Basic Inferential Data Analysis

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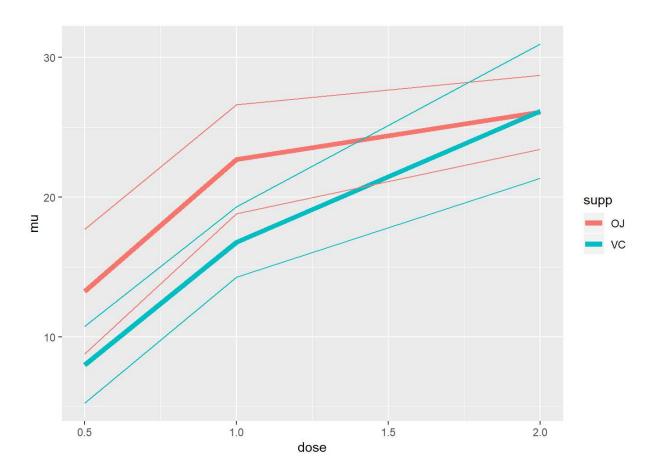
Data description

The response is the length of odontoblasts (cells responsible for tooth growth) in 60 guinea pigs. Each animal received one of three dose levels of vitamin C (0.5, 1, and 2 mg/day) by one of two delivery methods, orange juice or ascorbic acid (a form of vitamin C and coded as VC).

```
##
        len
                   supp
                                dose
##
   Min.
          : 4.20
                   OJ:30
                          Min.
                                  :0.500
##
   1st Qu.:13.07
                   VC:30
                          1st Qu.:0.500
                           Median :1.000
##
   Median :19.25
         :18.81
                           Mean
                                :1.167
##
   Mean
   3rd Qu.:25.27
                           3rd Qu.:2.000
##
          :33.90
                                :2.000
##
   Max.
                           Max.
```

It looks like there are two dimensions (supplement and dosage) and one meassurement (length). Let's take a look at the groups combinations:

```
## # A tibble: 6 x 5
## # Groups:
              supp [2]
     supp
           dose subj_count
                             mu
    <fct> <dbl>
##
                   <int> <dbl> <dbl>
## 1 OJ
            0.5
                      10 13.2 4.46
## 2 OJ
            1
                       10 22.7
                                 3.91
## 3 OJ
            2
                       10 26.1
                                2.66
## 4 VC
## 5 VC
            0.5
                       10 7.98 2.75
            1
                       10 16.8
                                 2.52
## 6 VC
                       10 26.1
                                4.80
```



Null Hypotesis - The supplement is more irrelevant the bigger dosage is.

```
OJ05 <- filter(ToothGrowth, supp == "OJ" & dose == 0.5)
OJ10 <- filter(ToothGrowth, supp == "OJ" & dose == 1.0)
OJ20 <- filter(ToothGrowth, supp == "OJ" & dose == 2.0)
VC05 <- filter(ToothGrowth, supp == "VC" & dose == 0.5)
VC10 <- filter(ToothGrowth, supp == "VC" & dose == 1.0)
VC20 <- filter(ToothGrowth, supp == "VC" & dose == 2.0)
```

Dose 0.5

```
t.test(0J05$len, VC05$len)
```

```
##
## Welch Two Sample t-test
##
## data: 0J05$len and VC05$len
## t = 3.1697, df = 14.969, p-value = 0.006359
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 1.719057 8.780943
## sample estimates:
## mean of x mean of y
## 13.23 7.98
```

Dose 1.0

```
t.test(OJ10$len, VC10$len)
```

```
##
## Welch Two Sample t-test
##
## data: OJ10$len and VC10$len
## t = 4.0328, df = 15.358, p-value = 0.001038
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 2.802148 9.057852
## sample estimates:
## mean of x mean of y
## 22.70 16.77
```

Dose 2.0

```
t.test(0J20$len, VC20$len)
```

```
##
## Welch Two Sample t-test
##
## data: OJ20$len and VC20$len
## t = -0.046136, df = 14.04, p-value = 0.9639
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -3.79807 3.63807
## sample estimates:
## mean of x mean of y
## 26.06 26.14
```

Conclustion

As we can see Dose 0.5 and Dose 1.0 have significant difference in teeth growth with p_values 0.0064 and 0.001 respectively. But picture totally changes for Dose 2.0 where it's doesn't matter what supplements we are using the progress is pretty similar with p-value 0.934. The supplement used for the teeth grows is less relevant the bigger dosage. OJ gives more precise result with less variance then VC.