Course6-2

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Overview

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

1. Load the ToothGrowth data and explore data analyses

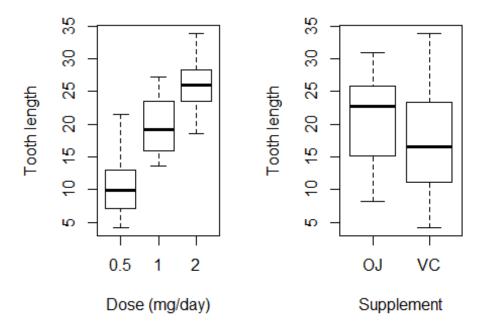
```
data(ToothGrowth)
str(ToothGrowth)
                   60 obs. of 3 variables:
## 'data.frame':
## $ len : num 4.2 11.5 7.3 5.8 6.4 10 11.2 11.2 5.2 7 ...
## $ supp: Factor w/ 2 levels "OJ", "VC": 2 2 2 2 2 2 2 2 2 2 ...
## $ dose: num 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
head(ToothGrowth)
##
     len supp dose
## 1 4.2
           VC 0.5
## 2 11.5
           VC 0.5
           VC 0.5
## 3 7.3
## 4 5.8
           VC 0.5
## 5 6.4
           VC 0.5
## 6 10.0
           VC 0.5
unique(ToothGrowth$dose)
## [1] 0.5 1.0 2.0
```

2. Provide a basic summary of the data.

```
summary(ToothGrowth)
##
                                dose
        len
                   supp
## Min. : 4.20
                   OJ:30
                                  :0.500
                           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
                           Max. :2.000
```

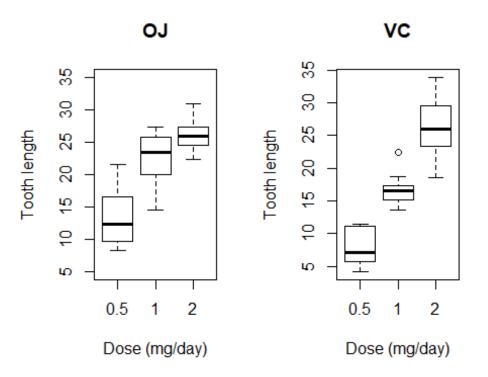
Plot box and whisker diagram

```
par(mfrow = c(1,2))
plot(len ~ as.factor(dose), data = ToothGrowth, xlab = "Dose (mg/day)", ylab
= "Tooth length")
plot(len ~ supp, data = ToothGrowth, xlab = "Supplement", ylab = "Tooth length")
```



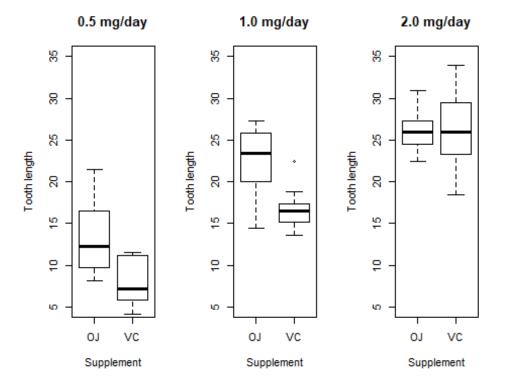
Plot box and whisker diagram according to supplement type subgroups

```
par(mfrow = c(1,2))
plot(len[ToothGrowth$supp == "0J"] ~ as.factor(dose[ToothGrowth$supp == "0J"]), data = ToothGrowth, xlab = "Dose (mg/day)", ylab = "Tooth length", main = "0J", ylim = c(5,35))
plot(len[ToothGrowth$supp == "VC"] ~ as.factor(dose[ToothGrowth$supp == "0J"]), data = ToothGrowth, xlab = "Dose (mg/day)", ylab = "Tooth length", main = "VC")
```



Plot box and whisker diagram according to dose subgroups

```
par(mfrow = c(1,3))
plot(len[ToothGrowth$dose == 0.5] ~ supp[ToothGrowth$dose == 0.5], data = Too
thGrowth, xlab = "Supplement", ylab = "Tooth length", main = "0.5 mg/day", yl
im = c(5, 35))
plot(len[ToothGrowth$dose == 1.0] ~ supp[ToothGrowth$dose == 1.0], data = Too
thGrowth, xlab = "Supplement", ylab = "Tooth length", main = "1.0 mg/day", yl
im = c(5, 35))
plot(len[ToothGrowth$dose == 2.0] ~ supp[ToothGrowth$dose == 2.0], data = Too
thGrowth, xlab = "Supplement", ylab = "Tooth length", main = "2.0 mg/day", yl
im = c(5, 35))
```



3. Hypothesis tests to compare tooth growth by supp and dose.

3-1. Supplement

See whether the test meets equal variance assumption

```
var.test(len~supp, data = ToothGrowth)

##

## F test to compare two variances

##

## data: len by supp

## F = 0.6386, num df = 29, denom df = 29, p-value = 0.2331

## alternative hypothesis: true ratio of variances is not equal to 1

## 95 percent confidence interval:

## 0.3039488 1.3416857

## sample estimates:

## ratio of variances

## 0.6385951
```

No significant difference in variance are shown.

Perform t.test function to compare two supplement groups

```
t.test(len~supp, data = ToothGrowth, paired = FALSE, var.equal = TRUE)
```

```
##
## Two Sample t-test
##
## data: len by supp
## t = 1.9153, df = 58, p-value = 0.06039
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.1670064 7.5670064
## sample estimates:
## mean in group OJ mean in group VC
## 20.66333 16.96333
```

3-2. Dose

See whether the test meets equal variance assumption

```
bartlett.test(len~as.factor(dose), data = ToothGrowth)

##

## Bartlett test of homogeneity of variances

##

## data: len by as.factor(dose)

## Bartlett's K-squared = 0.66547, df = 2, p-value = 0.717
```

No significant difference in variance are shown.

Perform and function to compare three dose groups

Then, which group shows significant difference in tooth length?

```
TukeyHSD(ANOVA)
    Tukey multiple comparisons of means
##
##
      95% family-wise confidence level
##
## Fit: aov(formula = len ~ as.factor(dose), data = ToothGrowth)
## $`as.factor(dose)`
##
           diff
                      lwr
                                upr
                                       p adj
## 1-0.5 9.130 5.901805 12.358195 0.00e+00
## 2-0.5 15.495 12.266805 18.723195 0.00e+00
## 2-1 6.365 3.136805 9.593195 4.25e-05
```

4. conclusions

As shown above, we can conclude

- 1. Tooth length does not differ according to supplement delivery type.
- 2. Tooth length does differ among the three dose levels of vitamin C.