

# Homework #11

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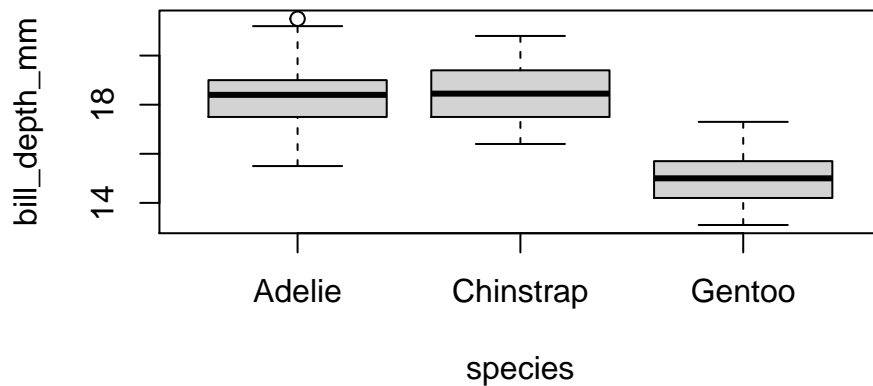
10 May 2021

## Chapter 12

### Problem 04

- a. Looks like the data satisfies ANOVA assumptions

```
pen<-palmerpenguins::penguins  
boxplot(bill_depth_mm ~ species, data=pen)
```



- b. p-value  $< 2.2 \times 10^{-16}$ , showing significance at the  $\alpha = 0.01$  level. Reject the null, species does significantly predict bill depth.

```
depth_mod<-lm(bill_depth_mm ~ species, data=pen)  
anova(depth_mod)
```

```
## Analysis of Variance Table  
##  
## Response: bill_depth_mm  
##           Df Sum Sq Mean Sq F value    Pr(>F)  
## species     2  903.97   451.98  359.79 < 2.2e-16 ***  
## Residuals 339  425.87     1.26  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

### Problem 05

P-value: 0.00362, reject the null. There are significant differences in the predictions of 300g objects by age group.

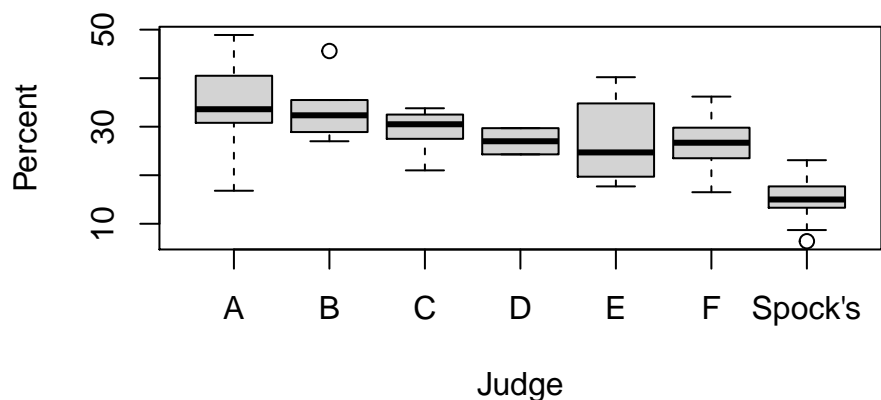
```
anova(lm(mean300 ~ age, data=fosdata::weight_estimate))

## Analysis of Variance Table
##
## Response: mean300
##          Df Sum Sq Mean Sq F value    Pr(>F)
## age         3  60786  20262.1    4.9018 0.00362 **
## Residuals  76 314155   4133.6
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

### Problem 06

- a. Spock's Lawyers may have a point, as Spock's mean percentage is visibly less than the means of the other 7 lawyers.

```
judge_perc <- Sleuth3::case0502
boxplot(Percent ~ Judge, data=judge_perc)
```



- b. P-value: 0.3239, do not reject the null. The means of the percentage of women jurors in the 6 other judges courts are significantly similar.

```
judge_perc %>% filter(Judge != "Spock's") %>% lm(Percent ~ Judge, data=.) %>% anova()

## Analysis of Variance Table
##
## Response: Percent
##          Df Sum Sq Mean Sq F value    Pr(>F)
## Judge      5  326.46   65.292    1.2183 0.3239
## Residuals  31 1661.33   53.591
```

- c. The significantly similar means of the other judges justifies comparing Spock's judge to the other 6. P-value:  $1.303 \times 10^{-6}$ , reject the null. There is significant difference in the expected percentage of women jurors in Spock's judge's court when compared to the other judges.

```
judge_perc %>% mutate(Spock = (Judge=="Spock's")) %>%  
  t.test(Percent ~ Spock, data=.)
```

```
##  
## Welch Two Sample t-test  
##  
## data: Percent by Spock  
## t = 7.1597, df = 17.608, p-value = 1.303e-06  
## alternative hypothesis: true difference in means is not equal to 0  
## 95 percent confidence interval:  
## 10.49935 19.23999  
## sample estimates:  
## mean in group FALSE mean in group TRUE  
## 29.49189 14.62222
```

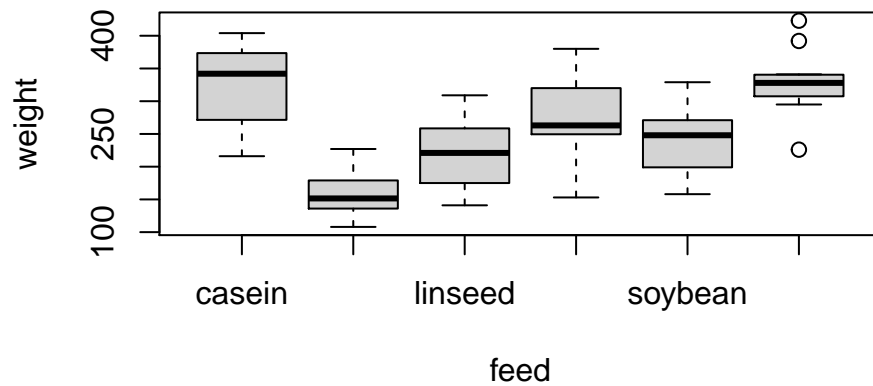
### Problem 08

The p-value:  $5.93 \times 10^{-10}$  looks to show significant difference in weight based on feed, however the boxplot shows inconsistent variance in the different groups with the most extreme being when you compare "meatmeal" with "sunflower".

```
anova(lm(weight ~ feed, data=chickwts))
```

```
## Analysis of Variance Table  
##  
## Response: weight  
##          Df Sum Sq Mean Sq F value    Pr(>F)  
## feed      5 231129   46226  15.365 5.936e-10 ***  
## Residuals 65 195556    3009  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
boxplot(weight ~ feed, data=chickwts)
```



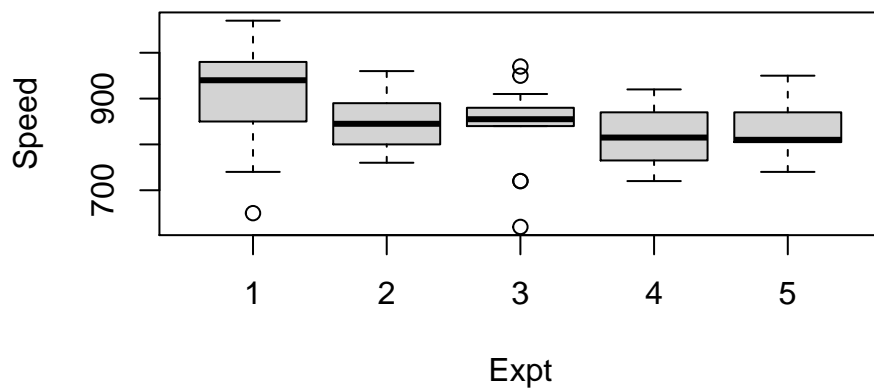
### Problem 09

With P-value: 0.0004827, it looks like there is significant difference in the mean based on experiment number. However, looking at the boxplots the data does not satisfy the anova assumptions, especially experiments 3 and 5.

```
anova(lm(Speed ~ Expt, data=morley))
```

```
## Analysis of Variance Table
##
## Response: Speed
##          Df Sum Sq Mean Sq F value    Pr(>F)
## Expt      1  72581    72581   13.041 0.0004827 ***
## Residuals 98 545444     5566
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

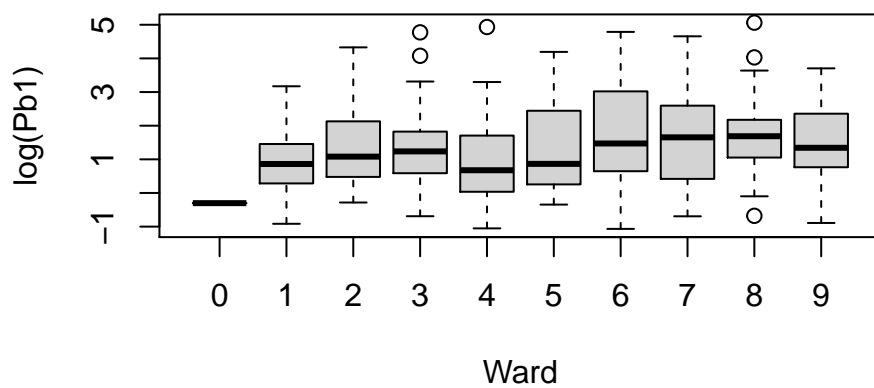
```
boxplot(Speed ~ Expt, data=morley)
```



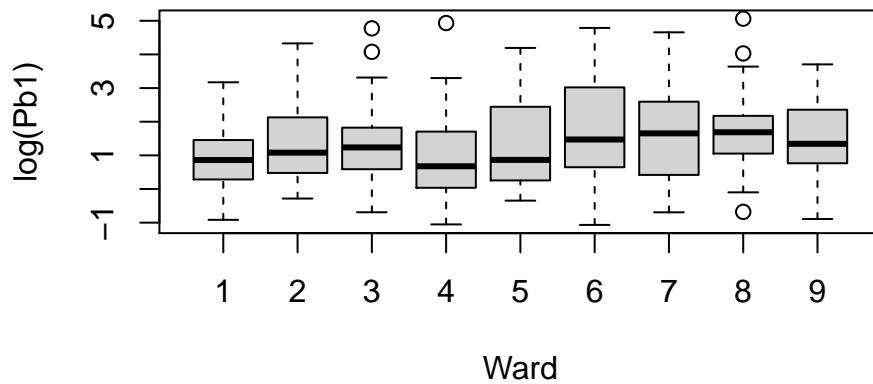
### Problem 15

a.

```
flint_pb <- fosdata::flint
boxplot(log(Pb1) ~ Ward, data=flint_pb)
```



```
flint_pb <- flint_pb %>% filter(Ward != 0)
boxplot(log(Pb1) ~ Ward, data=flint_pb)
```



b. The log levels look normally distributed and the variances are similar enough to justify anova.  
c. P-value: 0.03137, there is a significant difference in the mean of the log of the amount of lead in the water based on ward in Flint, MI.

```
anova(lm(log(Pb1) ~ Ward, data=flint_pb))
```

```
## Analysis of Variance Table
##
## Response: log(Pb1)
##           Df Sum Sq Mean Sq F value    Pr(>F)
## Ward         1   7.76   7.7563    4.6815 0.03137 *
## Residuals 268 444.03   1.6568
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```