Stat-Inference Assignment-02

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```
library(ggplot2)

## Warning: package 'ggplot2' was built under R version 3.2.2

### Load the dataset
data(ToothGrowth)
```

Explore 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
```

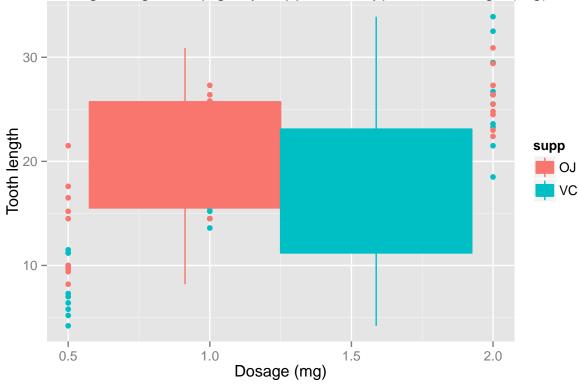
summary(ToothGrowth)

```
##
                                   dose
         len
                     supp
##
   \mathtt{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
##
           :33.90
                                     :2.000
   Max.
                             Max.
```

```
# visualize the data
```

qplot(dose,len,data=ToothGrowth, col=supp, main="Tooth length of guinea pigs by supplement type and dos





##Hypothesis Testing

H0: There is no significant difference in treatment from both the medicine OJ and VC

H1: alternate of H0

Tooth length in VC = OJ H0

Tooth length in VC NOT equal to OJ treaments

Split the two treatment groups

```
OJ = ToothGrowth$len[ToothGrowth$supp == 'OJ']
VC = ToothGrowth$len[ToothGrowth$supp == 'VC']
```

T- Test

one tail test

```
t.test(OJ, VC, alternative = "greater", paired = FALSE, var.equal = FALSE, conf.level = 0.95)
```

##

Hypothesis for dosage

HO: there is no sifnificant difference in dosage in two sizes

H1: There is difference in the two dosage

```
##Divide the dataset with dosage group
OJ2 = ToothGrowth$len[ToothGrowth$supp == 'OJ' & ToothGrowth$dose == 2]
VC2 = ToothGrowth$len[ToothGrowth$supp == 'VC' & ToothGrowth$dose == 2]
half = ToothGrowth$len[ToothGrowth$dose == 0.5]
one = ToothGrowth$len[ToothGrowth$dose == 1]
two = ToothGrowth$len[ToothGrowth$dose == 2]
# T-test
##One-tailed independent t-test with unequal variance.
t.test(half, one, alternative = "less", paired = FALSE, var.equal = FALSE, conf.level = 0.95)
##
   Welch Two Sample t-test
##
## data: half and one
## t = -6.4766, df = 37.986, p-value = 6.342e-08
## alternative hypothesis: true difference in means is less than 0
## 95 percent confidence interval:
         -Inf -6.753323
##
## sample estimates:
## mean of x mean of y
      10.605
                19.735
```

p-value = 6.342e-08 is less than the 5% significance level .05 Hence accepting the Alternate hypothesis as there is difference in tooth growth.

Note that the echo = FALSE parameter was added to the code chunk to prevent printing of the R code that generated the plot.