Title Analysis of the ToothGrowth Data from R Datasets Package

Overview

This project explores the *ToothGrowth* data from the R *datasets* library by performing some preliminary analysis on the data and drawing some initial conclusions.

Exploratory Data Analysis

```
library(dplyr)  # Data manipulation (filter, mutate, group_by, etc.)
library(ggplot2)  # Plotting (qplot, ggplot etc.)
library(knitr)  # Dynamic Report Creation
library(datasets)  # R included sample data sets
data(ToothGrowth)
```

To setup the analysis we load the required libraries and the *ToothGrowth* data set. Per the R documentation for the *datasets* package [1], *ToothGrowth* is described as follows:

The response is the length of odontoblasts (teeth) in each of 10 guinea pigs at each of three dose levels of Vitamin C (0.5, 1, and 2 mg) with each of two delivery methods (orange juice or ascorbic acid).

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 2 2 ...
## $ dose: num 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
                   VC:30
                           1st Qu.:0.500
  1st Qu.:13.07
                           Median :1.000
## Median :19.25
## Mean
         :18.81
                           Mean :1.167
## 3rd Qu.:25.27
                           3rd Qu.:2.000
## Max.
          :33.90
                           Max. :2.000
```

table(ToothGrowth\$supp)

```
## UJ VC ## 30 30
```

```
table(ToothGrowth$dose)
```

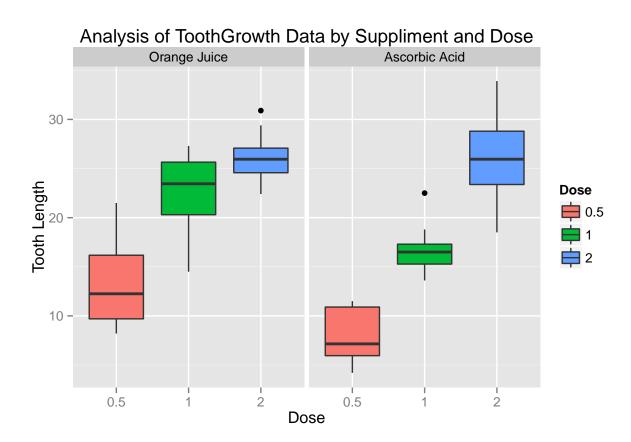
```
##
## 0.5
             2
        1
   20 20 20
table(ToothGrowth %>% select(supp, dose))
##
       dose
## supp 0.5 1 2
##
     OJ 10 10 10
     VC 10 10 10
##
## Transformation/Renaming for clarity
names(ToothGrowth) = c("Length", "Suppliment", "Dose")
ToothGrowth$Dose = factor(ToothGrowth$Dose)
ToothGrowth$Suppliment <- factor(</pre>
    ToothGrowth$Suppliment
    ,labels = c("Orange Juice", "Ascorbic Acid")
)
tgBySuppDose <- ToothGrowth %>%
    group_by(Suppliment, Dose) %>%
    summarize(Length = mean(Length)) %>%
    arrange(Suppliment, Dose, desc(Length))
kable(tgBySuppDose
      ,format = "markdown"
      ,caption = "ToothGrowth by Suppliment and Dose"
)
```

Suppliment	Dose	Length
Orange Juice	0.5	13.23
Orange Juice	1	22.70
Orange Juice	2	26.06
Ascorbic Acid	0.5	7.98
Ascorbic Acid	1	16.77
Ascorbic Acid	2	26.14

The exploratory analysis of the ToothGrowth data frame demonstrates that the data consist of 30 observations each of supp OJ (Orange Juice) and VC (Ascorbic Acid); the 30 observations of each supp consist of 10 observations each of the three dose levels (0.5, 1 and 2).

Data Summary

Confidence Intervals and Hypothesis Test



Conclusions

Reference

 $[1] \ https://stat.ethz.ch/R-manual/R-devel/library/datasets/html/ToothGrowth.html$