## Statistical Inference Project, Part 2

First we will load the ToothGrowth built-in dataset that examines the effect of Vitamin C on tooth growth in guinea pigs

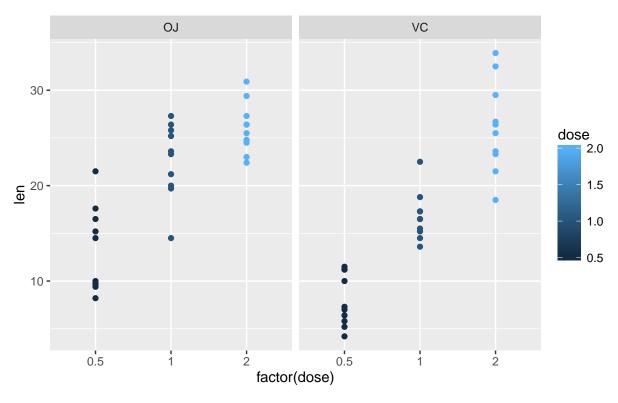
```
data("ToothGrowth")
summary(ToothGrowth)
##
                                 dose
         len
                    supp
          : 4.20
                    OJ:30
                                   :0.500
##
                            Min.
    Min.
                            1st Qu.:0.500
                    VC:30
##
   1st Qu.:13.07
##
  Median :19.25
                            Median :1.000
          :18.81
                                  :1.167
  Mean
                            Mean
   3rd Qu.:25.27
                            3rd Qu.:2.000
           :33.90
                            Max. :2.000
  Max.
library("dplyr")
library("ggplot2")
library("knitr")
```

kable(summarise(grouped,min=min(len),max=max(len),median=median(len),mean=mean(len),sd=sd(len)))

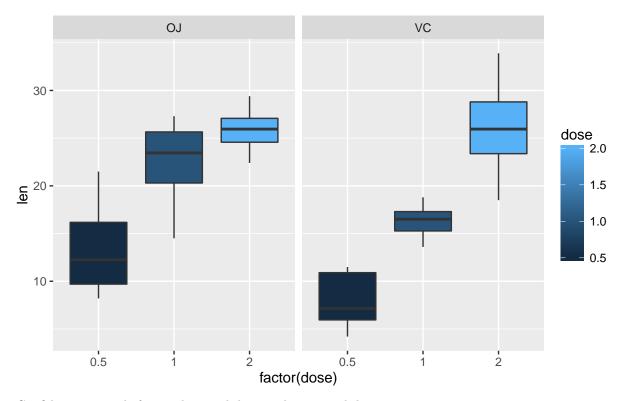
| supp | dose | min  | max  | median | mean  | $\operatorname{sd}$ |
|------|------|------|------|--------|-------|---------------------|
| OJ   | 0.5  | 8.2  | 21.5 | 12.25  | 13.23 | 4.459708            |
| OJ   | 1.0  | 14.5 | 27.3 | 23.45  | 22.70 | 3.910953            |
| OJ   | 2.0  | 22.4 | 30.9 | 25.95  | 26.06 | 2.655058            |
| VC   | 0.5  | 4.2  | 11.5 | 7.15   | 7.98  | 2.746634            |
| VC   | 1.0  | 13.6 | 22.5 | 16.50  | 16.77 | 2.515309            |
| VC   | 2.0  | 18.5 | 33.9 | 25.95  | 26.14 | 4.797731            |

grouped = group\_by(ToothGrowth, supp, dose)

```
ggplot(ToothGrowth,aes(x=factor(dose),y=len)) +
  geom_point(aes(color=dose)) +
  facet_wrap(~ supp)
```



```
ggplot(ToothGrowth,aes(x=factor(dose),y=len)) +
  geom_boxplot(outlier.shape = NA,aes(fill=dose)) +
  facet_wrap(~ supp)
```



Confidence intervals for tooth growth by supplement and dose

```
tests = list()
tests[['supp']] = t.test(len~supp, data=ToothGrowth,paired=FALSE)
for (d in c(0.5,1,2)) {
   tests[[toString(d)]] = t.test(len~supp, data=filter(ToothGrowth,dose == d),paired=FALSE)
}
options(digits=2)
kable(sapply(tests,function(t){
   c(t$estimate[1],
        t$estimate[2],
        ci.lower = t$conf.int[1],
        ci.upper = t$conf.int[2],
        p.value = t$p.value)
}))
```

|                  | supp  | 0.5   | 1    | 2     |
|------------------|-------|-------|------|-------|
| mean in group OJ | 20.66 | 13.23 | 22.7 | 26.06 |
| mean in group VC | 16.96 | 7.98  | 16.8 | 26.14 |
| ci.lower         | -0.17 | 1.72  | 2.8  | -3.80 |
| ci.upper         | 7.57  | 8.78  | 9.1  | 3.64  |
| p.value          | 0.06  | 0.01  | 0.0  | 0.96  |

From these results, it appears that regarding supplement type we can not conclude whether the type of supplement at all doses can predict the magnitude of tooth growth since the p-value is close to 0.5 and the confidence interval includes 0 in its range. This can be seen in the graph since the distribution of length as a function of supplement type overlaps for OJ and VC.

Regarding dosage, it can be confirmed that at a dosage of 0.5 or 1.0, supplement type can predict the magnitude of growth. Specifically there could be a link between OJ and greater tooth growth relative to VC. This same conclusion can not be made for a dosage of 2.0 since the p-value is greater than 0.05. The graphs confirm this conclusion since at 0.5 and 1.0 dosages the observed growth range was not overlapping between the supplements.