# Statistical\_Infer\_Part2.Rmd

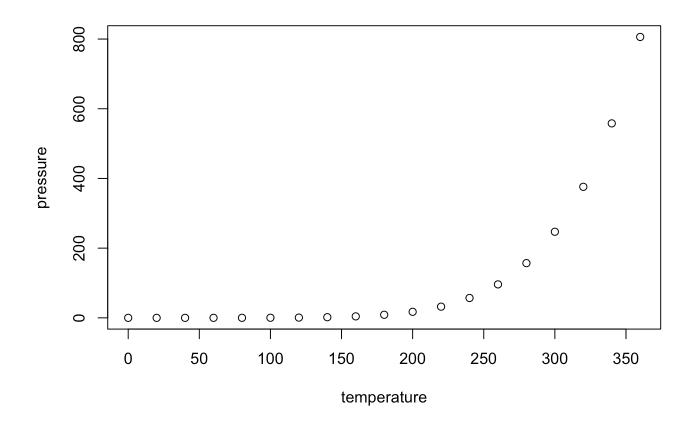
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### Instructions

- 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.

## **Including Plots**

You can also embed plots, for example:



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

## **Exploratory data Analysis**

First me load the packages, and dataset

```
library(ggplot2)
library(knitr)
library(datasets)
```

Load the ToothGrowth data and perform basic Exploratory Data Analysis

```
data(ToothGrowth)
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 2 ...
## $ dose: num 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
```

```
head(ToothGrowth, 4)
```

```
## 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
```

```
tail(ToothGrowth, 4)
```

```
## len supp dose

## 57 26.4 0J 2

## 58 27.3 0J 2

## 59 29.4 0J 2

## 60 23.0 0J 2
```

Calculate the summary of the data

```
summary(ToothGrowth)
```

```
##
         len
                                  dose
                    supp
##
  Min.
           : 4.20
                    0J: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
```

Calculate the mean of the length

```
suppl_mean = split(ToothGrowth$len, ToothGrowth$supp)
sapply(suppl_mean, mean)
```

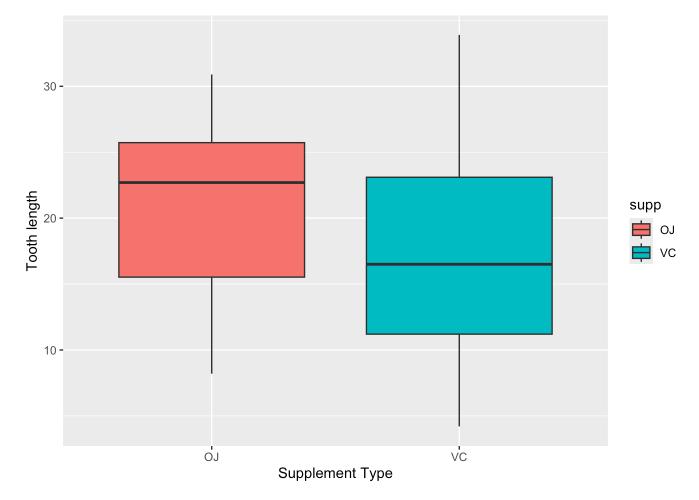
```
## 0J VC
## 20.66333 16.96333
```

#### suppl\_mean

```
## $0J
## [1] 15.2 21.5 17.6 9.7 14.5 10.0 8.2 9.4 16.5 9.7 19.7 23.3 23.6 26.4 20.0
## [16] 25.2 25.8 21.2 14.5 27.3 25.5 26.4 22.4 24.5 24.8 30.9 26.4 27.3 29.4 23.0
##
## $VC
## [1] 4.2 11.5 7.3 5.8 6.4 10.0 11.2 11.2 5.2 7.0 16.5 16.5 15.2 17.3 22.5
## [16] 17.3 13.6 14.5 18.8 15.5 23.6 18.5 33.9 25.5 26.4 32.5 26.7 21.5 23.3 29.5
```

## Basic Exploratory Analysis, Graph below

ggplot(aes(x=supp, y=len), data=ToothGrowth) + geom\_boxplot(aes(fill=supp))+
 xlab("Supplement Type") +ylab("Tooth length")



Get the confidence intervals

unique(ToothGrowth\$dose)

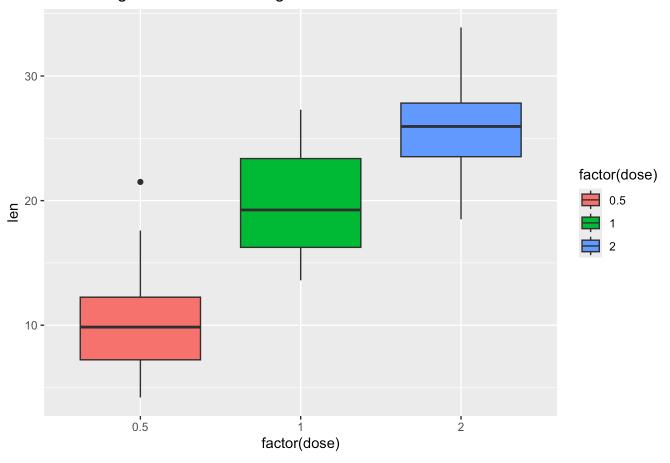
```
## [1] 0.5 1.0 2.0
```

Unique dose groups are 0.5, 1, 2

Graph below, shows the relationship between Tooth Length and Dosages

```
ggplot(aes(x = factor(dose), y = len), data = ToothGrowth) +
geom_boxplot(aes(fill = factor(dose))) +
ggtitle("Tooth length relation to Dosage")
```

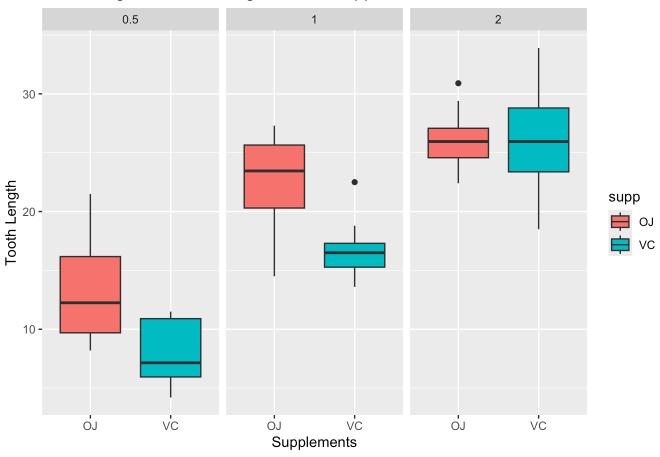
### Tooth length relation to Dosage



Graph below show the tooth Length realation to dosage of each supplement

```
ggplot(aes(x=supp, y=len), data=ToothGrowth) +
  geom_boxplot(aes(fill=supp)) + xlab("Supplements") +
  ylab("Tooth Length") + facet_grid(~ dose) +
  ggtitle("Tooth length relation dosage of each Supplement")
```

### Tooth length relation dosage of each Supplement



Hypothesis test defined below:

H0: tooth length does not depend of different supplements Ha: tooth length are effected by different supplement

```
t.test(len ~ supp, ToothGrowth[ToothGrowth$dose == .5, ])
```

```
##
## Welch Two Sample t-test
##
## data: len by supp
## t = 3.1697, df = 14.969, p-value = 0.006359
## alternative hypothesis: true difference in means between group OJ and group VC is not equal to 0
## 95 percent confidence interval:
## 1.719057 8.780943
## sample estimates:
## mean in group OJ mean in group VC
## 13.23 7.98
```

```
t.test(len ~ supp, ToothGrowth[ToothGrowth$dose == 1, ])
```

```
##
## Welch Two Sample t-test
##
## data: len by supp
## t = 4.0328, df = 15.358, p-value = 0.001038
## alternative hypothesis: true difference in means between group OJ and group VC is not equal to 0
## 95 percent confidence interval:
## 2.802148 9.057852
## sample estimates:
## mean in group OJ mean in group VC
## 22.70 16.77
```

```
t.test(len ~ supp, ToothGrowth[ToothGrowth$dose == 2, ])
```

```
##
## Welch Two Sample t-test
##
## data: len by supp
## t = -0.046136, df = 14.04, p-value = 0.9639
## alternative hypothesis: true difference in means between group OJ and group VC is not equal to 0
## 95 percent confidence interval:
## -3.79807 3.63807
## sample estimates:
## mean in group OJ mean in group VC
## 26.06 26.14
```

### Conclusion

we reject the Null Hypothesis, give more explanation on each test, CHATGPT use.