = 9, p-value = 0.967
```

```
## alternative hypothesis: true difference in means is not equal to 0
```

```
## 95 percent confidence interval:
```

```
## -4.328976 4.168976
```

```
## sample estimates:
```

```
## mean of the differences
```

```
## -0.08
```

```
# perform t test between supp types where dose != 2
```

```
t.test(ToothGrowth$len[ToothGrowth$supp=="OJ" & ToothGrowth$dose!=2], ToothGrowth$len[ToothGrowth$supp=="VC" & ToothGrowth$dose!=2], paired = TRUE)
```

```
##
```

```
## Paired t-test
```

```
##
```

```
## data: ToothGrowth$len[ToothGrowth$supp == "OJ" & ToothGrowth$dose != 2] and ToothGrowth$len[ToothGrowth$supp == "VC" & ToothGrowth$dose != 2]
```

```
## t = 4.6042, df = 19, p-value = 0.0001936
```

```
## alternative hypothesis: true difference in means is not equal to 0
```

```
## 95 percent confidence interval:
```

```
## 3.048852 8.131148
```

```
## sample estimates:
```

```
## mean of the differences
```

```
## 5.59
```

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

If we consider Null Hypothesis ( $H_0$ ) to be; mean is almost equal per supp per dose

1- We Fail to reject  $H_0$  where dose = 2, as t-value is very small and is equal to -0.042592

2- We reject  $H_0$  where dose does not equal to 2, as t-value is large enough and is equal to 4.6042202