# Stat-Inference Course Project B - Tooth Growth

Steven

## 1 - Synopsis

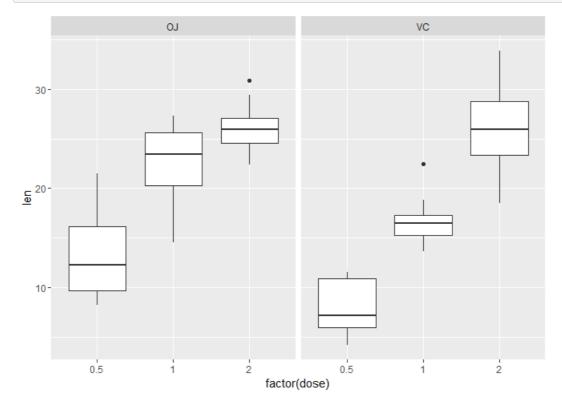
In this analysis, we're going to analyze the ToothGrowth data in the R datasets package, summarise the data and sse confidence intervals and hypothesis tests to compare tooth growth by supp and dose.

#### 2 - Data Summarization

```
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
```

The dose column is num type now, we will treat it as a factor kind variable. Let's have a look at their distribution.

```
g <- ggplot(ToothGrowth, aes(factor(dose), len))
g + geom_boxplot() + facet_grid(. ~ supp)</pre>
```



### 3 - Hypothesis Test

Now we are going to compare tooth growth by supply and dose. There is 3 kind of dose and 2 kind of supply. From the above analysis, the average len value of OJ may be larger than VC's. So we make the following hypothesis (with omega0) as the mean of OJ and the omega0):

```
\(H_0\): \(\mu_0\) = \(\mu_1\)\(H_a\): \(\mu_0\) > \(\mu_1\)
```

## data: OJ.5 and VC.5

we will compare our hypothesis under all three levels of dose.

## t = 3.1697, df = 14.969, p-value = 0.003179

```
df <- ToothGrowth
OJ.5 <- df[df$dose==0.5,]$len[11:20]
VC.5 <- df[df$dose==0.5,]$len[1:10]
t.test(OJ.5, VC.5, alternative = "greater", var.equal = FALSE)

##
## Welch Two Sample t-test
##</pre>
```

```
OJ1 <- df[df$dose==1,]$len[11:20]
VC1 <- df[df$dose==1,]$len[1:10]
t.test(OJ1, VC1, alternative = "greater", var.equal = FALSE)
```

```
OJ2 <- df[df$dose==2,]$len[11:20]
VC2 <- df[df$dose==2,]$len[1:10]
t.test(OJ2, VC2, alternative = "greater", var.equal = FALSE)
```

```
# t.test(df[df$dose==0.5 && df$supp=="OJ",]$len, df[df$dose==0.5 && df$supp=="VC",]$len)
```

From the three tests above, we can see the len if OJ is larger than VC's under the 0.5 and 1.0 levels. However, under the 2.0 level, they are almost the same.

## 4 - Summary

This analysis is under the guidance of statistical inference course on Coursera.

```
sessionInfo()
```

```
## R version 3.2.5 (2016-04-14)
  ## Platform: x86 64-w64-mingw32/x64 (64-bit)
  ## Running under: Windows 10 x64 (build 10586)
  ##
  ## locale:
  ## [1] LC COLLATE=Chinese (Simplified) China.936
  ## [2] LC CTYPE=Chinese (Simplified) China.936
  ## [3] LC_MONETARY=Chinese (Simplified)_China.936
  ## [4] LC NUMERIC=C
  ## [5] LC_TIME=Chinese (Simplified)_China.936
  ##
  ## attached base packages:
  ## [1] stats graphics grDevices utils datasets methods base
  ## other attached packages:
  ## [1] ggplot2 2.1.0
  \#\,\#
  ## loaded via a namespace (and not attached):
 ## [1] Rcpp_0.12.7 digest_0.6.10 plyr_1.8.4 grid_3.2.5
## [5] gtable_0.2.0 formatR_1.4 magrittr_1.5 evaluate_0.9
## [9] scales_0.4.0 stringi_1.1.2 reshape2_1.4.1 rmarkdown_1.0
## [13] labeling_0.3 tools_3.2.5 stringr_1.1.0 munsell_0.4.3
## [17] yaml_2.1.13 colorspace_1.2-6 htmltools_0.3.5 knitr_1.14
Loading [Contrib]/a11y/accessibility-menu.js
```