

# Basic-inferential-data-analysis.R

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```
setwd("C:\\Users\\Mahesha\\Desktop\\Desktop\\Data_Science\\Courseera\\Statistical Inference")
getwd()
```

```
## [1] "C:/Users/Mahesha/Desktop/Desktop/Data_Science/Courseera/Statistical Inference"
```

```
library(ggplot2)
```

```
# Load ToothGrowth data
data("ToothGrowth")
```

```
# Display a summary of the data
summary(ToothGrowth)
```

```
##           len           supp           dose
##  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
##  Max.   :33.90                      Max.   :2.000
```

```
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
```

```
#Unique Values
unique(ToothGrowth$len)
```

```
## [1] 4.2 11.5 7.3 5.8 6.4 10.0 11.2 5.2 7.0 16.5 15.2 17.3 22.5 13.6
## [15] 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 17.6
## [29] 9.7 8.2 9.4 19.7 20.0 25.2 25.8 21.2 27.3 22.4 24.5 24.8 30.9 29.4
## [43] 23.0
```

```
unique(ToothGrowth$supp)
```

```
## [1] VC OJ
## Levels: OJ VC
```

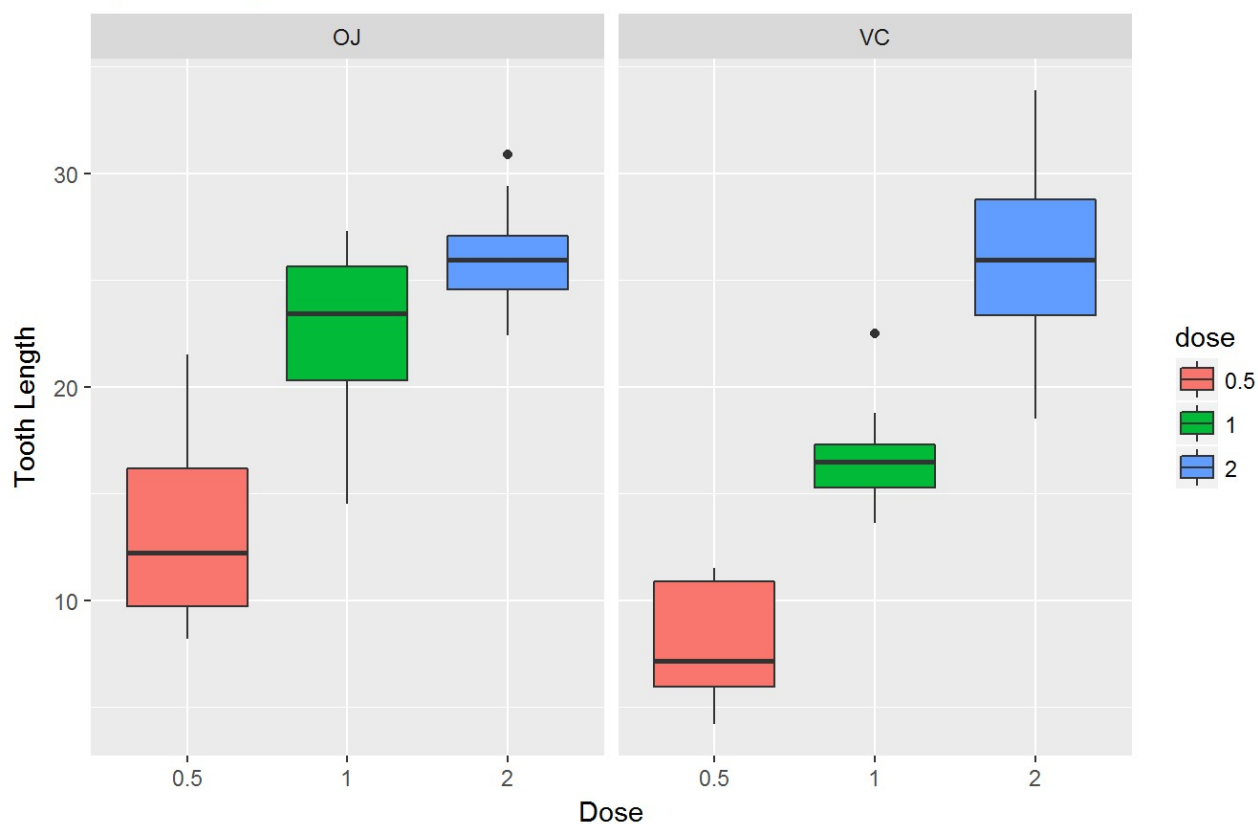
```
unique(ToothGrowth$dose)
```

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

```
# Convert dose to a factor
ToothGrowth$dose<-as.factor(ToothGrowth$dose)

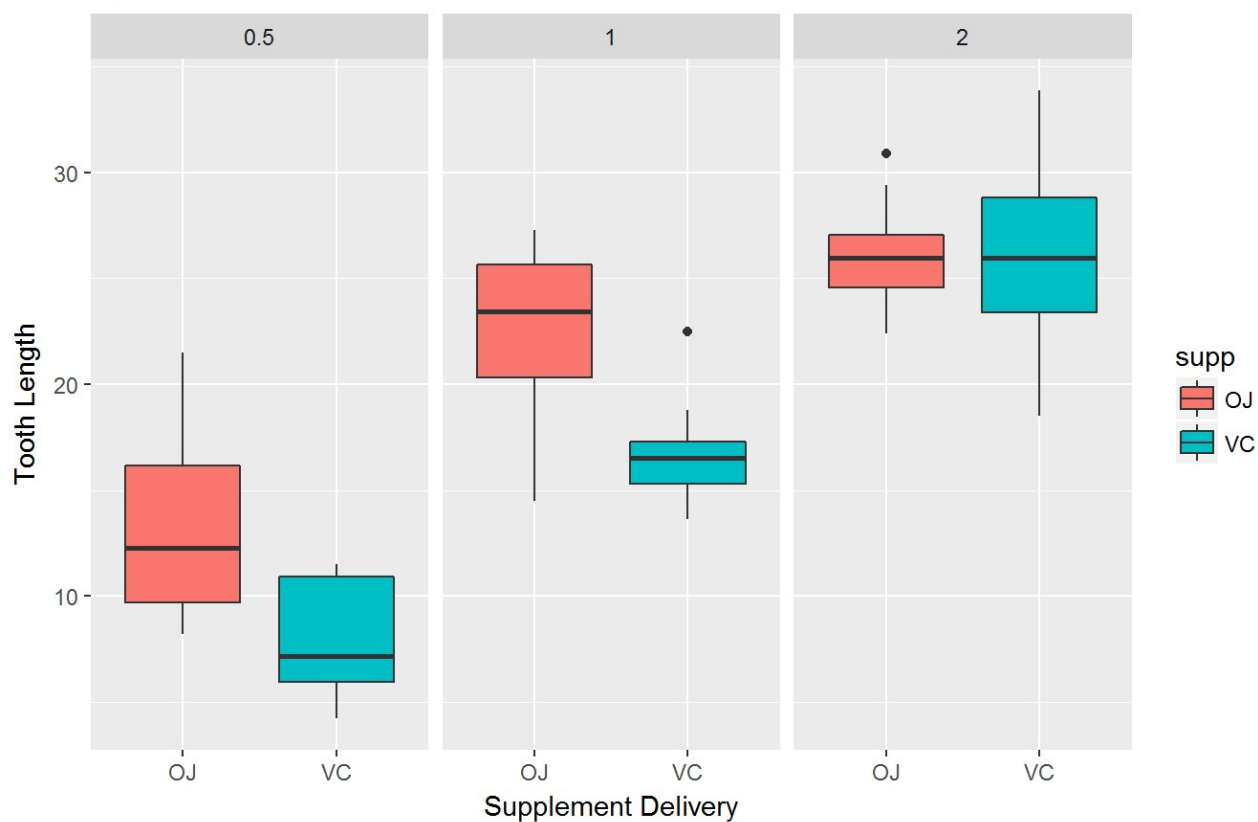
# Plot tooth length ('len') vs. the dose amount ('dose'), broken out by supplement delivery method ('supp')
ggplot(aes(x=dose, y=len), data=ToothGrowth) + geom_boxplot(aes(fill=dose)) + xlab("Dose") + ylab("Tooth Length") + facet_grid(~ supp) + ggtitle("Tooth Length vs. Dose Amount \nby Delivery Method") + theme(plot.title = element_text(lineheight=.9, face="bold"))
```

## Tooth Length vs. Dose Amount by Delivery Method



```
# Plot tooth length ('len') vs. supplement delivery method ('supp') broken out  
by the dose amount ('dose')  
ggplot(aes(x=supp, y=len), data=ToothGrowth) + geom_boxplot(aes(fill=supp)) + x  
lab("Supplement Delivery") + ylab("Tooth Length") + facet_grid(~ dose) + ggtitle  
("Tooth Length vs. Delivery Method \nby Dose Amount") +  
theme(plot.title = element_text(lineheight=.9, face="bold"))
```

## Tooth Length vs. Delivery Method by Dose Amount



```
# run t-test to compare tooth growth by supplement
t.test(len~supp,data=ToothGrowth)
```

```
##
##  Welch Two Sample t-test
##
## data:  len by supp
## t = 1.9153, df = 55.309, p-value = 0.06063
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -0.1710156  7.5710156
## sample estimates:
## mean in group OJ mean in group VC
##      20.66333      16.96333
```

*#Since the p-value is greater than 0.05 and the confidence interval of the test contains zero we can say that supplement types seems to have no impact on Tooth growth based on this test.*

*#Now we'll compare tooth growth by dose, looking at the different pairs of dose values.*

*# run t-test using dose amounts 0.5 and 1.0*

```
ToothGrowth_sub <- subset(ToothGrowth, ToothGrowth$dose %in% c(1.0,0.5))
t.test(len~dose,data=ToothGrowth_sub)
```

```
##
##  Welch Two Sample t-test
##
## data:  len by dose
## 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 in group 0.5    mean in group 1
##           10.605           19.735
```

*# run t-test using dose amounts 0.5 and 2.0*

```
ToothGrowth_sub <- subset(ToothGrowth, ToothGrowth$dose %in% c(0.5,2.0))
t.test(len~dose,data=ToothGrowth_sub)
```

```
##
##  Welch Two Sample t-test
##
## data:  len by dose
## t = -11.799, df = 36.883, p-value = 4.398e-14
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -18.15617 -12.83383
## sample estimates:
## mean in group 0.5    mean in group 2
##           10.605           26.100
```

*# run t-test using dose amounts 1.0 and 2.0*

```
ToothGrowth_sub <- subset(ToothGrowth, ToothGrowth$dose %in% c(1.0,2.0))
t.test(len~dose,data=ToothGrowth_sub)
```

```
##
## Welch Two Sample t-test
##
## data: len by dose
## t = -4.9005, df = 37.101, p-value = 1.906e-05
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -8.996481 -3.733519
## sample estimates:
## mean in group 1 mean in group 2
##          19.735          26.100
```

*#As can be seen, the p-value of each test was essentially zero and the confidence interval of each test does not cross over zero (0).*

*#Based on this result we can assume that the average tooth length increases with an increasing dose, and therefore the null hypothesis can be rejected.*

*#Conclusions*

*#Given the following assumptions:*

*#1. The sample is representative of the population*

*#2. The distribution of the sample means follows the Central Limit Theorem*

*#In reviewing our t-test analysis from above, we can conclude that supplement delivery method has no effect on tooth growth/length, however increased dosages do result in increased tooth length.*