

STAT430 Assignment3

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October 22, 2019

Question 5.8

```
chemical_data <- data.frame(Temperature=factor(c(rep(150,6), rep(160,6),
                                                rep(170,6))),
                             Pressure=factor(rep(c(200,200,215,215,230,230),3)),
                             Chem=c(90.4,90.2,90.7,90.6,90.2,90.4,90.1,90.3,
                                       90.5,90.6,89.9,90.1,90.5,90.7,90.8,90.9,
                                       90.4,90.1))
```

(a) First denote a effects model:

$$y_{ijk} = \mu + \tau_i + \beta_j + (\tau\beta)_{ij} + \epsilon_{ijk}$$

where τ_i is the effect of i th level of the row factor, β_j is the effect of j th level of the column factor, $(\tau\beta)_{ij}$ is the effect of the interaction between τ_i and β_j , and ϵ_{ijk} is a random error component. And k is the numbers of replicate in each factor.

Now, we are interested in testing hypotheses about the equality of row treatment effects by stating:

$$H_0 : \tau_1 = \tau_2 = \tau_3$$

$$H_a : \text{at least one } \tau_i \text{ is different than the other, where } i \in \{1, 2, 3\}$$

Similarly, we can set our hypotheses for column treatment effects and interaction effects between row and column treatments.

```

anova(lm(Chem~Temperature*Pressure,data=chemical_data))

## Analysis of Variance Table
##
## Response: Chem
##
##           Df  Sum Sq Mean Sq F value    Pr(>F)
## Temperature      2  0.30111  0.15056   8.4687 0.0085392 **
## Pressure          2  0.76778  0.38389  21.5937 0.0003673 ***
## Temperature:Pressure  4  0.06889  0.01722   0.9687 0.4700058
## Residuals        9  0.16000  0.01778
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Based on the result, we can see that the main effects of temperature and pressure are significant since we have both p-values less than 0.05. However, we see that the interaction effect is not significant as we obtained a p-value of $0.47 > 0.05$. Since We know both treatments are significant, we can further perform a LSD test to see which level of treatments are different.

```

library(agricolae)
# Fisher LSD: Compare the temperature regardless of pressure.
LSD.test(chemical_data$Chem, chemical_data$Temperature, DError=9,
          MSError=0.01778, console=T)

##
## Study: chemical_data$Chem ~ chemical_data$Temperature
##
## LSD t Test for chemical_data$Chem
##
## Mean Square Error:  0.01778
##

```

```

## chemical_data$Temperature, means and individual ( 95 %) CI
##
##      chemical_data.Chem      std r      LCL      UCL  Min  Max
## 150          90.41667 0.2041241 6 90.29352 90.53981 90.2 90.7
## 160          90.25000 0.2664583 6 90.12686 90.37314 89.9 90.6
## 170          90.56667 0.2943920 6 90.44352 90.68981 90.1 90.9
##
## Alpha: 0.05 ; DF Error: 9
## Critical Value of t: 2.262157
##
## least Significant Difference: 0.1741518
##
## Treatments with the same letter are not significantly different.
##
##      chemical_data$Chem groups
## 170          90.56667      a
## 150          90.41667     ab
## 160          90.25000      b

# Fisher LSD: Compare the pressure regardless of temperautre.
LSD.test(chemical_data$Chem, chemical_data$Pressure, DError=9,
         MSerror=0.01778, console=T)

##
## Study: chemical_data$Chem ~ chemical_data$Pressure
##
## LSD t Test for chemical_data$Chem
##
## Mean Square Error: 0.01778
##
## chemical_data$Pressure, means and individual ( 95 %) CI

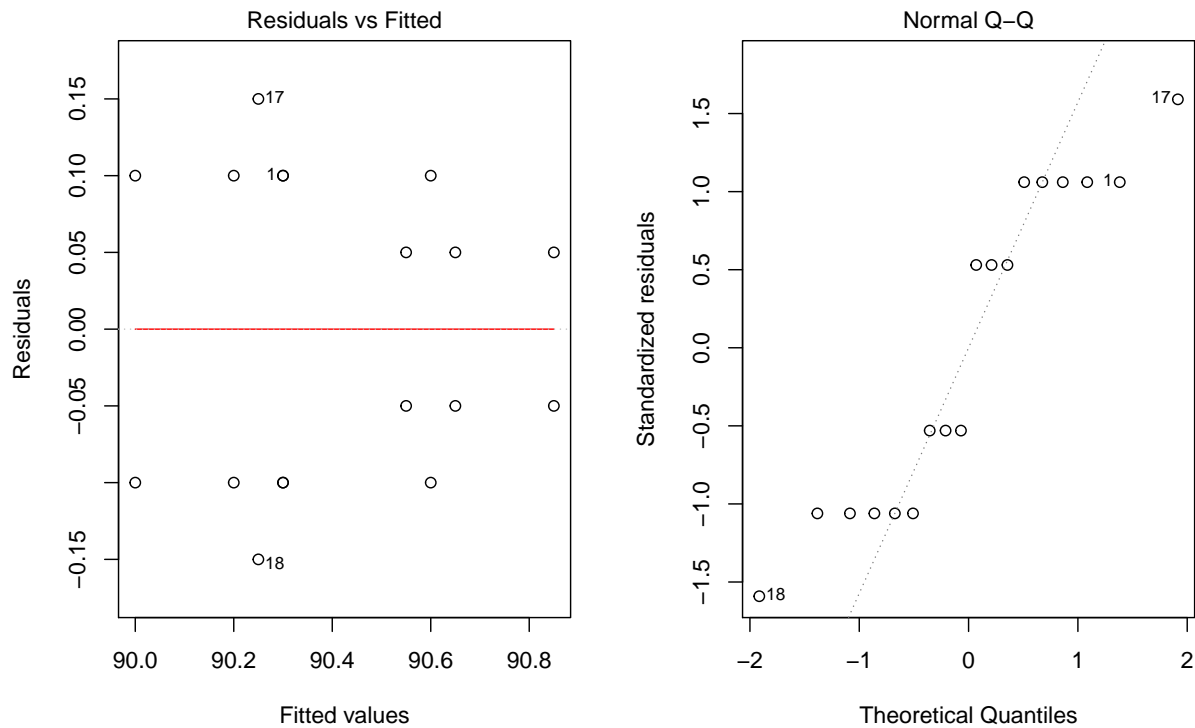
```

```
##
##      chemical_data.Chem      std r      LCL      UCL  Min  Max
## 200      90.36667 0.2160247 6 90.24352 90.48981 90.1 90.7
## 215      90.68333 0.1471960 6 90.56019 90.80648 90.5 90.9
## 230      90.18333 0.1940790 6 90.06019 90.30648 89.9 90.4
##
## Alpha: 0.05 ; DF Error: 9
## Critical Value of t: 2.262157
##
## least Significant Difference: 0.1741518
##
## Treatments with the same letter are not significantly different.
##
##      chemical_data$Chem groups
## 215      90.68333      a
## 200      90.36667      b
## 230      90.18333      c
```

From the LSD test results, we can see that temperature level of 160 °C is different than temperature level of 170 °C. On the other hand, we observe that all the pressure levels are significantly different than the other.

(b)

```
fit <- aov(lm(Chem~Temperature*Pressure,data=chemical_data))
par(mfrow=c(1,2))
plot(fit, 1); plot(fit, 2)
```



There is nothing unusual with the residual vs fitted value plot as since we can observe from the above plot that there is a constant band throughout the graph. The normal QQplot shows that many points are off the theoretical line. However, in order to check whether the normality assumption fails, we can perform a Shapiro-Wilk's test.

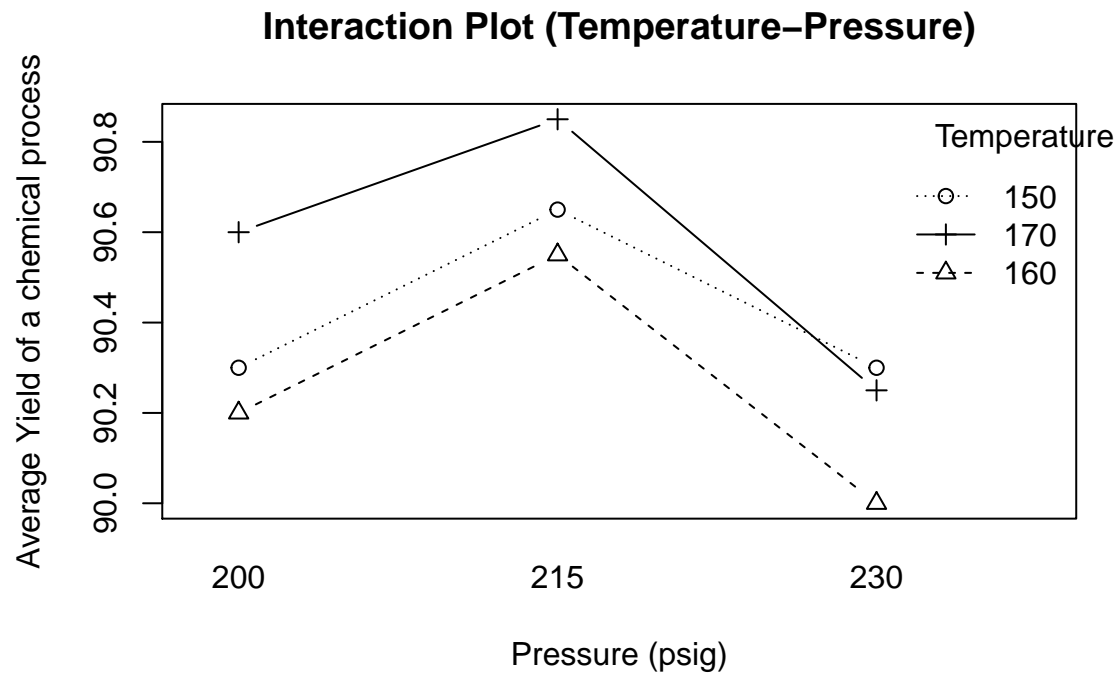
```
# H0: Data follows a normal distribution
shapiro.test(chemical_data$Chem)

##
##  Shapiro-Wilk normality test
##
## data:  chemical_data$Chem
## W = 0.97363, p-value = 0.8625
```

Based on the result, we can conclude that the assumption of normality is satisfied.

(c)

```
with(chemical_data, interaction.plot(x.factor=Pressure,
    trace.factor=Temperature, response=Chem, fun=mean,
    type="b", legend=T, pch=c(1,2,3), xlab="Pressure (psig)",
    ylab="Average Yield of a chemical process",
    main="Interaction Plot (Temperature-Pressure)"))
```



From the interaction graph, we can see that under the condition temperature of 170 °C and pressure of 215 psig, we will have the highest chemical yield.

Question 5.14

We can construct a effects model same as the previous question. For drill speed effect:

$$H_0 : \tau_1 = \tau_2$$

$$H_a : \tau_1 \text{ is different than } \tau_2$$

Similarly, we can set our hypotheses for column treatment effects and interaction effects between row and column treatments.

```
mechanic_data <- data.frame(Drill.Speed=factor(c(rep(125,8), rep(200,8))),
                             Feed.Rate=factor(rep(c(0.015,0.015,0.030,0.030,
                                                     0.045,0.045,0.060,0.060),2)),
                             Thrust.Force=c(2.70,2.78,2.45,2.49,2.60,2.72,2.75,
                                             2.86,2.83,2.86,2.85,2.80,2.86,2.87,
                                             2.94,2.88))

anova(lm(Thrust.Force~Drill.Speed*Feed.Rate, data=mechanic_data))

## Analysis of Variance Table
##
## Response: Thrust.Force
##
##           Df    Sum Sq  Mean Sq F value    Pr(>F)
## Drill.Speed      1 0.148225  0.148225  57.0096 6.605e-05 ***
## Feed.Rate        3 0.092500  0.030833  11.8590  0.002582 **
## Drill.Speed:Feed.Rate 3 0.041875  0.013958   5.3686  0.025567 *
## Residuals       8 0.020800  0.002600
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Based on the result, we can conclude that both treatments drill speed, feed rate and their interaction are significant since we have p-values less than 0.05. Therefore, we can perform LSD tests on both treatments and interaction.

```

# Fisher LSD: Compare the drill speed regardless of feed rate.
LSD.test(mechanic_data$Thrust.Force, mechanic_data$Drill.Speed, DError=8,
         MSError=0.0026, console=T)

##
## Study: mechanic_data$Thrust.Force ~ mechanic_data$Drill.Speed
##
## LSD t Test for mechanic_data$Thrust.Force
##
## Mean Square Error: 0.0026
##
## mechanic_data$Drill.Speed, means and individual ( 95 %) CI
##
##      mechanic_data.Thrust.Force      std r      LCL      UCL  Min  Max
## 125                2.66875 0.14327172 8 2.627178 2.710322 2.45 2.86
## 200                2.86125 0.04051014 8 2.819678 2.902822 2.80 2.94
##
## Alpha: 0.05 ; DF Error: 8
## Critical Value of t: 2.306004
##
## least Significant Difference: 0.0587918
##
## Treatments with the same letter are not significantly different.
##
##      mechanic_data$Thrust.Force groups
## 200                2.86125      a
## 125                2.66875      b

# Fisher LSD: Compare the feed rate regardless of drill speed.
LSD.test(mechanic_data$Thrust.Force, mechanic_data$Feed.Rate, DError=8,
         MSError=0.0026, console=T)

```



```
##
## Study: mechanic_data$Thrust.Force ~ mechanic_data$Feed.Rate
##
## LSD t Test for mechanic_data$Thrust.Force
##
## Mean Square Error: 0.0026
##
## mechanic_data$Feed.Rate, means and individual ( 95 %) CI
##
##      mechanic_data.Thrust.Force      std r      LCL      UCL  Min  Max
## 0.015                2.7925 0.06994045 4 2.733708 2.851292 2.70 2.86
## 0.03                 2.6475 0.20661962 4 2.588708 2.706292 2.45 2.85
## 0.045                2.7625 0.12816006 4 2.703708 2.821292 2.60 2.87
## 0.06                 2.8575 0.07932003 4 2.798708 2.916292 2.75 2.94
##
## Alpha: 0.05 ; DF Error: 8
## Critical Value of t: 2.306004
##
## least Significant Difference: 0.08314416
##
## Treatments with the same letter are not significantly different.
##
##      mechanic_data$Thrust.Force groups
## 0.06                2.8575      a
## 0.015               2.7925     ab
## 0.045               2.7625      b
## 0.03                2.6475      c
```

From the LSD test results, we can see that the two drill speed levels are significant different. And for feed rate, we observe that feed rate of 0.06 and rate of 0.045, rate of 0.06

and 0.03, and rate of 0.045 and 0.03 are significant different.

```
mechanic_data$Interact <- with(mechanic_data,
                                interaction(Drill.Speed, Feed.Rate))
int_fit <- aov(Thrust.Force~Interact, data=mechanic_data)
LSD.test(int_fit, "Interact", group=FALSE, console = TRUE)

##
## Study: int_fit ~ "Interact"
##
## LSD t Test for Thrust.Force
##
## Mean Square Error: 0.0026
##
## Interact, means and individual ( 95 %) CI
##
##          Thrust.Force      std r      LCL      UCL  Min  Max
## 125.0.015      2.740 0.056568542 2 2.656856 2.823144 2.70 2.78
## 125.0.03       2.470 0.028284271 2 2.386856 2.553144 2.45 2.49
## 125.0.045      2.660 0.084852814 2 2.576856 2.743144 2.60 2.72
## 125.0.06       2.805 0.077781746 2 2.721856 2.888144 2.75 2.86
## 200.0.015      2.845 0.021213203 2 2.761856 2.928144 2.83 2.86
## 200.0.03       2.825 0.035355339 2 2.741856 2.908144 2.80 2.85
## 200.0.045      2.865 0.007071068 2 2.781856 2.948144 2.86 2.87
## 200.0.06       2.910 0.042426407 2 2.826856 2.993144 2.88 2.94
##
## Alpha: 0.05 ; DF Error: 8
## Critical Value of t: 2.306004
##
## Comparison between treatments means
##
```

##	difference	pvalue	signif.	LCL	UCL
## 125.0.015 - 125.0.03	0.270	0.0007	***	0.1524164	0.387583601
## 125.0.015 - 125.0.045	0.080	0.1553		-0.0375836	0.197583601
## 125.0.015 - 125.0.06	-0.065	0.2382		-0.1825836	0.052583601
## 125.0.015 - 200.0.015	-0.105	0.0734	.	-0.2225836	0.012583601
## 125.0.015 - 200.0.03	-0.085	0.1341		-0.2025836	0.032583601
## 125.0.015 - 200.0.045	-0.125	0.0398	*	-0.2425836	-0.007416399
## 125.0.015 - 200.0.06	-0.170	0.0103	*	-0.2875836	-0.052416399
## 125.0.03 - 125.0.045	-0.190	0.0058	**	-0.3075836	-0.072416399
## 125.0.03 - 125.0.06	-0.335	0.0002	***	-0.4525836	-0.217416399
## 125.0.03 - 200.0.015	-0.375	0.0001	***	-0.4925836	-0.257416399
## 125.0.03 - 200.0.03	-0.355	0.0001	***	-0.4725836	-0.237416399
## 125.0.03 - 200.0.045	-0.395	0.0001	***	-0.5125836	-0.277416399
## 125.0.03 - 200.0.06	-0.440	0.0000	***	-0.5575836	-0.322416399
## 125.0.045 - 125.0.06	-0.145	0.0217	*	-0.2625836	-0.027416399
## 125.0.045 - 200.0.015	-0.185	0.0067	**	-0.3025836	-0.067416399
## 125.0.045 - 200.0.03	-0.165	0.0120	*	-0.2825836	-0.047416399
## 125.0.045 - 200.0.045	-0.205	0.0038	**	-0.3225836	-0.087416399
## 125.0.045 - 200.0.06	-0.250	0.0012	**	-0.3675836	-0.132416399
## 125.0.06 - 200.0.015	-0.040	0.4554		-0.1575836	0.077583601
## 125.0.06 - 200.0.03	-0.020	0.7051		-0.1375836	0.097583601
## 125.0.06 - 200.0.045	-0.060	0.2731		-0.1775836	0.057583601
## 125.0.06 - 200.0.06	-0.105	0.0734	.	-0.2225836	0.012583601
## 200.0.015 - 200.0.03	0.020	0.7051		-0.0975836	0.137583601
## 200.0.015 - 200.0.045	-0.020	0.7051		-0.1375836	0.097583601
## 200.0.015 - 200.0.06	-0.065	0.2382		-0.1825836	0.052583601
## 200.0.03 - 200.0.045	-0.040	0.4554		-0.1575836	0.077583601
## 200.0.03 - 200.0.06	-0.085	0.1341		-0.2025836	0.032583601
## 200.0.045 - 200.0.06	-0.045	0.4032		-0.1625836	0.072583601

And by performing a LSD test on the treatments interaction, we can observe some significant difference between interactions by looking at the p-values result above. More importantly, we can see from the result that when the drill speed is 200 and feed rate is 0.06, the **thrust force is maximized with an average value of 2.91**. This concludes that we shall operate the process under this condition.

Question 5.24

We can construct a effects model same as the previous question. For cycle time effect:

$$H_0 : \tau_1 = \tau_2 = \tau_3$$

H_a : at least one τ_i is different than the other, where $i \in \{1, 2, 3\}$

Similarly, we can set our hypotheses for the other two treatment effects, two-way interaction between each of the two treatments and three-way interaction.

```
#Three-factor factorial design
cloth_data <- data.frame(Cycle.Time=factor(c(rep(40,18),rep(50,18),rep(60,18))),
                        Temperature=factor(rep(c(300,350),each=9,times=3)),
                        Operator=factor(rep(c(1,2,3),each=3,times=6)),
                        Cloth=c(23,24,25,27,28,26,31,32,29,24,23,28,38,36,35,
                                   34,36,39,36,35,36,34,38,39,33,34,35,37,39,35,
                                   34,38,36,34,36,31,28,24,27,35,35,34,26,27,25,
                                   26,29,25,36,37,34,28,26,24))

anova(lm(Cloth~Cycle.Time*Temperature*Operator, data=cloth_data))

## Analysis of Variance Table
##
## Response: Cloth
##
##              Df Sum Sq Mean Sq F value    Pr(>F)
## Cycle.Time      2  436.00  218.000  66.5085 8.141e-13 ***
## Temperature      1   50.07   50.074  15.2768 0.0003934 ***
## Operator         2  261.33  130.667  39.8644 7.439e-10 ***
## Cycle.Time:Temperature  2   78.81   39.407  12.0226 0.0001002 ***
## Cycle.Time:Operator    4  355.67   88.917  27.1271 1.982e-10 ***
## Temperature:Operator   2   11.26    5.630   1.7175 0.1938948
## Cycle.Time:Temperature:Operator  4   46.19   11.546   3.5226 0.0158701 *
## Residuals          36  118.00    3.278
## ---
```

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

Based on the result, we can see that only the interaction between temperature and operator is not significant since we have a p-value of $0.19 > 0.05$. Other than that, every other terms are significant. Therefore, we can perform LSD tests on three treatments.

```
# Fisher LSD: Compare the cycle time regardless of temperature, operator.
```

```
LSD.test(cloth_data$Cloth, cloth_data$Cycle.Time, DError=36,  
         MSerror=3.278, console=T)
```

```
##
```

```
## Study: cloth_data$Cloth ~ cloth_data$Cycle.Time
```

```
##
```

```
## LSD t Test for cloth_data$Cloth
```

```
##
```

```
## Mean Square Error:  3.278
```

```
##
```

```
## cloth_data$Cycle.Time,  means and individual ( 95 %) CI
```

```
##
```

	cloth_data.Cloth	std	r	LCL	UCL	Min	Max
## 40	29.88889	5.378485	18	29.02341	30.75437	23	39
## 50	35.55556	2.120550	18	34.69008	36.42103	31	39
## 60	29.22222	4.557548	18	28.35674	30.08770	24	37

```
##
```

```
## Alpha: 0.05 ; DF Error: 36
```

```
## Critical Value of t: 2.028094
```

```
##
```

```
## least Significant Difference: 1.223971
```

```
##
```

```
## Treatments with the same letter are not significantly different.
```

```
##
```

```
##      cloth_data$Cloth groups
## 50      35.55556      a
## 40      29.88889      b
## 60      29.22222      b

# Fisher LSD: Compare the temperature regardless of cycle time, operator.
LSD.test(cloth_data$Cloth, cloth_data$Temperature, DError=36,
          MSerror=3.278, console=T)

##
## Study: cloth_data$Cloth ~ cloth_data$Temperature
##
## LSD t Test for cloth_data$Cloth
##
## Mean Square Error:  3.278
##
## cloth_data$Temperature, means and individual ( 95 %) CI
##
##      cloth_data.Cloth      std  r      LCL      UCL Min Max
## 300      30.59259 4.837826 27 29.88593 31.29925 23 39
## 350      32.51852 5.184076 27 31.81186 33.22518 23 39
##
## Alpha: 0.05 ; DF Error: 36
## Critical Value of t: 2.028094
##
## least Significant Difference: 0.9993685
##
## Treatments with the same letter are not significantly different.
##
##      cloth_data$Cloth groups
## 350      32.51852      a
```

```
## 300          30.59259      b

# Fisher LSD: Compare the operator regardless of cycle time, temperature.
LSD.test(cloth_data$Cloth, cloth_data$Operator, DError=36,
          MSerror=3.278, console=T)

##
## Study: cloth_data$Cloth ~ cloth_data$Operator
##
## LSD t Test for cloth_data$Cloth
##
## Mean Square Error:  3.278
##
## cloth_data$Operator,  means and individual ( 95 %) CI
##
##   cloth_data.Cloth      std  r      LCL      UCL Min Max
## 1          29.11111 5.571871 18 28.24563 29.97659  23  39
## 2          34.44444 3.776432 18 33.57897 35.30992  26  39
## 3          31.11111 4.377602 18 30.24563 31.97659  24  39
##
## Alpha: 0.05 ; DF Error: 36
## Critical Value of t: 2.028094
##
## least Significant Difference: 1.223971
##
## Treatments with the same letter are not significantly different.
##
##   cloth_data$Cloth groups
## 2          34.44444      a
## 3          31.11111      b
## 1          29.11111      c
```


From the LSD test results, we can first see that cycle time level of 50 is significant different than levels of 40 and 60. Then, two temperature levels are significant different than each other. Moreover, three operator levels are also significant different than each other.

Then, we can also perform LSD tests on the two significant interactions.

```
cloth_data$CycleTemp <- with(cloth_data,
                             interaction(Cycle.Time, Temperature))
CycleTemp_fit <- aov(Cloth~CycleTemp, data=cloth_data)
LSD.test(CycleTemp_fit, "CycleTemp", group=FALSE, console = TRUE)

##
## Study: CycleTemp_fit ~ "CycleTemp"
##
## LSD t Test for Cloth
##
## Mean Square Error: 16.50926
##
## CycleTemp, means and individual ( 95 %) CI
##
##          Cloth      std r      LCL      UCL Min Max
## 40.300 27.22222 3.073181 9 24.49905 29.94540 23 32
## 40.350 32.55556 6.002314 9 29.83238 35.27873 23 39
## 50.300 35.55556 1.943651 9 32.83238 38.27873 33 39
## 50.350 35.55556 2.403701 9 32.83238 38.27873 31 39
## 60.300 29.00000 4.415880 9 26.27682 31.72318 24 35
## 60.350 29.44444 4.952553 9 26.72127 32.16762 24 37
##
## Alpha: 0.05 ; DF Error: 48
## Critical Value of t: 2.010635
##
## Comparison between treatments means
```

##		difference	pvalue	signif.	LCL	UCL
##	40.300 - 40.350	-5.3333333	0.0076	**	-9.1844859	-1.4821808
##	40.300 - 50.300	-8.3333333	0.0001	***	-12.1844859	-4.4821808
##	40.300 - 50.350	-8.3333333	0.0001	***	-12.1844859	-4.4821808
##	40.300 - 60.300	-1.7777778	0.3580		-5.6289303	2.0733748
##	40.300 - 60.350	-2.2222222	0.2517		-6.0733748	1.6289303
##	40.350 - 50.300	-3.0000000	0.1239		-6.8511526	0.8511526
##	40.350 - 50.350	-3.0000000	0.1239		-6.8511526	0.8511526
##	40.350 - 60.300	3.5555556	0.0696	.	-0.2955970	7.4067081
##	40.350 - 60.350	3.1111111	0.1109		-0.7400414	6.9622637
##	50.300 - 50.350	0.0000000	1.0000		-3.8511526	3.8511526
##	50.300 - 60.300	6.5555556	0.0013	**	2.7044030	10.4067081
##	50.300 - 60.350	6.1111111	0.0025	**	2.2599586	9.9622637
##	50.350 - 60.300	6.5555556	0.0013	**	2.7044030	10.4067081
##	50.350 - 60.350	6.1111111	0.0025	**	2.2599586	9.9622637
##	60.300 - 60.350	-0.4444444	0.8175		-4.2955970	3.4067081

From this result, we can observe from the result that when the cycle time is 50 and temperatures of 300 and 350, the **cloth scores are maximized with both average values of 35.556**.

```
cloth_data$CycleOperator <- with(cloth_data,
                                interaction(Cycle.Time, Operator))
CycleOperator_fit <- aov(Cloth~CycleOperator, data=cloth_data)
LSD.test(CycleOperator_fit, "CycleOperator", group=FALSE, console = TRUE)

##
## Study: CycleOperator_fit ~ "CycleOperator"
##
## LSD t Test for Cloth
```

```

##
## Mean Square Error: 6.762963
##
## CycleOperator, means and individual ( 95 %) CI
##
##          Cloth          std r          LCL          UCL Min Max
## 40.1 24.50000 1.870829 6 22.36167 26.63833 23 28
## 40.2 31.66667 5.240865 6 29.52834 33.80500 26 38
## 40.3 33.50000 3.619392 6 31.36167 35.63833 29 39
## 50.1 36.33333 1.505545 6 34.19500 38.47166 35 39
## 50.2 36.50000 2.167948 6 34.36167 38.63833 34 39
## 50.3 33.83333 1.722401 6 31.69500 35.97166 31 36
## 60.1 26.50000 1.870829 6 24.36167 28.63833 24 29
## 60.2 35.16667 1.169045 6 33.02834 37.30500 34 37
## 60.3 26.00000 1.414214 6 23.86167 28.13833 24 28
##
## Alpha: 0.05 ; DF Error: 45
## Critical Value of t: 2.014103
##
## Comparison between treatments means
##
##          difference pvalue signif.          LCL          UCL
## 40.1 - 40.2 -7.1666667 0.0000 *** -10.1907213 -4.14261200
## 40.1 - 40.3 -9.0000000 0.0000 *** -12.0240547 -5.97594534
## 40.1 - 50.1 -11.8333333 0.0000 *** -14.8573880 -8.80927867
## 40.1 - 50.2 -12.0000000 0.0000 *** -15.0240547 -8.97594534
## 40.1 - 50.3 -9.3333333 0.0000 *** -12.3573880 -6.30927867
## 40.1 - 60.1 -2.0000000 0.1895 -5.0240547 1.02405466
## 40.1 - 60.2 -10.6666667 0.0000 *** -13.6907213 -7.64261200
## 40.1 - 60.3 -1.5000000 0.3231 -4.5240547 1.52405466

```

## 40.2 - 40.3	-1.8333333	0.2284		-4.8573880	1.19072133
## 40.2 - 50.1	-4.6666667	0.0033	**	-7.6907213	-1.64261200
## 40.2 - 50.2	-4.8333333	0.0024	**	-7.8573880	-1.80927867
## 40.2 - 50.3	-2.1666667	0.1559		-5.1907213	0.85738800
## 40.2 - 60.1	5.1666667	0.0013	**	2.1426120	8.19072133
## 40.2 - 60.2	-3.5000000	0.0243	*	-6.5240547	-0.47594534
## 40.2 - 60.3	5.6666667	0.0005	***	2.6426120	8.69072133
## 40.3 - 50.1	-2.8333333	0.0656	.	-5.8573880	0.19072133
## 40.3 - 50.2	-3.0000000	0.0518	.	-6.0240547	0.02405466
## 40.3 - 50.3	-0.3333333	0.8253		-3.3573880	2.69072133
## 40.3 - 60.1	7.0000000	0.0000	***	3.9759453	10.02405466
## 40.3 - 60.2	-1.6666667	0.2729		-4.6907213	1.35738800
## 40.3 - 60.3	7.5000000	0.0000	***	4.4759453	10.52405466
## 50.1 - 50.2	-0.1666667	0.9121		-3.1907213	2.85738800
## 50.1 - 50.3	2.5000000	0.1028		-0.5240547	5.52405466
## 50.1 - 60.1	9.8333333	0.0000	***	6.8092787	12.85738800
## 50.1 - 60.2	1.1666667	0.4412		-1.8573880	4.19072133
## 50.1 - 60.3	10.3333333	0.0000	***	7.3092787	13.35738800
## 50.2 - 50.3	2.6666667	0.0825	.	-0.3573880	5.69072133
## 50.2 - 60.1	10.0000000	0.0000	***	6.9759453	13.02405466
## 50.2 - 60.2	1.3333333	0.3792		-1.6907213	4.35738800
## 50.2 - 60.3	10.5000000	0.0000	***	7.4759453	13.52405466
## 50.3 - 60.1	7.3333333	0.0000	***	4.3092787	10.35738800
## 50.3 - 60.2	-1.3333333	0.3792		-4.3573880	1.69072133
## 50.3 - 60.3	7.8333333	0.0000	***	4.8092787	10.85738800
## 60.1 - 60.2	-8.6666667	0.0000	***	-11.6907213	-5.64261200
## 60.1 - 60.3	0.5000000	0.7407		-2.5240547	3.52405466
## 60.2 - 60.3	9.1666667	0.0000	***	6.1426120	12.19072133

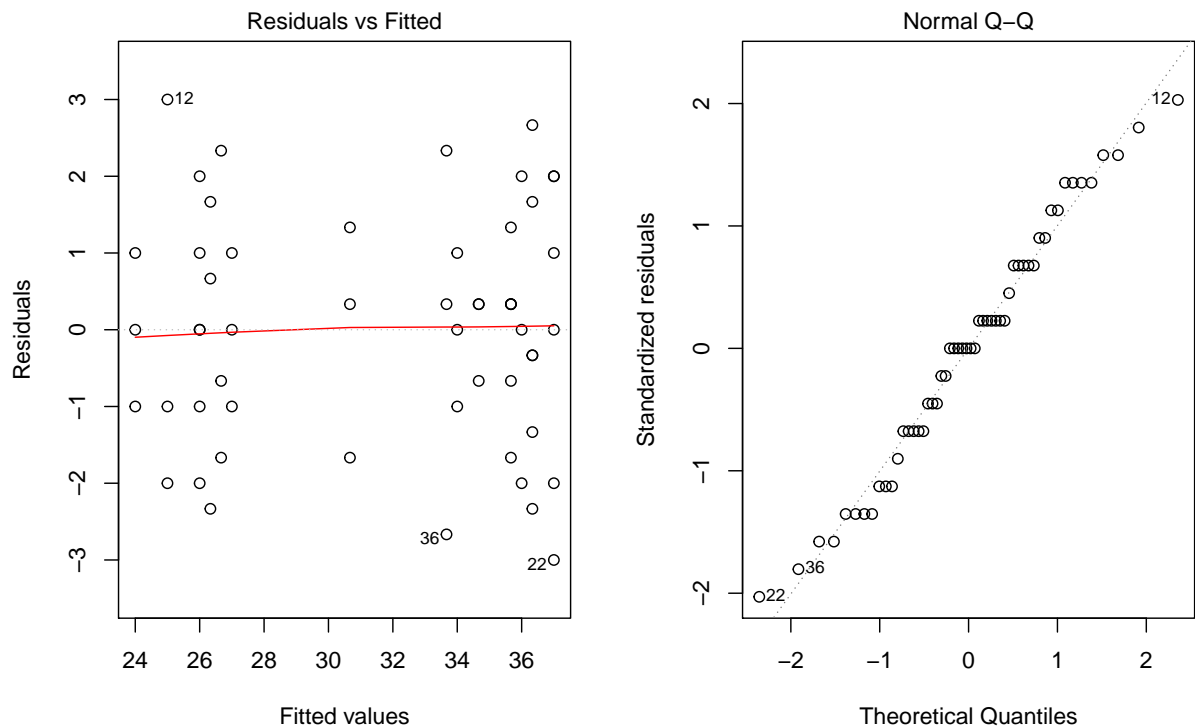
From this result, we can observe from the result that when the cycle time is 50 and

operator is number 2, the **cloth score is maximized with an average value of 36.5**.

Therefore, we shall consider operating the process by the conditions we concluded above.

Lastly, we need to check the model's adequacy(constant variance and normality assumptions).

```
fit <- aov(lm(Cloth~Cycle.Time*Temperature*Operator, data=cloth_data))
par(mfrow=c(1,2))
plot(fit, 1); plot(fit, 2)
```



The residuals vs fitted values plot and QQplot do not seem to have any unusual patterns. We can observe a constant band throughout the residuals vs fitted values plot and most of the points follow the theoretical line in the normal QQplot.

Question 5.46

(a)

$$SS_A = 118.667 - (96.333 + 12.167 + 10.000) = 0.167$$

(b)

$$DF_A = \frac{0.167}{0.0833} = 2$$

(c)

$$DF_B = \frac{96.333}{96.3333} = 1$$

(d)

$$MS_E = \frac{10.000}{6} = 1.667$$

(e)

```
pf(3.65, df1=2, df2=6, lower.tail=F)
```

```
## [1] 0.09181187
```

The p-value for the interaction test statistic is 0.09.

(f) We know the degree freedom for factor A is 2. Therefore, there are $2 + 1 = 3$ levels.

(g) Similarly, we know there are $1 + 1 = 2$ levels in factor B.

(h) We know the $DF_{error} = ab(n - 1)$, and we know $a = 3$, $b = 2$, and $DF_{error} = 6$. Therefore, $n = 2$ which suggests that there are 2 replicates in this experiment.

(i) No, since we don't find the interaction between factor B and factor A is significant where we obtained a p-value greater than 0.05.

(j)

$$\hat{\sigma} = \sqrt{\frac{10}{6}} = 1.29$$

Question 5.52

(a) True.

$$SS_{error} = 185 - (50 + 80 + 30 + 10) = 15$$

$$DF_{error} = 11 - (1 + 2 + 2 + 1) = 5$$

$$\hat{\sigma} = \sqrt{\frac{15}{5}} = 1.73$$

(b)

$$DF_{error} = 11 - (1 + 2 + 2) = 6$$

(c)

$$SS_{error} = 185 - (50 + 80 + 30) = 25$$

$$MS_{error} = \frac{25}{6} = 4.167$$

(d) False, since $F_0 = 3.6$ less than the critical F value. Not significant.

```
#F critical
```

```
qf(0.95, 2, 6)
```

```
## [1] 5.143253
```

$$F_0 = \frac{MS_{AB}}{MS_{error}} = \frac{15}{4.167} = 3.6$$