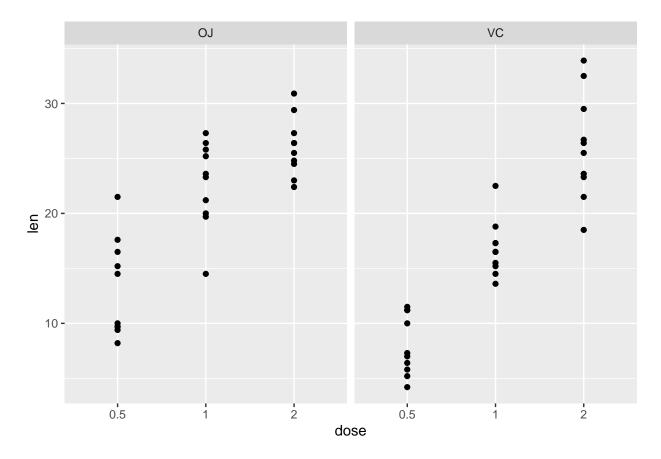
# Part 2

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Basic Inferential Data Analysis-We are going to analyze the ToothGrowth data in the R datasets package. ##Q1 Load the ToothGrowth data and perform some basic exploratory data analyses

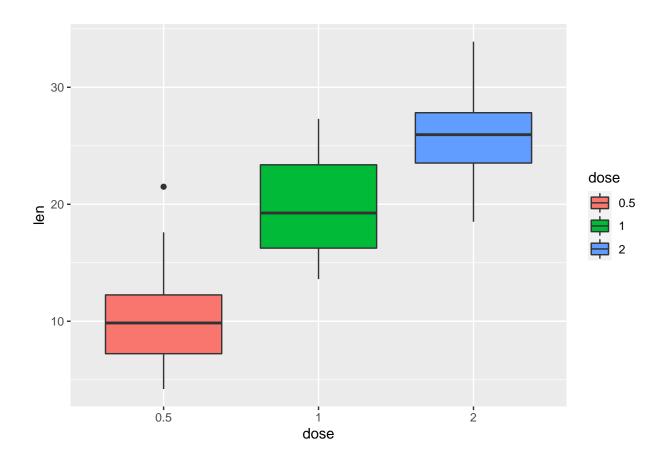
```
library(ggplot2)
data(ToothGrowth)
str(ToothGrowth)
## 'data.frame':
                   60 obs. of 3 variables:
## $ len : num 4.2 11.5 7.3 5.8 6.4 10 11.2 11.2 5.2 7 ...
## $ supp: Factor w/ 2 levels "OJ", "VC": 2 2 2 2 2 2 2 2 2 2 ...
## $ dose: num 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
head (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
ToothGrowth$dose <- as.factor(ToothGrowth$dose)</pre>
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 ...
## $ dose: Factor w/ 3 levels "0.5", "1", "2": 1 1 1 1 1 1 1 1 1 1 ...
qplot(dose, len, data=ToothGrowth, facets=.~supp, geom=c("point", "smooth"), method="loess")
## Warning: Ignoring unknown parameters: method
## 'geom_smooth()' using formula 'y ~ x'
```



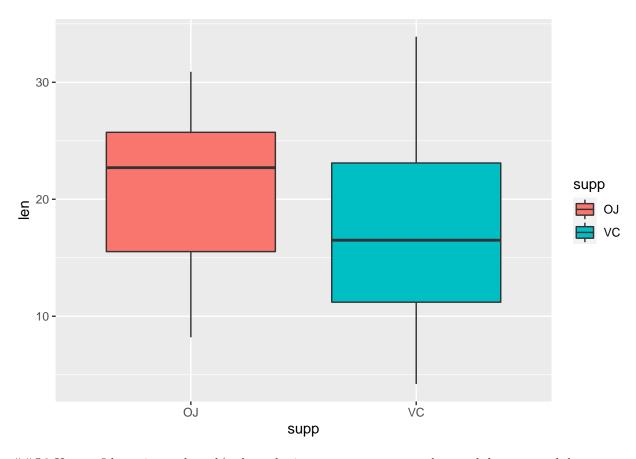
 $\#\#\mathrm{Q2}$  Provide a basic summary of the data.

## summary(ToothGrowth)

```
##
        len
                   supp
                           dose
## Min. : 4.20
                   OJ:30
                          0.5:20
##
   1st Qu.:13.07
                   VC:30
                          1 :20
  Median :19.25
                          2 :20
##
## Mean
         :18.81
   3rd Qu.:25.27
##
   Max.
         :33.90
ggplot(aes(x=dose, y=len), data=ToothGrowth) + geom_boxplot(aes(fill=dose))
```



ggplot(aes(x=supp, y=len), data=ToothGrowth) + geom\_boxplot(aes(fill=supp))



##Q3 Use confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose.

```
varsupp <- ToothGrowth$supp</pre>
vardose <- ToothGrowth$dose</pre>
varlen <- ToothGrowth$len</pre>
t.test(varlen[varsupp == "VC"], varlen[varsupp == "OJ"], paired=FALSE)
##
##
   Welch Two Sample t-test
##
## data: varlen[varsupp == "VC"] and varlen[varsupp == "OJ"]
## t = -1.9153, df = 55.309, p-value = 0.06063
\mbox{\tt \#\#} alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -7.5710156 0.1710156
## sample estimates:
## mean of x mean of y
## 16.96333 20.66333
t.test(varlen[vardose == 0.5], varlen[vardose == 1], paired=FALSE)
##
   Welch Two Sample t-test
##
```

```
## data: varlen[vardose == 0.5] and varlen[vardose == 1]
## t = -6.4766, df = 37.986, p-value = 1.268e-07
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -11.983781 -6.276219
## sample estimates:
## mean of x mean of y
## 10.605 19.735
```

 $\#\#\mathrm{Q4}$  State your conclusions and the assumptions needed for your conclusions.

#### Conclusion-

- 1. For guiena pigs Vitamin c is a factor of growth for their teeth.
- 2. Whereas the delivery mode does not have any imapact on their growth.

#### Assumption-

1. We assume that variance in all groups should be expected to be equal. 2. The underlying assumption is that sampling of Guinea Pigs to assign them to a supplement and a dose was done properly.