# The ToothGrowth data Analysis

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# 07/06/2020

#### Overview

We are going to analyze the ToothGrowth data in the R datasets package. We will study the change of the tooth length in pigs each type of supplement and changing dose quantity.

## **Exploratory Data Analysis**

First, loading the dataset and check its dimension

```
library(dplyr)
```

```
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
## filter, lag
## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union

tg <- ToothGrowth
dim(tg)</pre>
```

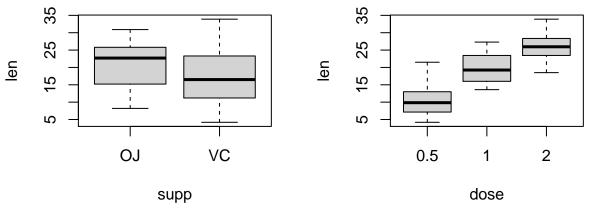
## [1] 60 3

The dataset contains 60 rows and 3 columns

Next, summarizing the basic data from the dataset

## summary(ToothGrowth)

```
##
        len
                                 dose
                    supp
         : 4.20
                   OJ:30
                                   :0.500
## Min.
                            Min.
  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
                                   :2.000
par(mfrow=c(1,2))
boxplot(len ~ supp, data=tg)
boxplot(len ~ dose, data=tg)
```



seem giving orange joice is better than an acid and give more dose is better

## Inferential Data Analysis

Create variables each type of supplement

```
OJ <- subset(tg, supp == 'OJ')
VC <- subset(tg, supp == 'VC')</pre>
```

It's

```
Next.
t.test(OJ$len, VC$len, alternative='greater', paired = FALSE, var.equal = FALSE)
##
##
   Welch Two Sample t-test
##
## data: OJ$len and VC$len
## t = 1.9153, df = 55.309, p-value = 0.03032
## alternative hypothesis: true difference in means is greater than 0
## 95 percent confidence interval:
## 0.4682687
                    Tnf
## sample estimates:
## mean of x mean of y
    20.66333 16.96333
half.dose <- subset(tg, dose == 0.5)
one.dose <- subset(tg, dose == 1)</pre>
two.dose <- subset(tg, dose == 2)</pre>
```

One dose effect greater than Half dose

10.605

##

19.735

Two dose effect greater than One dose