part2

Saurabh Yadav

PART-2 Basic inferential data analysis

Loading the tooth data

```
data("ToothGrowth")
library(ggplot2)
```

```
## Warning: package 'ggplot2' was built under R version 3.5.1
```

Subsetting the tooth growth data on the basis of the supp

```
vc<-subset(ToothGrowth, supp=="VC")
oj<-subset(ToothGrowth, supp=="0J")</pre>
```

Arranging the datasets on the basis of doses

```
newvc<-c()
newvc$dose0.5<-vc$len[vc$dose==0.5]
newvc$dose1<-vc$len[vc$dose==1]
newvc$dose2<-vc$len[vc$dose==2]
newvc<-data.frame(newvc)
newoj<-c()
newoj$dose0.5<-oj$len[oj$dose==0.5]
newoj$dose1<-oj$len[oj$dose==1]
newoj$dose2<-oj$len[oj$dose==2]
newoj<-data.frame(newoj)</pre>
```

Taking the mean of the length of the tooth growth for different doses

```
lengthmeanvc<-apply(newvc, 2, mean)
lengthmeanoj<-apply(newoj, 2, mean)</pre>
```

converting it into a marix

```
lengthmeanvc<-data.frame(matrix(lengthmeanvc))
lengthmeanoj<-data.frame(matrix(lengthmeanoj))</pre>
```

giving new column names

```
colnames(lengthmeanvc)<-"len"
rownames(lengthmeanvc)<-c("dose0.5", "dose1", "dose2")
colnames(lengthmeanoj)<-"len"
rownames(lengthmeanoj)<-c("dose0.5", "dose1", "dose2")</pre>
```

adding new coloumn

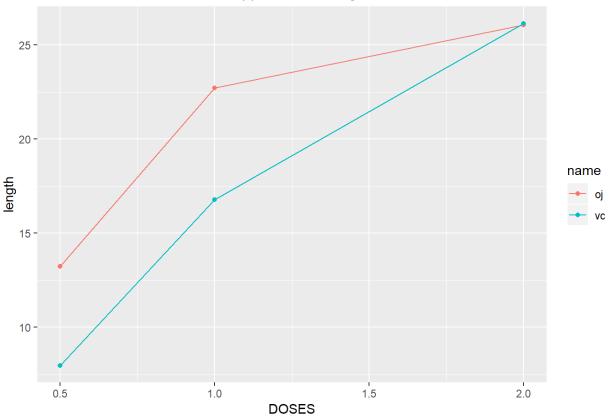
```
lengthmeanvc$dose<-c(0.5,1,2)
lengthmeanvc$name<-c("vc","vc","vc")
lengthmeanoj$dose<-c(0.5,1,2)</pre>
```

```
lengthmeanoj$name<-c("oj","oj","oj")
finaldata<-rbind(lengthmeanoj,lengthmeanvc)</pre>
```

plotting

```
\label{thm:color=name} ggplot(final data, aes(x=dose, y=len, color=name)) + geom\_point() + geom\_line() + labs(title="effect of doses of different supplies on tooth growth") + labs(x="DOSES", y="length")
```

effect of doses of different supplies on tooth growth



```
len<-lengthmeanvc
colnames(len)<-c("lenVC", "dose")
len$len0J<-lengthmeanoj$len</pre>
```

Using confidence intervals and/or hypothesis tests to compare tooth growth by supp and dose

1. Confidence interval for dose 0.5

```
t.test(oj$len[oj$dose==0.5], vc$len[vc$dose==0.5], paired = F)$conf.int

## [1] 1.719057 8.780943
## attr(,"conf.level")
## [1] 0.95
```

RESULT: The t test between the supp vc and oj for dose=0.5 gives the interval 1.719057 to 8.780943. Which means the dose of 0.5 for oj is way more effective for tooth growth.

2. Confidence interval for dose 1

```
t.test(oj$len[oj$dose==1],vc$len[vc$dose==1],paired = F)$conf.int
```

```
## [1] 2.802148 9.057852
## attr(,"conf.level")
## [1] 0.95
```

RESULT: The t test between the supp vc and oj for dose=1 gives the interval 2.802148 to 9.057852. Which means the dose of 1 for oj is way more effective for tooth growth.

3. Confidence interval for dose 2

```
t.test(oj$len[oj$dose==2],vc$len[vc$dose==2],paired = F)$conf.int
```

```
## [1] -3.79807 3.63807
## attr(,"conf.level")
## [1] 0.95
```

RESULT: The t test between the supp vc and oj for dose=2 gives the interval -3.79807 to 3.63807. Which means the dose of 2 for both vc and oj are approximately same. This can also be shown by the plot which clearly shows that dose 2 for both affect the tooth growth in a same way.

Conclusion

- 1. The effect of two medications are similar when the dose is 2.
- 2. The oj is more effective in tooth growth when the doses are 0.5 and 1