

Stat631 Extra Credit

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```
library(readr)
library(dplyr)
library(ggplot2)
library(phia)
library(emmeans)
library(car)

coffee <- read.table("coffee.txt")
head(coffee)
```

	Coffee	Setting	ReactionTime
1	Regular	Library	207
2	Decaf	Library	242
3	Regular	Home	234
4	Decaf	Home	236
5	Regular	Café	213
6	Decaf	Café	252

1. I would like this extra credit to be added to option 1: Midterm 2

2 (a)

2 factors: Coffee and Setting

Treatments: 2 (Coffee) x 3 (Setting) = 6 treatments.

```
coffee %>%
  group_by(Coffee, Setting) %>%
  summarise(n = n())
```

```
`summarise()` has grouped output by 'Coffee'. You can override using the  
.groups` argument.
```

```
# A tibble: 6 x 3  
# Groups: Coffee [2]  
  Coffee Setting n  
  <chr>   <chr>   <int>  
1 Decaf    Café      5  
2 Decaf    Home      5  
3 Decaf    Library   5  
4 Regular  Café      5  
5 Regular  Home     5  
6 Regular  Library  5
```

Replications per treatment = 5

There are 6 treatment groups, each with 5 replications, totaling 30 observations. Therefore, the design is balanced.

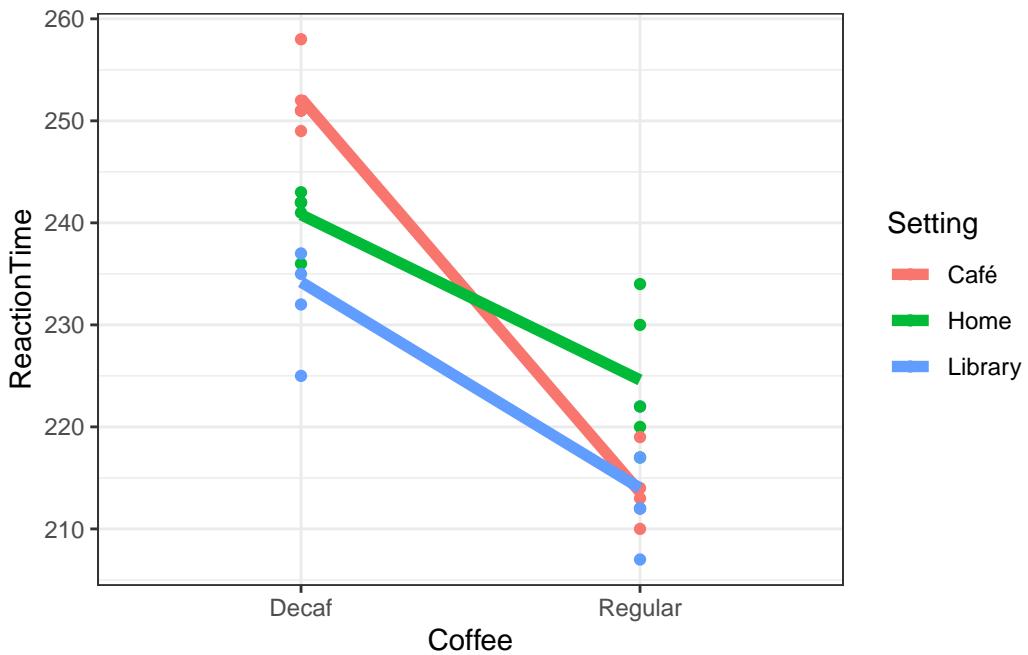
2 (b)

```
coffee <- within(coffee, {fCoffee = factor(Coffee); fSetting = factor(Setting)})  
head(coffee)  
  
  Coffee Setting ReactionTime fSetting fCoffee  
1 Regular  Library        207  Library  Regular  
2 Decaf    Library        242  Library  Decaf  
3 Regular  Home          234   Home   Regular  
4 Decaf    Home          236   Home   Decaf  
5 Regular  Café          213   Café   Regular  
6 Decaf    Café          252   Café   Decaf  
  
model1 <- lm(ReactionTime ~ fCoffee*fSetting, data = coffee)  
lsmFR <- lsmeans(model1, ~ fCoffee*fSetting)  
lsmFR  
  
fCoffee fSetting lsmean    SE df lower.CL upper.CL  
Decaf    Café      252 2.26 24      248      257  
Regular  Café      214 2.26 24      209      218
```

Decaf	Home	241	2.26	24	236	245
Regular	Home	225	2.26	24	220	229
Decaf	Library	234	2.26	24	230	239
Regular	Library	214	2.26	24	209	219

Confidence level used: 0.95

```
ggplot(coffee, aes(x = Coffee, y = ReactionTime, color = Setting)) +
  geom_point() +
  stat_summary(fun = mean, geom = "line",
               aes(group = Setting), linewidth = 2) +
  theme_bw()
```



The interaction plot suggests there may be an interaction between the type of coffee type and study location. Specifically, the effect of coffee type on reaction time varies across different settings. This effect appears to be strongest in the Café, where the difference in reaction times between Decaf and Regular coffee is most pronounced. The differences are smaller in the Home and Library setting. Since the lines in the plot are not parallel, this indicates that the effect of coffee type is not consistent across the different study settings.

2 (c)

```
anova(aov(ReactionTime ~ fCoffee * fSetting, data = coffee))
```

Analysis of Variance Table

Response: ReactionTime

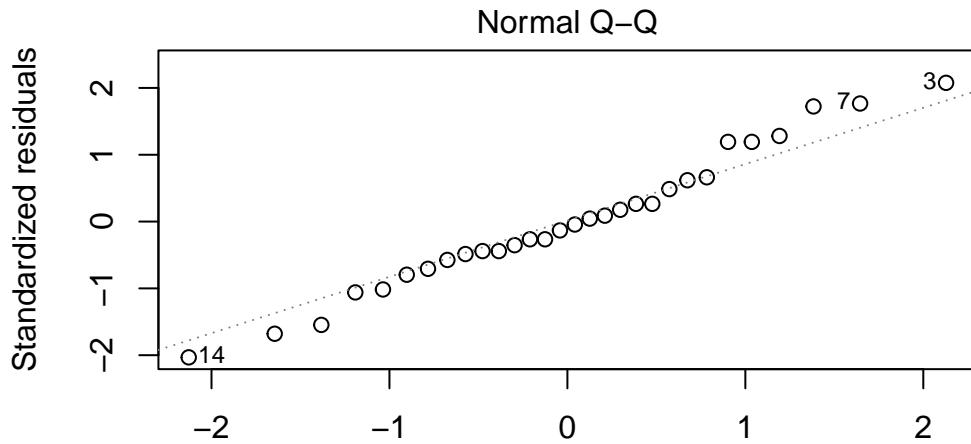
	Df	Sum Sq	Mean Sq	F value	Pr(>F)
fCoffee	1	4687.5	4687.5	182.986	1.007e-12 ***
fSetting	2	504.8	252.4	9.853	0.0007517 ***
fCoffee:fSetting	2	713.6	356.8	13.928	9.657e-05 ***
Residuals	24	614.8	25.6		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

From the ANOVA table all terms in the model are statistically significant. The main effect of coffee is highly significant which indicates that the reaction time is significantly different between Decaf and Regular coffee. We also can see that the main effect of setting is also significant. This shows us that study setting has an impact on reaction time as well. Finally, the interaction between Coffee and Setting is also significant. This confirms the effect of Coffee type and it depends on the study setting which does align with what we saw in the interaction plot. This matches our interaction plot above.

2 (d)

```
# Q-Q plot
plot(model1, 2)
```



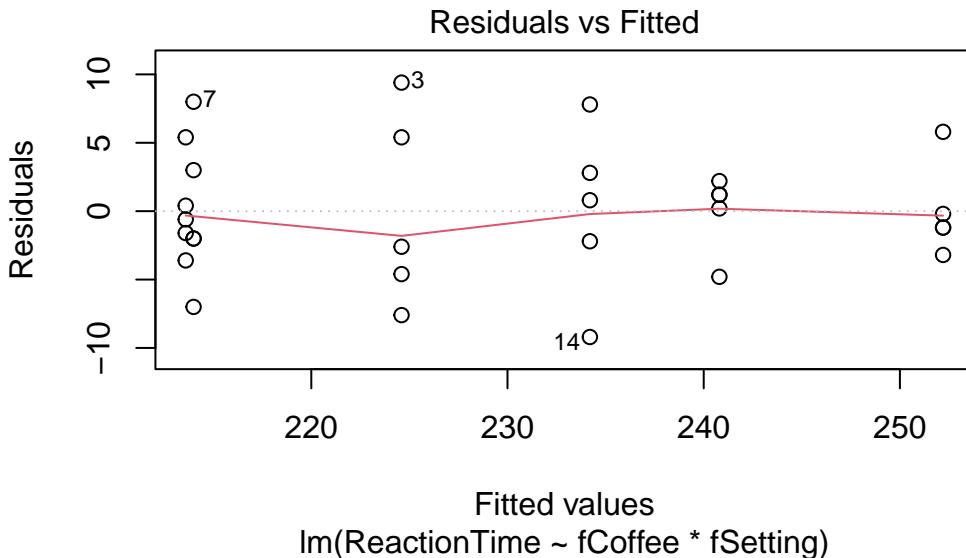
Theoretical Quantiles
 $\text{lm}(\text{ReactionTime} \sim \text{fCoffee} * \text{fSetting})$

```
# Shapiro test for Normality
shapiro.test(model1$residuals) # p-value > 0.05 means that Normality satisfied
```

```
Shapiro-Wilk normality test
```

```
data: model1$residuals
W = 0.97976, p-value = 0.8192
```

```
# Residuals vs Fitted values
plot(model1, 1)
```



```
# Levene Test
leveneTest(ReactionTime ~ fCoffee*fSetting, data = coffee) # p-value > 0.05 Equal Variance
```

```
Levene's Test for Homogeneity of Variance (center = median)
Df F value Pr(>F)
group 5 0.9256 0.4817
24
```

The Normal Q-Q plot shows that most of the residuals follow the diagonal line which indicates the normality assumption is satisfied. While there is some slight deviation at the upper end with a few outliers, the pattern is not too concerning. The Shapiro-Wilk test confirms this with a p-value of 0.8192, which is greater than our significance level of $\alpha = 0.05$ which suggest that we fail to reject the null hypothesis of normality. The residuals vs. Fitted plot shows no clear pattern or funnel shape; residuals appear to be randomly scattered with constant spread across fitted values. The levene's test also confirms this with a p-value of 0.4817, which again is greater than our significance level of $\alpha = 0.05$ Equal variance assumption is satisfied. Thus the ANVOA model and results from part c are valid and reliable.

2 (e)

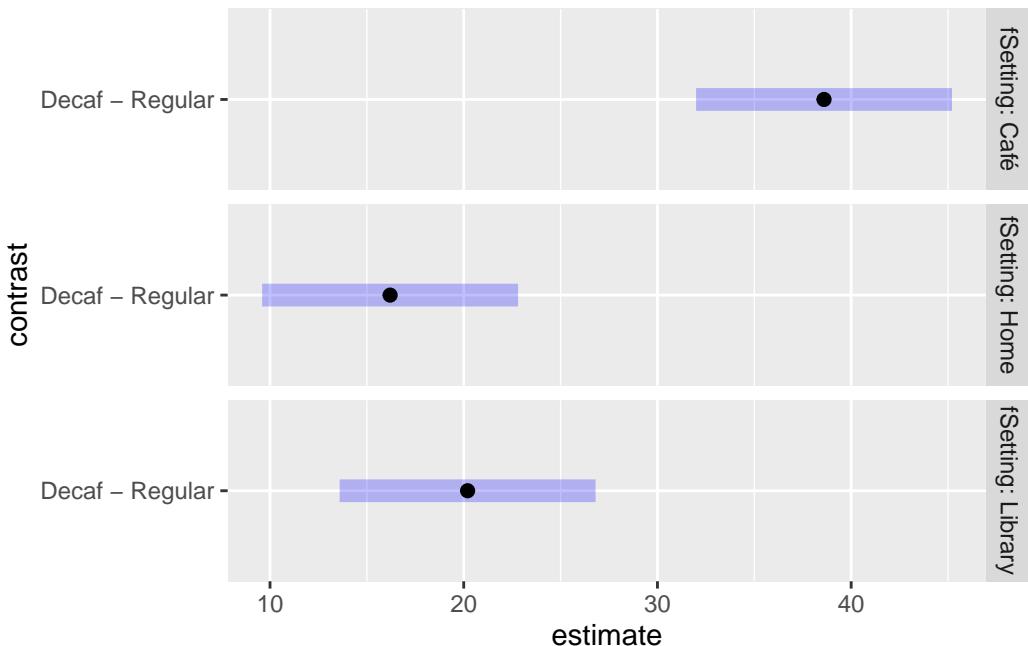
```
em <- emmeans(model1, pairwise ~ fCoffee | fSetting)
em$contrasts

fSetting = Café:
contrast      estimate SE df t.ratio p.value
Decaf - Regular     38.6 3.2 24   12.059  <.0001

fSetting = Home:
contrast      estimate SE df t.ratio p.value
Decaf - Regular     16.2 3.2 24    5.061  <.0001

fSetting = Library:
contrast      estimate SE df t.ratio p.value
Decaf - Regular     20.2 3.2 24    6.310  <.0001

plot(em$contrasts)
```



From the pairwise comparison plot, we observe that in all three settings students who consumed decaf coffee had significantly slower reaction times compared to those who drank regular

coffee. The largest difference occurred in the Cafe setting, with a difference of 38.6 milliseconds. These results align with the earlier findings from both the interaction plot and the ANOVA table, confirming a significant interaction between coffee type and study setting on reaction time.