

# Statistical Inference Course Project

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August 27, 2018

## Part 2: Basic Inferential Data Analysis Instructions

Now 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 perform some basic exploratory data analysis

```
library(datasets)
data(ToothGrowth)
library(ggplot2)

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 0.5 ...
```

```
head(ToothGrowth)
```

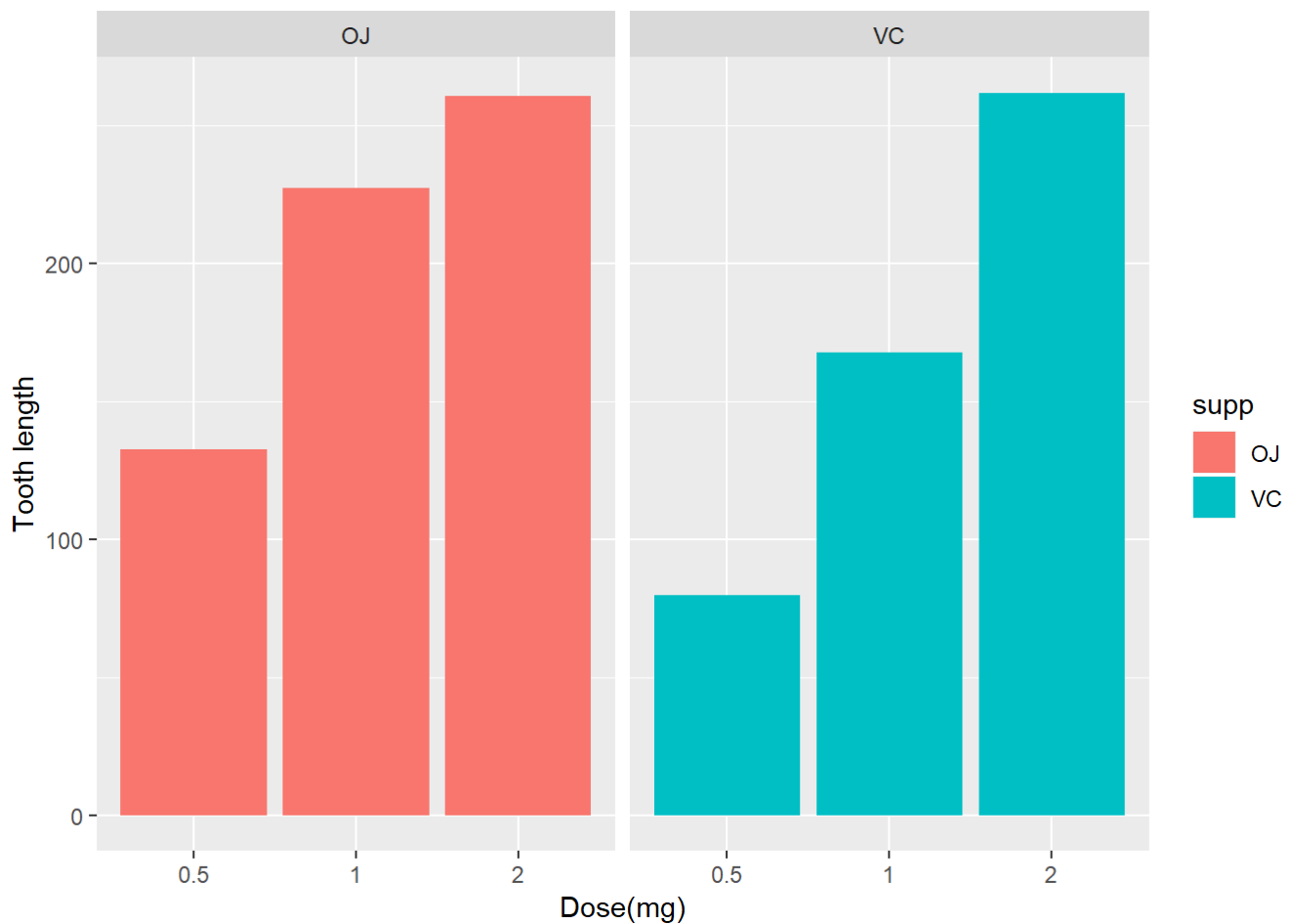
```
##   len supp dose
## 1  4.2   VC  0.5
## 2 11.5   VC  0.5
## 3  7.3   VC  0.5
## 4  5.8   VC  0.5
## 5  6.4   VC  0.5
## 6 10.0   VC  0.5
```

2. Provide a basic 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
```

```
ggplot(data=ToothGrowth, aes(x=as.factor(dose), y=len, fill=supp)) +
  geom_bar(stat="identity") +
  facet_grid(. ~ supp) +
  xlab("Dose(mg)") +
  ylab("Tooth length")
```



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)

```
hypothesis1 <- t.test(len ~ supp, data = ToothGrowth)
hypothesis1$conf.int
```

```
## [1] -0.1710156 7.5710156
## attr(,"conf.level")
## [1] 0.95
```

```
hypothesis1$p.value
```

```
## [1] 0.06063451
```

```
hypothesis2<-t.test(len ~ supp, data = subset(ToothGrowth, dose == 0.5))
hypothesis2$conf.int
```

```
## [1] 1.719057 8.780943
## attr(,"conf.level")
## [1] 0.95
```

```
hypothesis2$p.value
```

```
## [1] 0.006358607
```

```
hypoth3<-t.test(len ~ supp, data = subset(ToothGrowth, dose == 1))  
hypoth3$conf.int
```

```
## [1] 2.802148 9.057852  
## attr(,"conf.level")  
## [1] 0.95
```

```
hypoth3$p.value
```

```
## [1] 0.001038376
```

```
hypoth4<-t.test(len ~ supp, data = subset(ToothGrowth, dose == 2))  
hypoth4$conf.int
```

```
## [1] -3.79807 3.63807  
## attr(,"conf.level")  
## [1] 0.95
```

```
hypoth4$p.value
```

```
## [1] 0.9638516
```

## Conclusions

OJ ensures more tooth growth than VC for dosages 0.5 & 1.0. OJ and VC gives the same amount of tooth growth for dose amount 2.0 mg/day. For the entire trial we cannot conclude OJ is more effective than VC for all scenarios.