### Statistical Inference: Tooth Growth Analysis

Henrique Souza (github.com/htssouza)

#### Goals

1-) Load the ToothGrowth data and perform some basic exploratory data analyses 2-) Provide a basic summary of the data 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) 4-) State your conclusions and the assumptions needed for your conclusions

## 1-) Load the ToothGrowth data and perform some basic exploratory data analyses

```
# Load data
data (ToothGrowth)
# Data summary
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 ...
   # A small sample of the data
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.

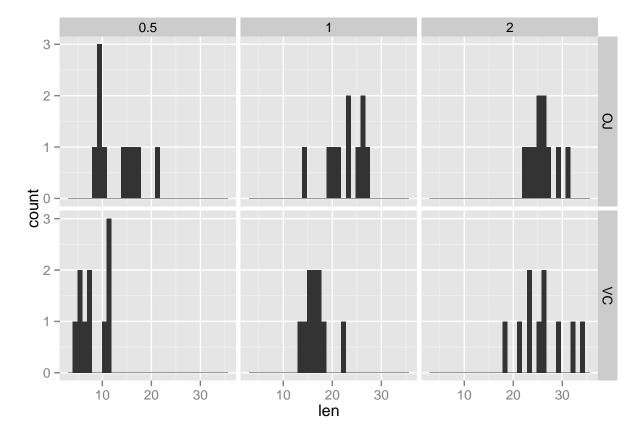
Dose is a factor (few different valyes) so make the preprocessing conversion:

```
ToothGrowth$dose <- as.factor(ToothGrowth$dose)
```

Plotting some histograms:

```
library(plyr)
library(ggplot2)
qplot(
  len,
  data = ToothGrowth,
  facets = .~supp~dose,
  geom="histogram")
```

```
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
```



# 3-) Use confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose.

Summarizing data to compute confidence intervals (using Student T interval):

```
## supp dose mean sd n
```

```
## 1
           0.5 13.23 4.459709 10
## 2
       OJ
             1 22.70 3.910953 10
## 3
       OJ
             2 26.06 2.655058 10
           0.5 7.98 2.746634 10
## 4
       VC
## 5
       VC
             1 16.77 2.515309 10
## 6
       VC
             2 26.14 4.797731 10
```

Computing confidence intervals (95% of confidence):

```
summaries$min <- summaries$mean - (qt(.975, (summaries$n-1) * summaries$sd / sqrt(summaries$n)))
summaries$max <- summaries$mean + (qt(.975, (summaries$n-1) * summaries$sd / sqrt(summaries$n)))
summaries</pre>
```

```
supp dose mean
                           sd n
## 1
           0.5 13.23 4.459709 10 11.064300 15.39570
## 2
             1 22.70 3.910953 10 20.502166 24.89783
## 3
      OJ
             2 26.06 2.655058 10 23.730220 28.38978
      VC
           0.5 7.98 2.746634 10 5.664568 10.29543
      VC
## 5
             1 16.77 2.515309 10 14.415961 19.12404
## 6
      VC
             2 26.14 4.797731 10 23.990112 28.28989
```

#### 4-) Conclusions

Based on the summary above we can say (with 95% of sure that):

- lens when using OJ (.5 dose) > VC (.5 dose)
- lens when using OJ (1.0 dose) > VC (1.0 dose)
- lens when using 2.0 of both OJ and VC are similar/equivalent

The assumptions are:

- current dataset (small, with only 60 rows)
- 95% of confidence