

ToothGrowth_Basic Inferential Data Analysis

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Overview

This part is going to analyze the ToothGrowth data in the R datasets package.

1. Load the ToothGrowth data, and provide a basic summary of the data.
2. Perform some basic exploratory data analyses.
3. Use confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose.
4. State your conclusions and the assumptions needed for your conclusions.

Loading the data and exploratory data analyses

```
data(ToothGrowth); attach(ToothGrowth)
```

Basic Summary of the Data

```
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 ...
```

```
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
```

```
unique(dose)
```

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

```
table(supp, dose)
```

```
##      dose
## supp 0.5  1  2
##   OJ  10 10 10
##   VC  10 10 10
```

```
ToothGrowth %>% group_by(supp, factor(dose)) %>% summarise(mean_len = mean(len), sd_len = sd(len))
```

```
## Source: local data frame [6 x 4]
```

```
## Groups: supp [?]
```

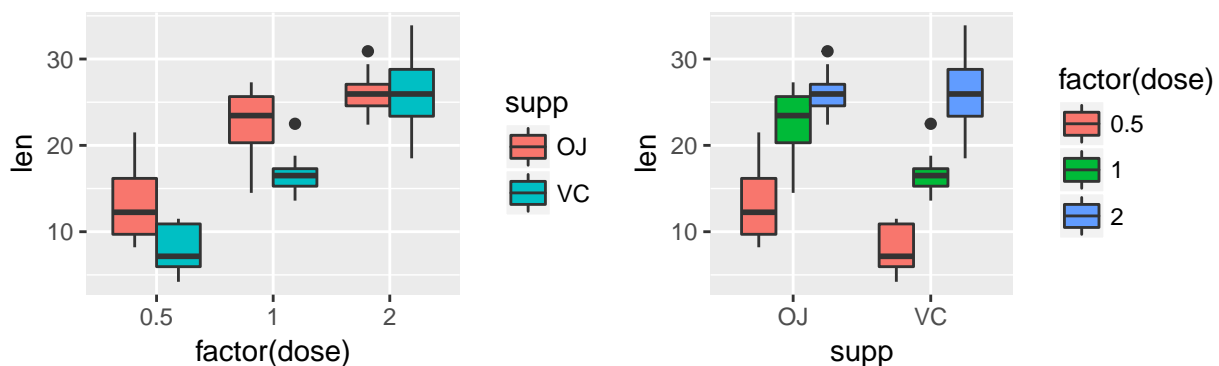
```
##
```

```
##      supp `factor(dose)` mean_len sd_len
```

```
##      <fctr>          <fctr>    <dbl>    <dbl>
## 1      OJ            0.5      13.23  4.459709
## 2      OJ            1        22.70  3.910953
## 3      OJ            2        26.06  2.655058
## 4      VC            0.5       7.98  2.746634
## 5      VC            1        16.77  2.515309
## 6      VC            2        26.14  4.797731
```

Exploratory Data Analysis

```
gg1<-ggplot(ToothGrowth, aes(x = factor(dose), y = len, fill = supp)) + geom_boxplot()
gg2 <- ggplot(ToothGrowth, aes(x = supp, y = len, fill = factor(dose))) + geom_boxplot()
grid.arrange(gg1, gg2, ncol = 2) # or interaction2wt(len ~ supp*factor(dose))
```



Confidence Interval and Hypothesis Testing

- Examining the impact of **supp** or **dose** as a factor separately.
- Examining the impact of supp, dose and the interaction of supp and dose by including both **supp** and **dose** as factors.

```
dose15 <- ToothGrowth %>% filter(dose %in% c("0.5", "1"))
dose12 <- ToothGrowth %>% filter(dose %in% c("1", "2"))
dose52 <- ToothGrowth %>% filter(dose %in% c("0.5", "2"))
t.test(len ~ supp)
```

```
##
## 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
t.test(len ~ factor(dose), data = dose15)
```

```
##
## Welch Two Sample t-test
```

```
##
## data: len by factor(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
t.test(len ~ factor(dose), data = dose12)

##
## Welch Two Sample t-test
##
## data: len by factor(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
t.test(len ~ factor(dose), data = dose52)

##
## Welch Two Sample t-test
##
## data: len by factor(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
# or summary(aov(len ~ dose)) and summary(aov(len ~ supp*factor(dose)))
detach(ToothGrowth)
```

Conclusion

- This is a balanced design(equal sample sizes in each cell of the design).
- The confidence interval is [-0.171, 7.571], which include zero. There is no significant correlation between delivery methods and tooth length.
- The confidence interval is [-11.98, -6.276] for doses 0.5 and 1.0, [-18.16, -12.83] for doses 0.5 and 2.0, and [-8.996, -3.734] for doses 1.0 and 2.0) and show that there is a significant correlation between tooth length and dose levels.
- T-test shows that two delivery methods (OJ or VC) have equal tooth length ($p > 0.05$). Three levels of dose aren't all equally effective. Increasing the dose level can lead to increased tooth growth.
- Both main effects and the interaction between these factors are significant.