

Julia Vineyard

Worked with: Andrew Gordon, Jessica Bonin

Lab 10 ANOVA By Hand

1. require(here)

```
rope= read.csv(here("data", "rope.csv"))
```

```
rm(list = ls())
```

```
rope = read.csv(here("data", "rope.csv"))  
rope$rope.type = factor(rope$rope.type)  
levels(rope$rope.type)
```

```
n_obs = nrow(rope)  
n_obs  
n_groups = length(levels(rope$rope.type))  
n_groups
```

```
mean_cut= mean(rope$p.cut)  
resids_cut= rope$p.cut - mean_cut  
ss_tot= sum(resids_cut^2)  
ss_tot  
df_tot = n_obs -1  
df_tot
```

```
agg_resids= aggregate(x=rope$p.cut, by=list(rope$rope.type),  
  FUN= function(x) {x-mean(x)})  
str(agg_resids)
```

```
agg_sq_resids = aggregate(x=rope$p.cut, by=list(rope$rope.type),  
  FUN= function(x) {sum((x-mean(x))^2)})
```

```
agg_sq_resids  
str(agg_sq_resids)  
ss_within = sum(agg_sq_resids$x)  
ss_within  
df_within = n_obs - n_groups  
df_within
```

```
ss_among = ss_tot - ss_within  
ss_among  
df_among = n_groups - 1  
df_among
```

```
ms_within =ss_within / df_within
```

```
ms_among = ss_among / df_among
```

```
f_ratio = ms_among / ms_within
```

```
f_ratio #if a big number then reject the null that all means are the same
```

```
f_pval = pf(f_ratio, df_among, df_within, lower.tail= FALSE)
```

```
f_pval
```

```
fit_1= lm(p.cut ~ rope.type, data=rope)
```

```
anova_fit_1 = anova(fit_1)
```

```
str(anova_fit_1)
```

```
anova_fit_1$`Sum Sq`
```

2. Based on the boxplot I think that there are not equal variances among the groups.
3. p-value = 0.00143
4. No, an ANOVA analysis is not appropriate for the raw data since there is not homogeneity among the variances. This is supported by the boxplot and the p-value of the Bartlett test. The significant p-value means that we can reject the null hypothesis that the variance is the same amongst the groups.
5. BLAZE is the base case.
6. 0.36714. The base case rope is just the Estimate value of the Intercept.
7. $0.36714 - 0.10164 = 0.2655$ (Intercept Estimate + rope.typeXTC Estimate)