

STAT 522 — Assignment 5

Subasish Das (sxd1684)

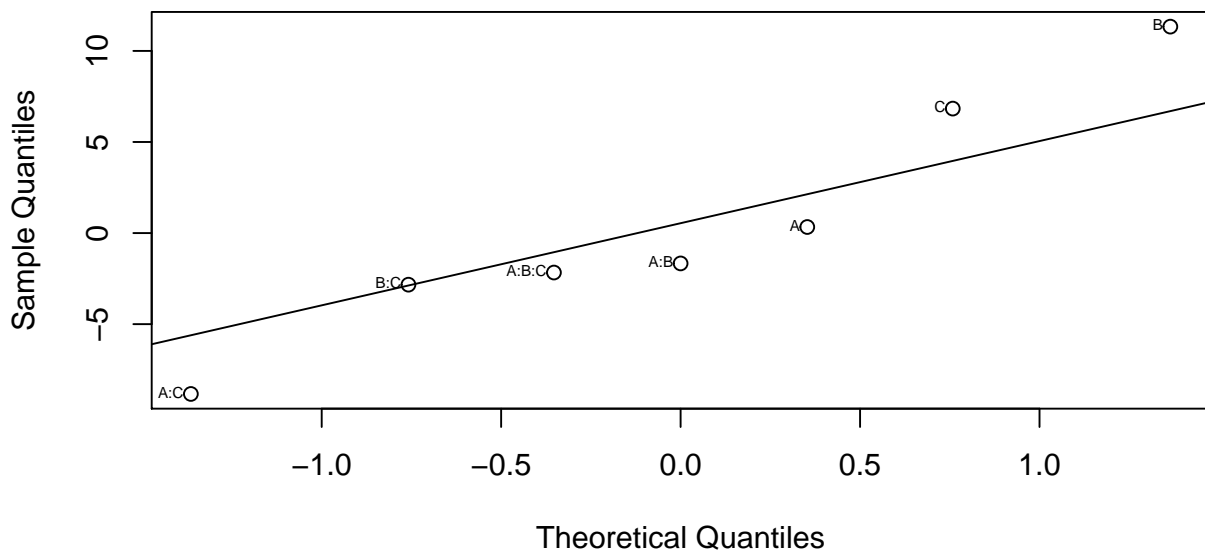
10th April 2014

1 Exercise 6.1

(a)

```
## Given
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5/effects")
tool <- read.csv("6.1a.csv")
mydata.lm = lm(Life.Hours ~ A * B * C, tool)
n = 3
k = 3
effects = coefficients(mydata.lm)[-c(1)] * 2
SS = effects^2 * n * 2^(k - 2)
percentage = SS/sum(SS) * 100
tem = qqnorm(effects)
qqline(effects)
text(tem$x, tem$y, names(effects), pos = 2, offset = 0.2, cex = 0.5)
```

Normal Q-Q Plot



```
cbind(effects, SS, percentage)
```

##	effects	SS	percentage
## A	0.3333	0.6667	0.04134
## B	11.3333	770.6667	47.78834
## C	6.8333	280.1667	17.37288
## A:B	-1.6667	16.6667	1.03348
## A:C	-8.8333	468.1667	29.03059
## B:C	-2.8333	48.1667	2.98677
## A:B:C	-2.1667	28.1667	1.74659

From the effect values and normality effect plot, it's observed that factors B, C and interaction AC are seemed to be significant.

(b)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
tool <- read.csv("6.1.csv")
hours <- tool$Life.Hours

tool[, 1:3] <- lapply(tool[, 1:3], factor)
tool1.aov <- aov(Life.Hours ~ A.Cutting.Speed * B.Tool.Geometry * C.Cutting.Angle,
  data = tool)
summary(tool1.aov)
```

	Df	Sum Sq	Mean Sq	F value
## A.Cutting.Speed	1	1	1	0.02
## B.Tool.Geometry	1	771	771	25.55
## C.Cutting.Angle	1	280	280	9.29
## A.Cutting.Speed:B.Tool.Geometry	1	17	17	0.55
## A.Cutting.Speed:C.Cutting.Angle	1	468	468	15.52
## B.Tool.Geometry:C.Cutting.Angle	1	48	48	1.60
## A.Cutting.Speed:B.Tool.Geometry:C.Cutting.Angle	1	28	28	0.93
## Residuals	16	483	30	

```
## Pr(>F)
## A.Cutting.Speed 0.88368
## B.Tool.Geometry 0.00012 ***
## C.Cutting.Angle 0.00768 **
## A.Cutting.Speed:B.Tool.Geometry 0.46808
## A.Cutting.Speed:C.Cutting.Angle 0.00117 **
## B.Tool.Geometry:C.Cutting.Angle 0.22448
## A.Cutting.Speed:B.Tool.Geometry:C.Cutting.Angle 0.34828
## Residuals
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The ANOVA table confirms the significance of factors B, C and interaction AC.

The ANOVA of the reduced model is performed below:

```
tool2.aov <- aov(Life.Hours ~ A.Cutting.Speed + B.Tool.Geometry + C.Cutting.Angle +
  A.Cutting.Speed * C.Cutting.Angle, data = tool)
summary(tool2.aov)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
## A.Cutting.Speed	1	1	1	0.02	0.8836
## B.Tool.Geometry	1	771	771	25.44	7.2e-05 ***
## C.Cutting.Angle	1	280	280	9.25	0.0067 **
## A.Cutting.Speed:C.Cutting.Angle	1	468	468	15.45	0.0009 ***
## Residuals	19	576	30		

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

Factor A is kept here to maintain the hierarchy. The factors B, C and interaction AC are significant at 0.01 level.

(c)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
tool_num <- read.csv("6.1.csv")
tool_num.lm <- lm(Life.Hours ~ A.Cutting.Speed + B.Tool.Geometry + C.Cutting.Angle +
  A.Cutting.Speed * C.Cutting.Angle, data = tool_num)
summary(tool_num.lm)
```

```
##
## Call:
## lm(formula = Life.Hours ~ A.Cutting.Speed + B.Tool.Geometry +
##     C.Cutting.Angle + A.Cutting.Speed * C.Cutting.Angle, data = tool_num)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7.333 -4.375 -0.417  2.958 11.500
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      40.833      1.124   36.34 < 2e-16 ***
## A.Cutting.Speed    0.167      1.124    0.15  0.8836
## B.Tool.Geometry    5.667      1.124    5.04  7.2e-05 ***
## C.Cutting.Angle    3.417      1.124    3.04  0.0067 **
## A.Cutting.Speed:C.Cutting.Angle -4.417      1.124   -3.93  0.0009 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.5 on 19 degrees of freedom
## Multiple R-squared:  0.725, Adjusted R-squared:  0.667
## F-statistic: 12.5 on 4 and 19 DF, p-value: 3.69e-05
```

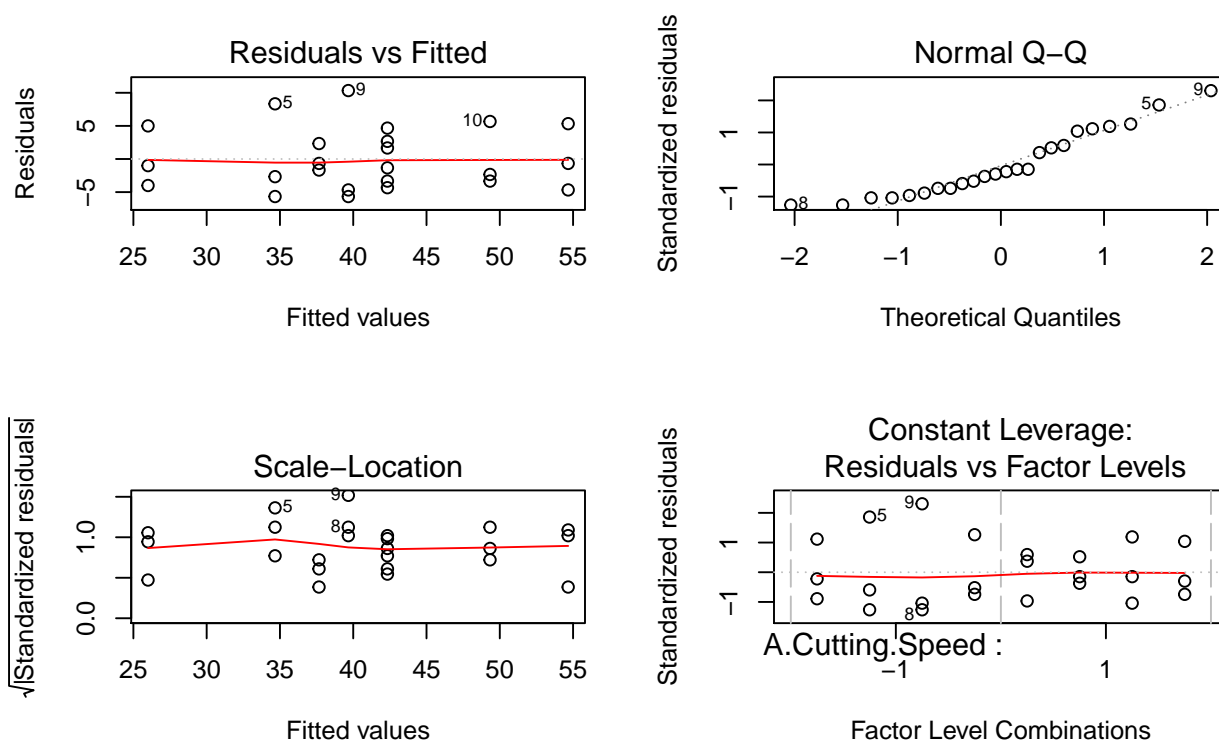
The regression model:

$$y_{ijk} = 40.8333 + 0.1667x_A + 5.667x_B + 3.417x_C - 4.4167x_Ax_C$$

The regression model is based on the significant factors B (tool geometry), C (cutting angle) and interaction of AC.

(d)

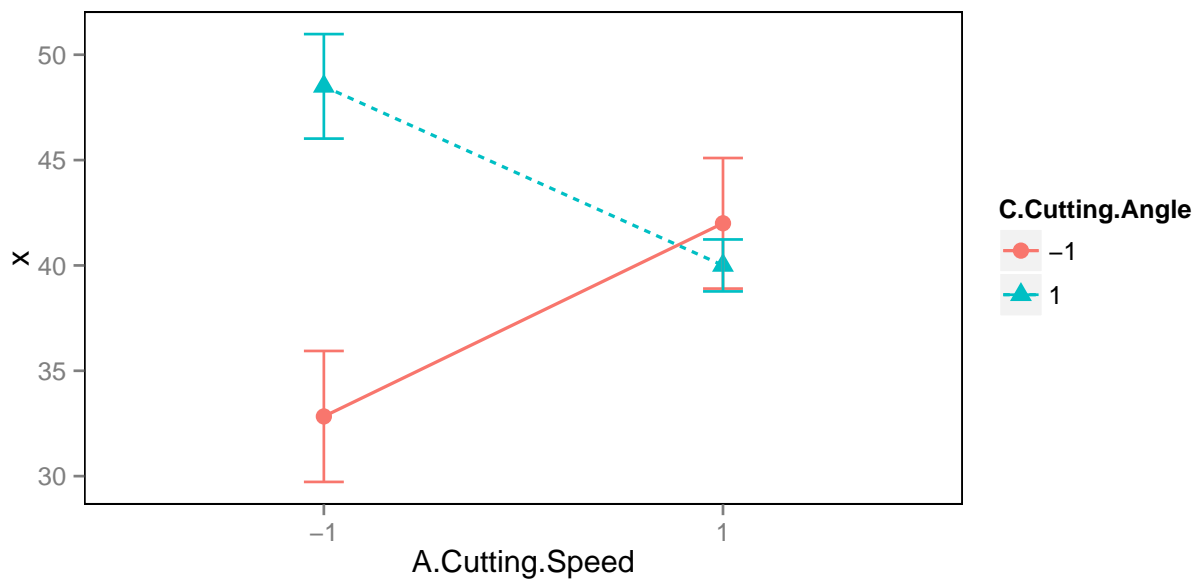
```
par(mfrow = c(2, 2))
plot(tool1.aov)
```



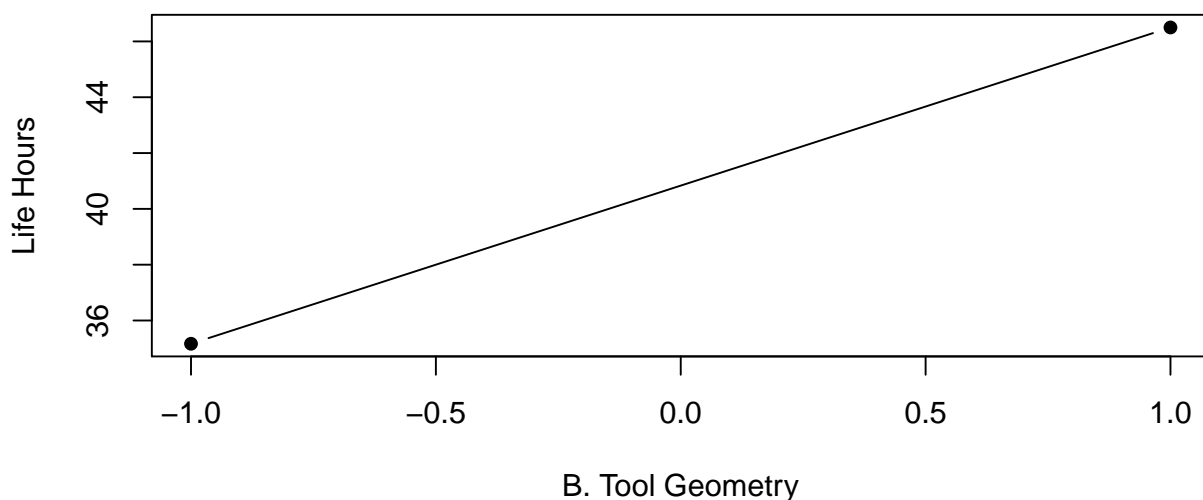
Nothing unusual is visible regarding the residual plots.

(e)

```
## Interaction Plot
tool[, 1:3] <- lapply(tool[, 1:3], factor)
library(ggplot2)
df <- with(tool, aggregate(Life.Hours, list(C.Cutting.Angle = C.Cutting.Angle,
  A.Cutting.Speed = A.Cutting.Speed), mean))
df$se <- with(tool, aggregate(Life.Hours, list(C.Cutting.Angle = C.Cutting.Angle,
  A.Cutting.Speed = A.Cutting.Speed), function(x) sd(x)/sqrt(10)))[, 3]
opar <- theme_update(panel.grid.major = element_blank(), panel.grid.minor = element_blank(),
  panel.background = element_rect(colour = "black"))
gp <- ggplot(df, aes(x = A.Cutting.Speed, y = x, colour = C.Cutting.Angle, group = C.Cutting.Angle))
gp + geom_line(aes(linetype = C.Cutting.Angle), size = 0.6) + geom_point(aes(shape = C.Cutting.Angle),
  size = 3) + geom_errorbar(aes(ymax = x + se, ymin = x - se), width = 0.1)
```



```
## One Factor Plot
with(tool, plot(c(-1, 1), tapply(Life.Hours, B.Tool.Geometry, mean), type = "b",
  pch = 16, xlab = "B. Tool Geometry", ylab = "Life Hours"))
```

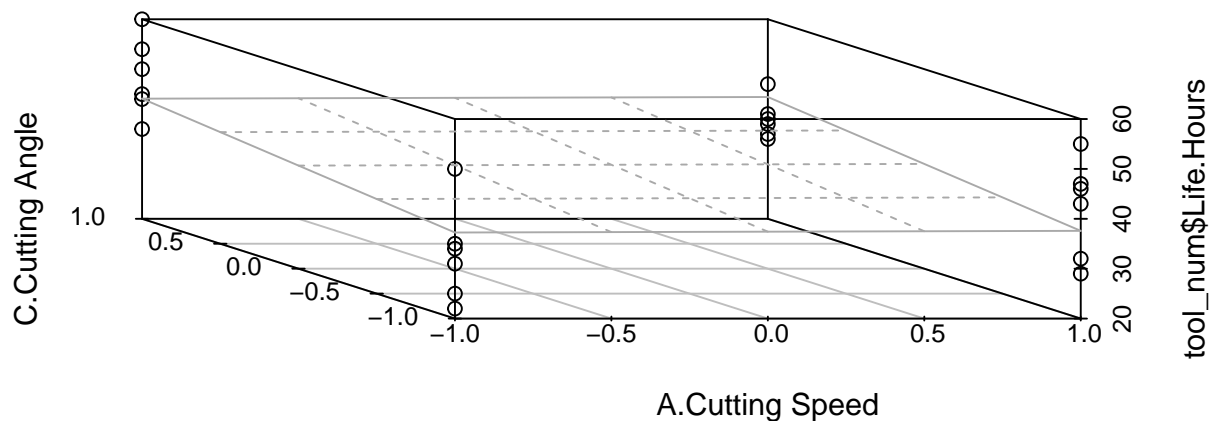


As B has positive effect, we can set B at a high level to increase the life hours. The interaction plot of AC indicates that life hours will be maximum at higher level of C and lower level of A.

2 Exercise 6.2

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
tool_num <- read.csv("6.1.csv")
tool_num.lm <- lm(Life.Hours ~ A.Cutting.Speed + C.Cutting.Angle, data = tool_num)

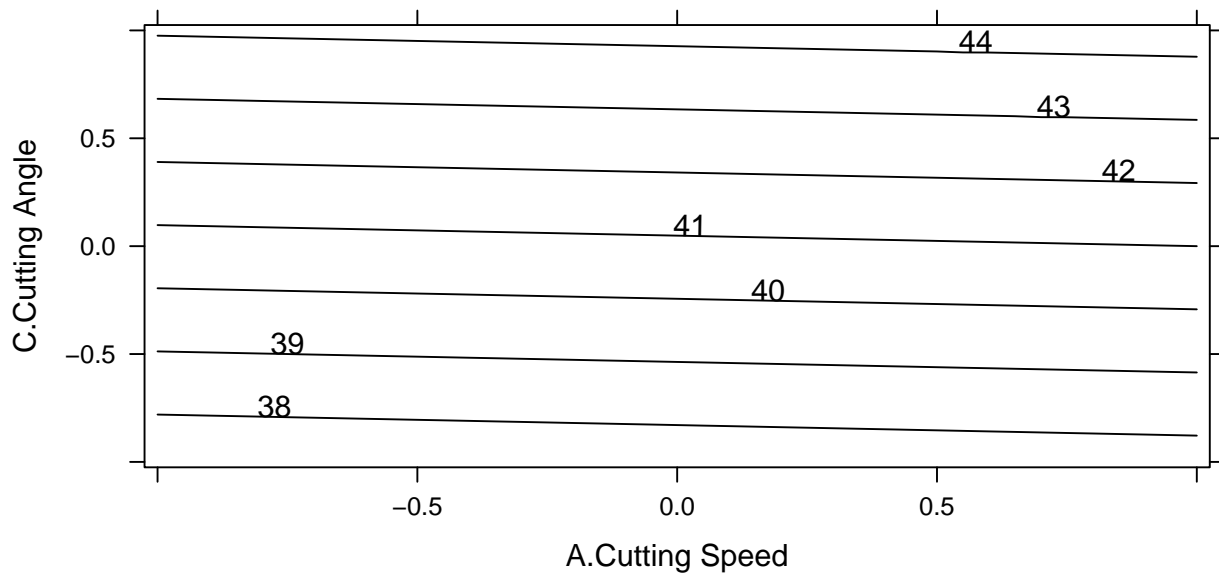
library(scatterplot3d)
s3d <- scatterplot3d(tool_num$A.Cutting.Speed, tool_num$C.Cutting.Angle, tool_num$Life.Hours,
  type = "p", angle = 135, scale.y = 1, xlab = "A.Cutting Speed", ylab = "C.Cutting Angle")
s3d$plane3d(tool_num.lm, lty.box = "solid", col = "darkgray")
```



```
tmp <- list(A.Cutting.Speed = seq(-1, 1, by = 0.05), C.Cutting.Angle = seq(-1,
  1, by = 0.05))
new.data <- expand.grid(tmp)
new.data$fit <- predict(tool_num.lm, new.data)
library(lme4)

## Loading required package: lattice
## Loading required package: Matrix
##
## Attaching package: 'lme4'
## The following object is masked from 'package:ggplot2':
##
## fortify

contourplot(fit ~ A.Cutting.Speed * C.Cutting.Angle, new.data, xlab = "A.Cutting Speed",
  ylab = "C.Cutting Angle")
```



The response surface plot and the contour plot are generated by using the regression model in problem 6.1 part (c). The curvature is visible due to the interaction of AC. Yes, these plots provide insight regarding the desirable operating conditions for this process.

3 Exercise 6.3

```
## Standard Error
n = 3
k = 3
S2 = 30.17
SE_effect = sqrt(S2/(n * 2^(k - 2)))
SE_effect

## [1] 2.242

## 95% Confidence Limit
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
tool <- read.csv("6.1.csv")
tool[, 1:3] <- lapply(tool[, 1:3], factor)
aov <- aov(Life.Hours ~ A.Cutting.Speed * B.Tool.Geometry * C.Cutting.Angle,
  data = tool)
confint(aov)
```

	2.5 %	97.5 %
## (Intercept)	19.2777	32.7223
## A.Cutting.Speed1	-0.8401	18.1735
## B.Tool.Geometry1	4.1599	23.1735
## C.Cutting.Angle1	6.8265	25.8401
## A.Cutting.Speed1:B.Tool.Geometry1	-12.4446	14.4446
## A.Cutting.Speed1:C.Cutting.Angle1	-26.7780	0.1113
## B.Tool.Geometry1:C.Cutting.Angle1	-14.7780	12.1113
## A.Cutting.Speed1:B.Tool.Geometry1:C.Cutting.Angle1	-27.6803	10.3469

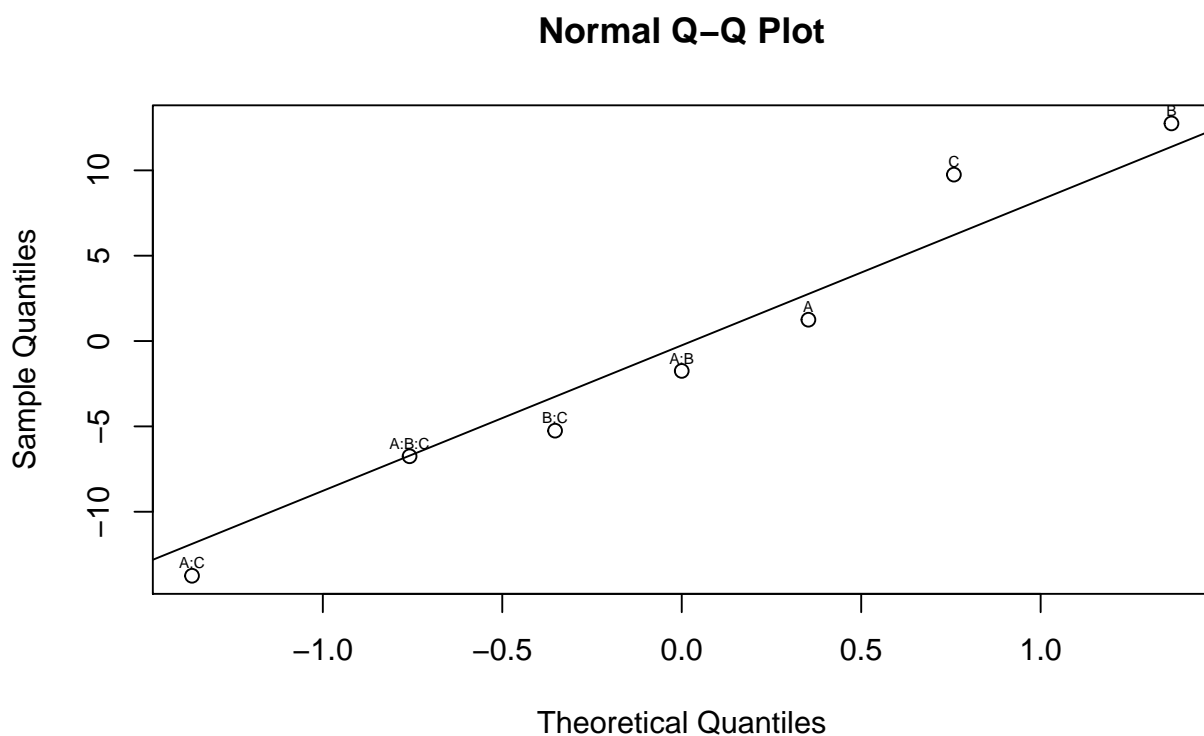
The standard error is 2.24.

The 95 percent confidence interval of the factors B and C and the interaction AC don't contain zero. This completely agrees with the variance approach.

4 Exercise 6.6

(a)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5/effects")
tool <- read.csv("6.6a.csv")
mydata.lm = lm(Life.Hours ~ A * B * C, tool)
n = 1
k = 3
effects = coefficients(mydata.lm)[-c(1)] * 2
SS = effects^2 * n * 2^(k - 2)
percentage = SS/sum(SS) * 100
tem = qqnorm(effects)
qqline(effects)
text(tem$x, tem$y, names(effects), pos = 3, offset = 0.2, cex = 0.5)
```



```
cbind(effects, SS, percentage)
```

##	effects	SS	percentage
## A	1.25	3.125	0.2979
## B	12.75	325.125	30.9975
## C	9.75	190.125	18.1266
## A:B	-1.75	6.125	0.5840
## A:C	-13.75	378.125	36.0505
## B:C	-5.25	55.125	5.2556
## A:B:C	-6.75	91.125	8.6879

From the normality plot for effects, factors B, C and interaction AC have larger effects.

(b)

```
## ANOVA including a check for pure quadratic curvature.
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
toola <- read.csv("6.6.csv")
```

```

tool1a.aov <- aov(Life.Hours ~ A.Cutting.Speed * B.Tool.Geometry * C.Cutting.Angle +
  I((A.Cutting.Speed)^2), data = toola)
summary(tool1a.aov)

##                               Df Sum Sq Mean Sq F value
## A.Cutting.Speed              1     3      3      0.20
## B.Tool.Geometry              1    325     325     21.20
## C.Cutting.Angle              1    190     190     12.40
## I((A.Cutting.Speed)^2)       1     0      0      0.00
## A.Cutting.Speed:B.Tool.Geometry 1     6      6      0.40
## A.Cutting.Speed:C.Cutting.Angle 1    378     378     24.66
## B.Tool.Geometry:C.Cutting.Angle 1     55      55      3.60
## A.Cutting.Speed:B.Tool.Geometry:C.Cutting.Angle 1     91      91      5.94
## Residuals                    3     46      15
##
##                               Pr(>F)
## A.Cutting.Speed              0.682
## B.Tool.Geometry              0.019 *
## C.Cutting.Angle              0.039 *
## I((A.Cutting.Speed)^2)       0.962
## A.Cutting.Speed:B.Tool.Geometry 0.572
## A.Cutting.Speed:C.Cutting.Angle 0.016 *
## B.Tool.Geometry:C.Cutting.Angle 0.154
## A.Cutting.Speed:B.Tool.Geometry:C.Cutting.Angle 0.093 .
## Residuals
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Inclusion of a pure quadratic curvature shows no significance. The ANOVA table confirms that the effects of the factors B, C and interaction AC are larger.

The ANOVA of the reduced model is performed below:

```

tool2a.aov <- aov(Life.Hours ~ A.Cutting.Speed + B.Tool.Geometry + C.Cutting.Angle +
  A.Cutting.Speed * C.Cutting.Angle, data = toola)
summary(tool2a.aov)

##                               Df Sum Sq Mean Sq F value Pr(>F)
## A.Cutting.Speed              1     3      3      0.11 0.7496
## B.Tool.Geometry              1    325     325     11.47 0.0117 *
## C.Cutting.Angle              1    190     190      6.71 0.0360 *
## A.Cutting.Speed:C.Cutting.Angle 1    378     378     13.34 0.0082 **
## Residuals                    7    198      28
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Factor A is kept here to maintain the hierarchy. The factors B, C and interaction AC are significant at 0.05 level.

(c)

```

setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
tool_numa <- read.csv("6.6.csv")
tool_numa.aov <- lm(Life.Hours ~ A.Cutting.Speed + B.Tool.Geometry + C.Cutting.Angle +
  A.Cutting.Speed * C.Cutting.Angle, data = tool_numa)
summary(tool_numa.aov)

##
## Call:
## lm(formula = Life.Hours ~ A.Cutting.Speed + B.Tool.Geometry +
##     C.Cutting.Angle + A.Cutting.Speed * C.Cutting.Angle, data = tool_numa)
##
## Residuals:

```



```
##      Min      1Q  Median      3Q      Max
## -6.917 -2.479 -0.042  2.583  6.833
##
## Coefficients:
##
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      40.917      1.537   26.62  2.7e-08 ***
## A.Cutting.Speed      0.625      1.882    0.33  0.7496
## B.Tool.Geometry      6.375      1.882    3.39  0.0117 *
## C.Cutting.Angle      4.875      1.882    2.59  0.0360 *
## A.Cutting.Speed:C.Cutting.Angle  -6.875      1.882   -3.65  0.0082 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.32 on 7 degrees of freedom
## Multiple R-squared:  0.819, Adjusted R-squared:  0.715
## F-statistic: 7.91 on 4 and 7 DF, p-value: 0.00979
```

The regression model:

$$y_{ijk} = 40.917 + 0.625x_A + 6.375x_B + 4.875x_C - 6.875x_Ax_C$$

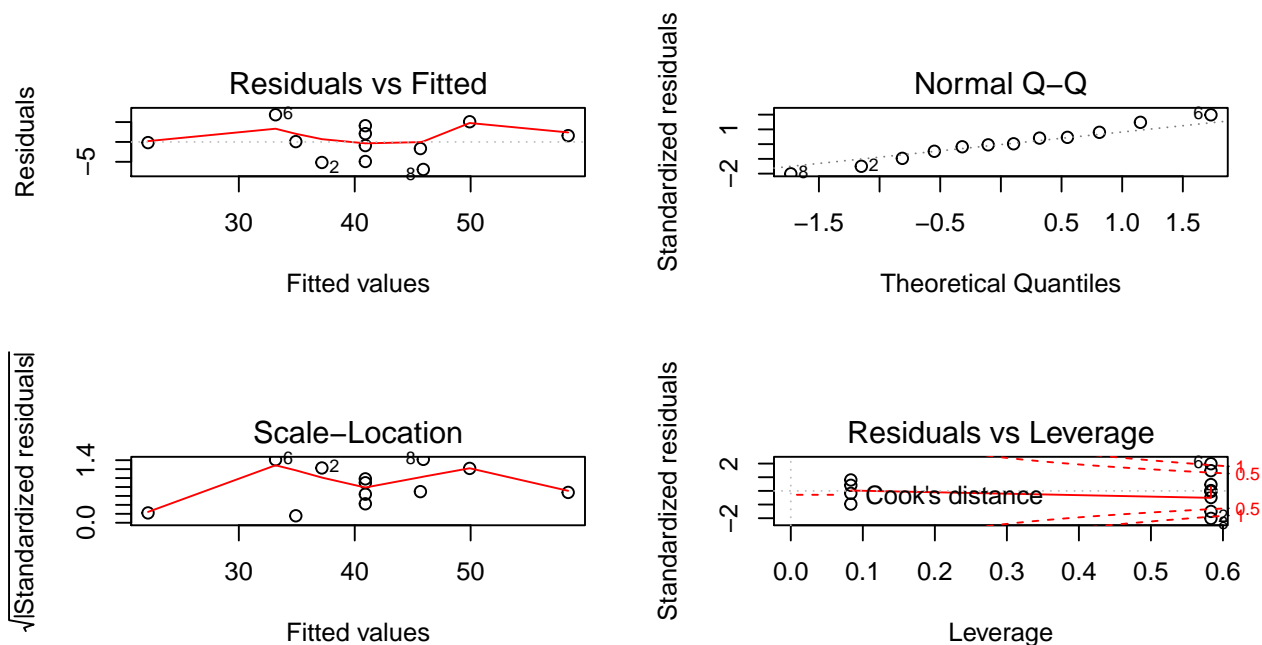
The regression model from 6.1 part (c):

$$y_{ijk} = 40.8333 + 0.1667x_A + 5.667x_B + 3.417x_C - 4.4167x_Ax_C$$

The regression model is based on the significant factors B,C and interaction of AC. The current model is not substantially different from the regression model of problem 6.1 part (c)

(d)

```
par(mfrow = c(2, 2))
plot(tool2a.aov)
```

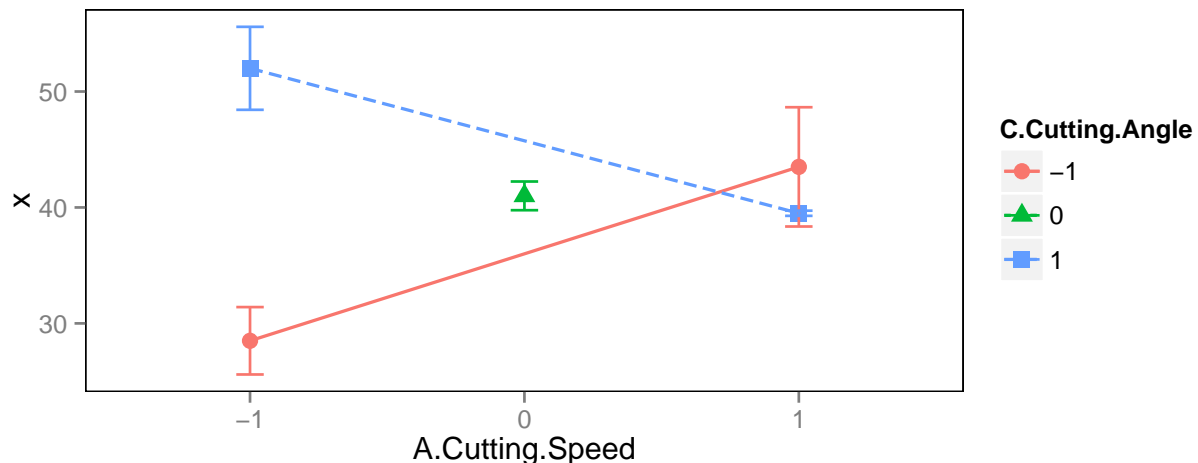


Nothing unusual is drastically visible regarding the residual plots.

(e)

```
## Interaction Graph
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
toola <- read.csv("6.6.csv")
toola[, 1:3] <- lapply(toola[, 1:3], factor)
library(ggplot2)
```

```
df <- with(toola, aggregate(Life.Hours, list(C.Cutting.Angle = C.Cutting.Angle,
  A.Cutting.Speed = A.Cutting.Speed), mean))
df$se <- with(toola, aggregate(Life.Hours, list(C.Cutting.Angle = C.Cutting.Angle,
  A.Cutting.Speed = A.Cutting.Speed), function(x) sd(x)/sqrt(10)))[, 3]
opar <- theme_update(panel.grid.major = element_blank(), panel.grid.minor = element_blank(),
  panel.background = element_rect(colour = "black"))
gp <- ggplot(df, aes(x = A.Cutting.Speed, y = x, colour = C.Cutting.Angle, group = C.Cutting.Angle))
gp + geom_line(aes(linetype = C.Cutting.Angle), size = 0.6) + geom_point(aes(shape = C.Cutting.Angle),
  size = 3) + geom_errorbar(aes(ymax = x + se, ymin = x - se), width = 0.1)
```



As B has positive effect, we can set at a high level to increase the life hours. The interaction plot of AC life hours will be maximum at higher level of C and lower level of A.

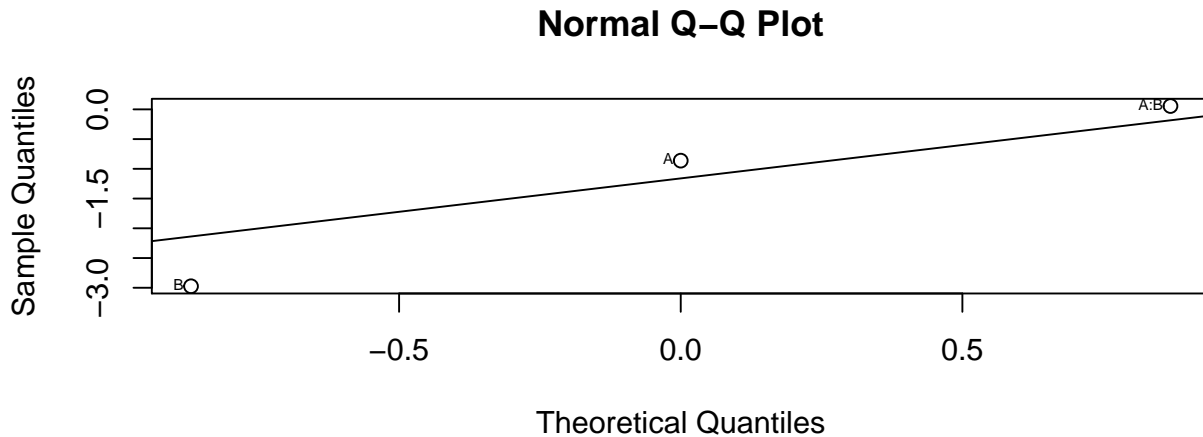
5 Exercise 6.12

(a)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5/effects")
circuit <- read.csv("6.12a.csv")
mydata.lm = lm(Thickness ~ A * B, circuit)
n = 2
k = 4
effects = coefficients(mydata.lm)[-c(1)] * 2
SS = effects^2 * n * 2^(k - 2)
percentage = SS/sum(SS) * 100
cbind(effects, SS, percentage)

##      effects      SS percentage
## A   -0.8624   5.94953    7.75025
## B   -2.9747  70.79072   92.21672
## A:B   0.0563   0.02536    0.03303

tem = qqnorm(effects)
qqline(effects)
text(tem$x, tem$y, names(effects), pos = 2, offset = 0.2, cex = 0.5)
```



From the factor effect values and the normality effect plot, we find that factor B (Deposition time) has significant effect.

(b)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
circuit <- read.csv("6.12.csv")
circuit.aov <- aov(Thickness ~ A.Flow.Rate * B.Dep.Time, data = circuit)
summary(circuit.aov)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
## A.Flow.Rate	1	0.40	0.403	1.26	0.28	
## B.Dep.Time	1	1.37	1.374	4.31	0.06 .	
## A.Flow.Rate:B.Dep.Time	1	0.32	0.317	0.99	0.34	
## Residuals	12	3.83	0.319			
## ---						
## Signif. codes:	0 '***'	0.001 '**'	0.01 '*'	0.05 '.'	0.1 ' '	1

From the values of the ANOVA table, we find that factor B (Deposition time) has significant effect at 0.1 level.

(c)

```
circuit.lm <- lm(Thickness ~ A.Flow.Rate * B.Dep.Time, data = circuit)
summary(circuit.lm)
```

```
##
## Call:
## lm(formula = Thickness ~ A.Flow.Rate * B.Dep.Time, data = circuit)
##
## Residuals:
```

	Min	1Q	Median	3Q	Max
##	-0.6133	-0.1443	-0.0056	0.1019	1.6447

```
##
## Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)		
## (Intercept)	37.6266	20.5334	1.83	0.092 .		
## A.Flow.Rate	-0.4312	0.3600	-1.20	0.254		
## B.Dep.Time	-1.4874	1.6108	-0.92	0.374		
## A.Flow.Rate:B.Dep.Time	0.0282	0.0282	1.00	0.339		
## ---						
## Signif. codes:	0 '***'	0.001 '**'	0.01 '*'	0.05 '.'	0.1 ' '	1
##						
## Residual standard error:	0.565	on 12 degrees of freedom				
## Multiple R-squared:	0.353	Adjusted R-squared:	0.192			
## F-statistic:	2.19	on 3 and 12 DF,	p-value:	0.142		

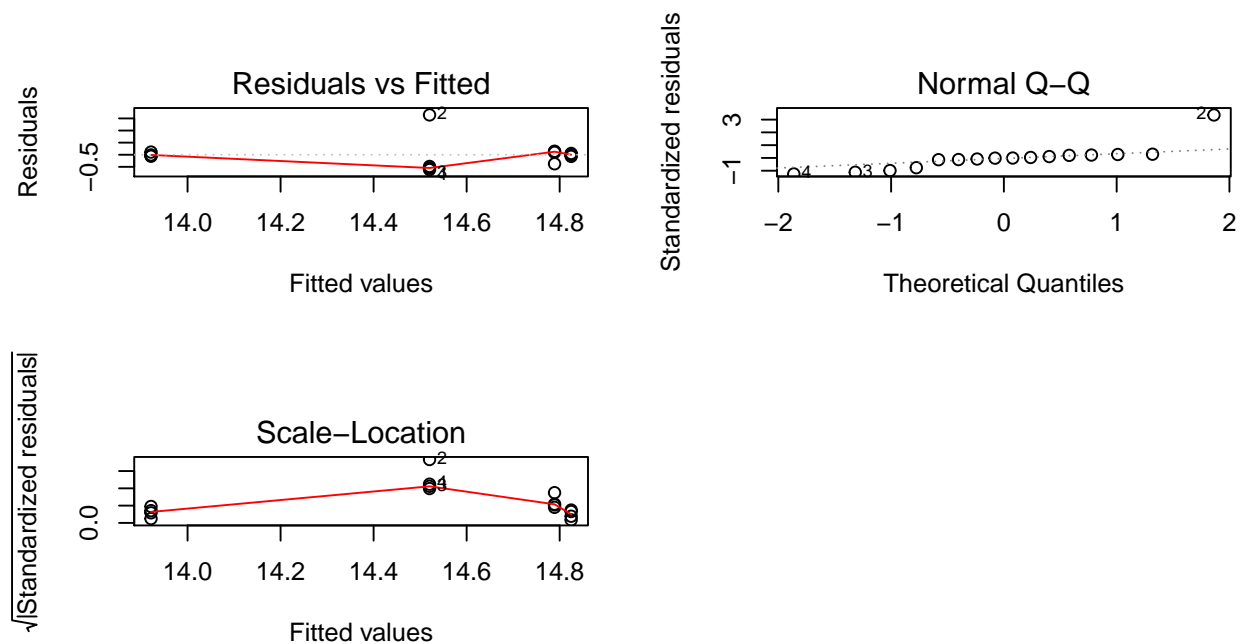
The regression equation:

$$y = 37.626 - 0.432x_A - 1.487x_B + 0.028x_Ax_B$$

(d)

```
par(mfrow = c(2, 2))
plot(circuit.aov)

## hat values (leverages) are all = 0.25
## and there are no factor predictors; no plot no. 5
```



From the residual plots, it's found that observation no. 2 (16.165) falls outside the groupings in the normal probability plot. This observation is also visible outside the groupings in residual versus predicted plot.

(e)

We can deal with the potential outlier by replacing that observation with the average of the observations from that particular cell. Another way is to validate that particular point. If validation produces findings out any error, it should be corrected.

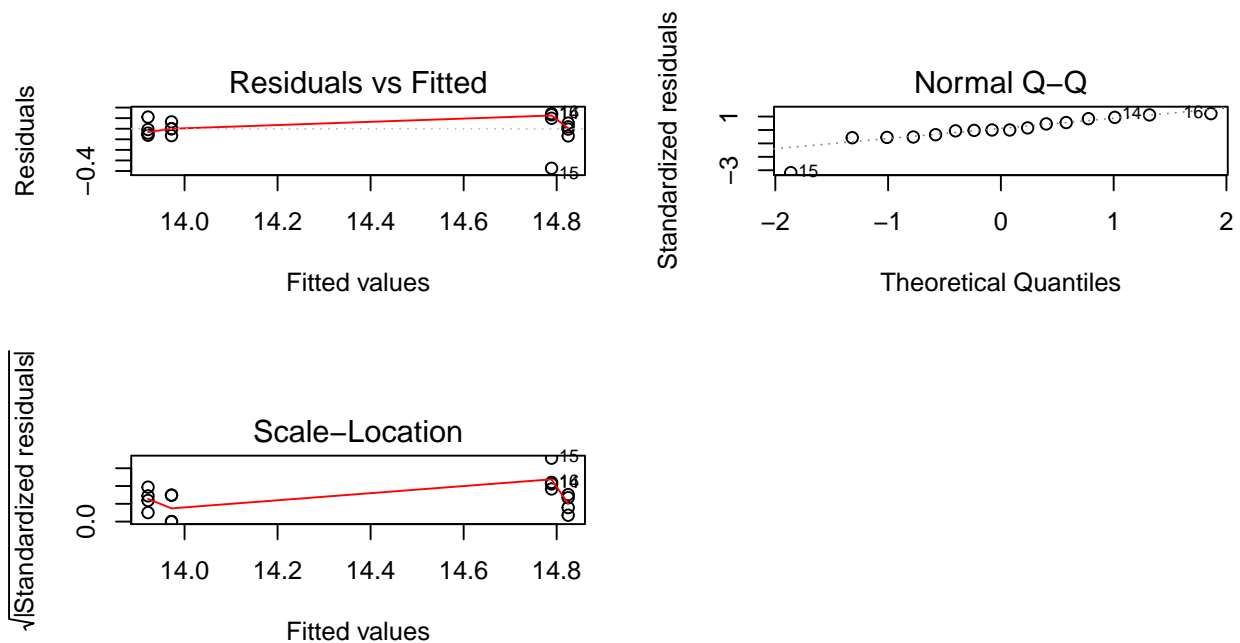
Observation 2 is replaced by the average of the remaining three other runs in the cell which is 13.972.

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
circuit_1 <- read.csv("6.12_1.csv")
circuit_1.aov <- aov(Thickness ~ A.Flow.Rate * B.Dep.Time, data = circuit_1)
summary(circuit_1.aov)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## A.Flow.Rate    1  0.007    0.007      0.40     0.54
## B.Dep.Time      1  2.959    2.959   160.29 2.7e-08 ***
## A.Flow.Rate:B.Dep.Time 1  0.000    0.000      0.01     0.92
## Residuals     12  0.222    0.018
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
par(mfrow = c(2, 2))
plot(circuit_1.aov)

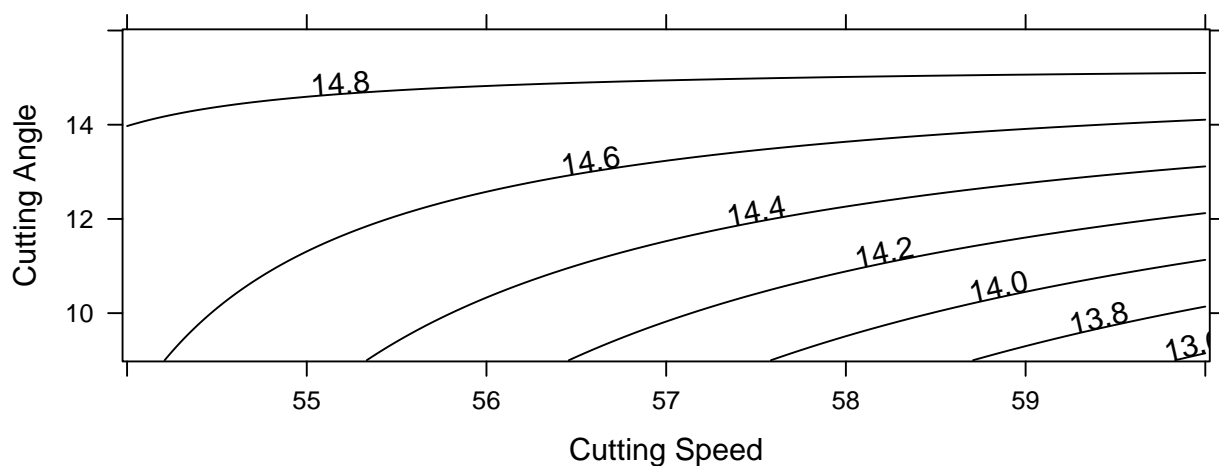
## hat values (leverages) are all = 0.25
## and there are no factor predictors; no plot no. 5
```



Another outlier is visible again by checking the normality plot and residual versus predicted plot. It's required to be taken care of again.

6 Exercise 6.13

```
tmp <- list(A.Flow.Rate = seq(54, 60, by = 0.05), B.Dep.Time = seq(9, 16, by = 0.05))
new.data <- expand.grid(tmp)
new.data$fit <- predict(circuit.lm, new.data)
contourplot(fit ~ A.Flow.Rate + B.Dep.Time, new.data, xlab = "Cutting Speed",
            ylab = "Cutting Angle")
```

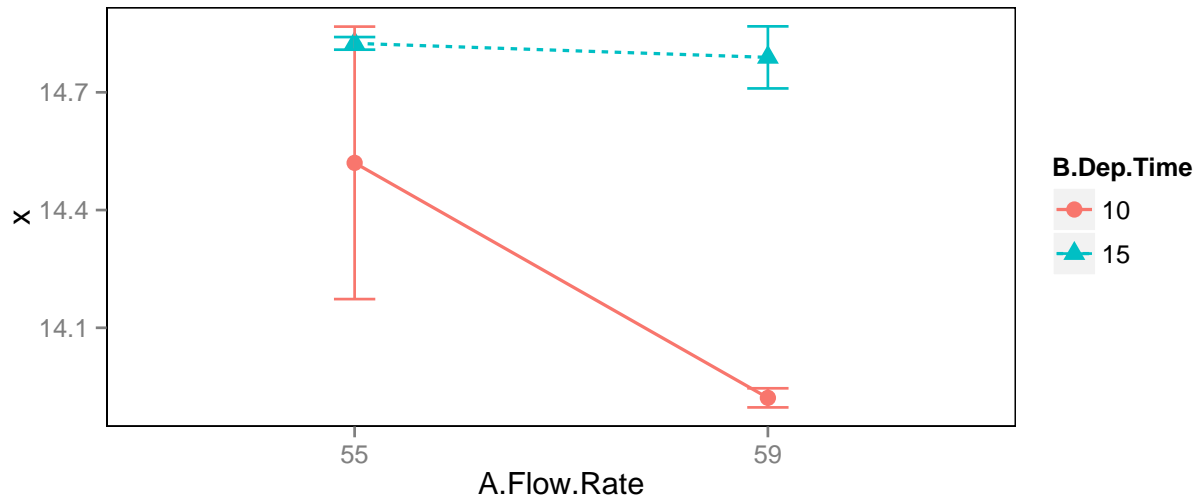


```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
circuit <- read.csv("6.12.csv")
circuit[, 1:2] <- lapply(circuit[, 1:2], factor)
library(ggplot2)
df <- with(circuit, aggregate(Thickness, list(B.Dep.Time = B.Dep.Time, A.Flow.Rate = A.Flow.Rate),
                                mean))
df$se <- with(circuit, aggregate(Thickness, list(B.Dep.Time = B.Dep.Time, A.Flow.Rate = A.Flow.Rate),
```

```

function(x) sd(x)/sqrt(10)))[, 3]
opar <- theme_update(panel.grid.major = element_blank(), panel.grid.minor = element_blank(),
  panel.background = element_rect(colour = "black"))
gp <- ggplot(df, aes(x = A.Flow.Rate, y = x, colour = B.Dep.Time, group = B.Dep.Time))
gp + geom_line(aes(linetype = B.Dep.Time), size = 0.6) + geom_point(aes(shape = B.Dep.Time),
  size = 3) + geom_errorbar(aes(ymax = x + se, ymin = x - se), width = 0.1)

```



By observing the contour plot and interaction plot, the deposition time can be recommended at 12.4 minutes (to obtain the required layer thickness). On the other hand the arsenic flow can be set at any of the experiment levels.

7 Exercise 6.14

From the countour plot and interaction plot of Problem 6.13, we observe that when the proccess would be run at a higher level of deposition time, there would be no change in the thickness value with the change of arsenic flow rate.

8 Exercise 6.26

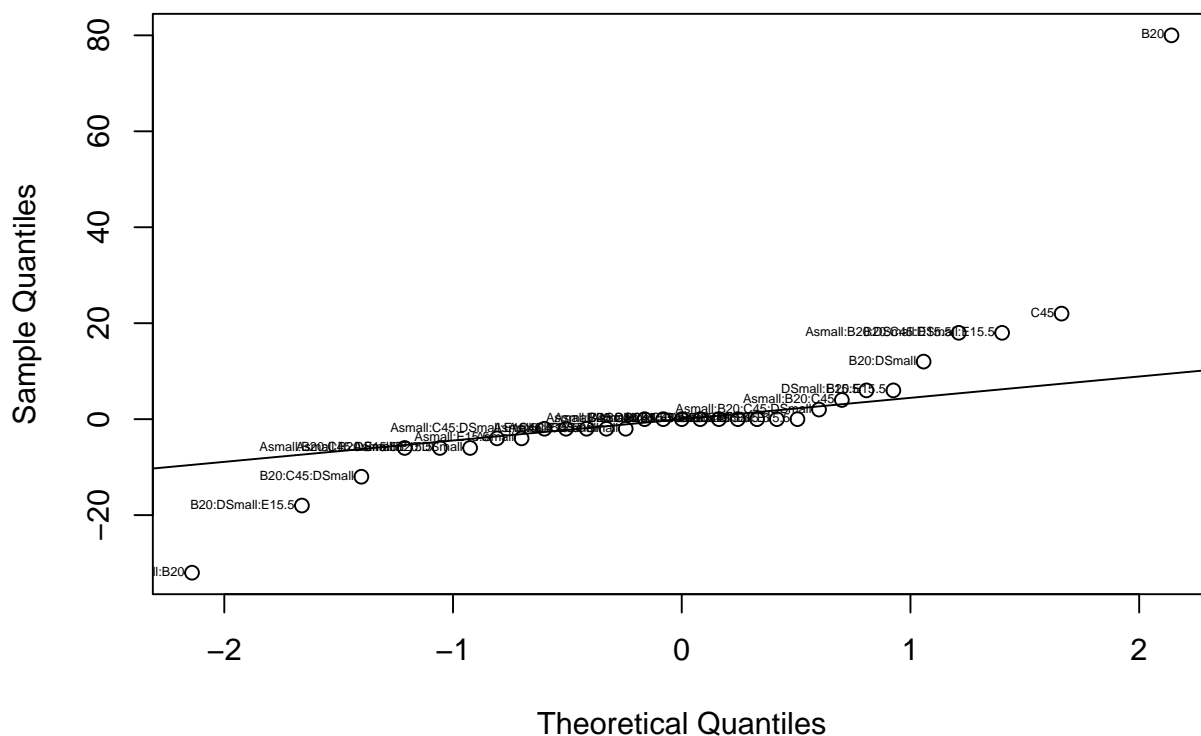
(a)

```

setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5/effects")
semicon <- read.csv("6.26a.csv")
semicon[, 1:5] <- lapply(semicon[, 1:5], factor)
mydata.lm = lm(Yield ~ A * B * C * D * E, data = semicon)
n = 1
k = 5
effects = coefficients(mydata.lm)[-c(1)] * 2
SS = effects^2 * n * 2^(k - 2)
percentage = SS/sum(SS) * 100
tem = qqnorm(effects)
qqline(effects)
text(tem$x, tem$y, names(effects), pos = 2, offset = 0.2, cex = 0.45)

```

Normal Q-Q Plot



```
cbind(effects, SS, percentage)
```

##	effects	SS	percentage
## Asmall	-4.000e+00	1.280e+02	1.699e-01
## B20	8.000e+01	5.120e+04	6.794e+01
## C45	2.200e+01	3.872e+03	5.138e+00
## DSmall	-2.000e+00	3.200e+01	4.246e-02
## E15.5	8.478e-15	5.751e-28	7.631e-31
## Asmall:B20	-3.200e+01	8.192e+03	1.087e+01
## Asmall:C45	-2.000e+00	3.200e+01	4.246e-02
## B20:C45	7.262e-15	4.219e-28	5.598e-31
## Asmall:DSmall	-1.479e-13	1.750e-25	2.322e-28
## B20:DSmall	1.200e+01	1.152e+03	1.529e+00
## C45:DSmall	-4.145e-14	1.375e-26	1.824e-29
## Asmall:E15.5	-4.000e+00	1.280e+02	1.699e-01
## B20:E15.5	6.000e+00	2.880e+02	3.822e-01
## C45:E15.5	-2.000e+00	3.200e+01	4.246e-02
## DSmall:E15.5	6.000e+00	2.880e+02	3.822e-01
## Asmall:B20:C45	4.000e+00	1.280e+02	1.699e-01
## Asmall:B20:DSmall	-6.000e+00	2.880e+02	3.822e-01
## Asmall:C45:DSmall	-2.000e+00	3.200e+01	4.246e-02
## B20:C45:DSmall	-1.200e+01	1.152e+03	1.529e+00
## Asmall:B20:E15.5	-6.000e+00	2.880e+02	3.822e-01
## Asmall:C45:E15.5	-7.154e-15	4.094e-28	5.433e-31
## B20:C45:E15.5	-3.219e-15	8.288e-29	1.100e-31
## Asmall:DSmall:E15.5	1.548e-13	1.917e-25	2.544e-28
## B20:DSmall:E15.5	-1.800e+01	2.592e+03	3.439e+00
## C45:DSmall:E15.5	4.496e-14	1.617e-26	2.146e-29
## Asmall:B20:C45:DSmall	2.000e+00	3.200e+01	4.246e-02
## Asmall:B20:C45:E15.5	-7.693e-15	4.735e-28	6.283e-31
## Asmall:B20:DSmall:E15.5	1.800e+01	2.592e+03	3.439e+00

```
## Asmall:C45:Dsmall:E15.5      -2.000e+00 3.200e+01 4.246e-02
## B20:C45:Dsmall:E15.5         1.800e+01 2.592e+03 3.439e+00
## Asmall:B20:C45:Dsmall:E15.5 -6.000e+00 2.880e+02 3.822e-01
```

From the normality plot for effects, factors B, C, A and interaction AB have larger effects.

(b)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
semicon <- read.csv("6.26.csv")
semicon.aov <- aov(Yield ~ A.Aperture + B.Exposure.Time + C.Develop.Time + A.Aperture *
  B.Exposure.Time, data = semicon)
summary(semicon.aov)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
A.Aperture	1	1116	1116	382	< 2e-16 ***
B.Exposure.Time	1	9214	9214	3155	< 2e-16 ***
C.Develop.Time	1	751	751	257	2.5e-15 ***
A.Aperture:B.Exposure.Time	1	504	504	173	3.0e-13 ***
Residuals	27	79	3		

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

The ANOVA table confirms the findings from part (a).

(c)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
semicon <- read.csv("6.26.csv")
semicon_small <- subset(semicon, A.Aperture == "small")
semicon_large <- subset(semicon, A.Aperture == "large")
semicon_small.lm <- lm(Yield ~ B.Exposure.Time + C.Develop.Time, data = semicon_small)
summary(semicon_small.lm)
```

```
##
## Call:
## lm(formula = Yield ~ B.Exposure.Time + C.Develop.Time, data = semicon_small)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.000 -1.250 -0.125  1.188  2.750
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    1.5000     2.2776   0.66    0.52
## B.Exposure.Time  0.6500     0.0223  29.10 3.2e-13 ***
## C.Develop.Time  0.6167     0.0596  10.35 1.2e-07 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.79 on 13 degrees of freedom
## Multiple R-squared:  0.987, Adjusted R-squared:  0.984
## F-statistic: 477 on 2 and 13 DF, p-value: 6.83e-13
```

```
semicon_large.lm <- lm(Yield ~ B.Exposure.Time + C.Develop.Time, data = semicon_large)
summary(semicon_large.lm)
```

```
##
## Call:
## lm(formula = Yield ~ B.Exposure.Time + C.Develop.Time, data = semicon_large)
##
```



```
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.438 -0.781 -0.375  0.875  2.688
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    11.1250     2.1158   5.26 0.00015 ***
## B.Exposure.Time  1.0469     0.0207  50.46 2.7e-16 ***
## C.Develop.Time   0.6750     0.0553  12.20 1.7e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.66 on 13 degrees of freedom
## Multiple R-squared:  0.995, Adjusted R-squared:  0.994
## F-statistic: 1.35e+03 on 2 and 13 DF, p-value: 8.48e-16
```

The regression equation (Aperture=Small):

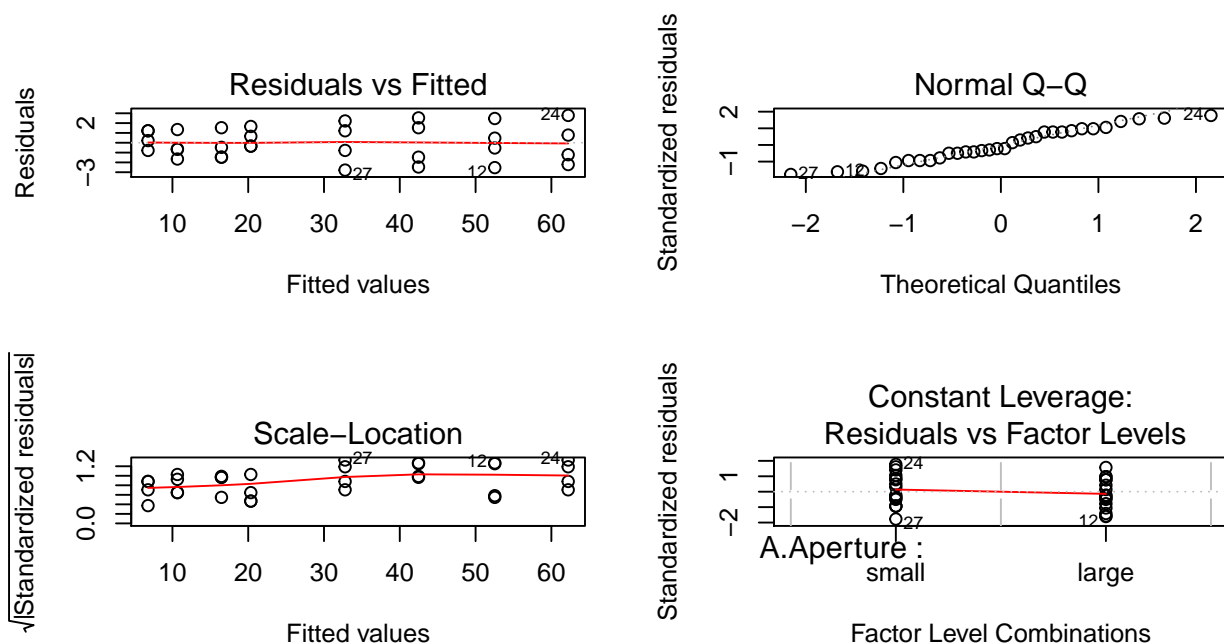
$$y = 1.5 + 0.650x_A + 0.645x_B$$

The regression equation (Aperture=Large):

$$y = 11.125 + 1.046x_A + 0.675x_B$$

(d)

```
par(mfrow = c(2, 2))
plot(semicon.aov)
```

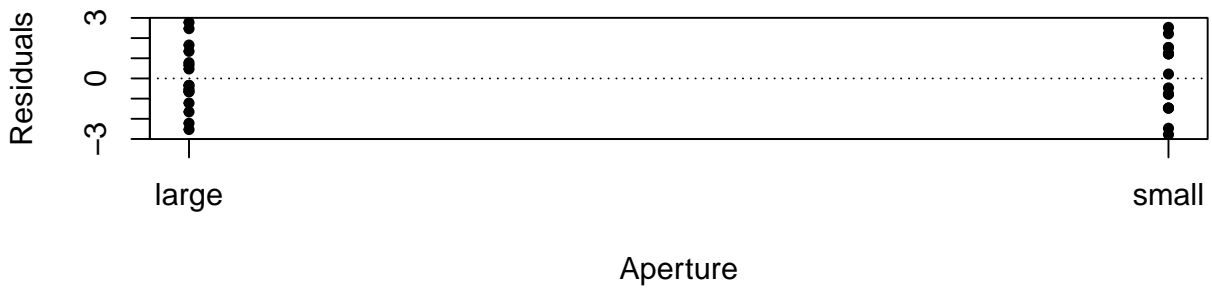


Nothing unusual is visible from the normality plot.

(e)

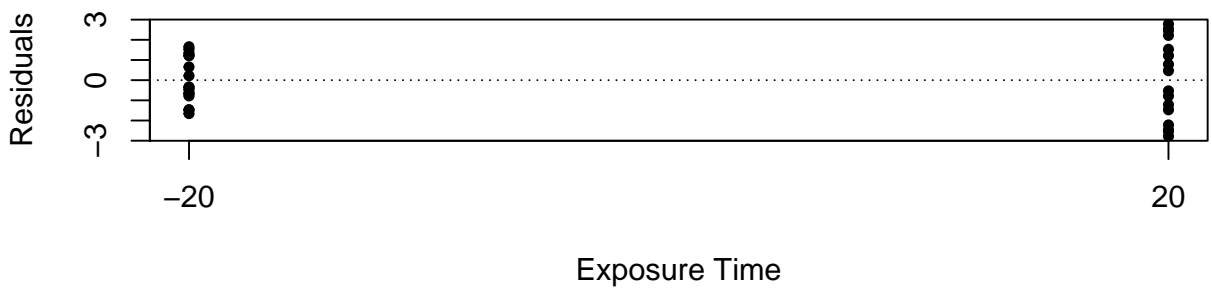
```
stripchart(residuals(semicon.aov) ~ semicon$A.Aperture, vertical = TRUE, jitter = 0,
           xlab = "Aperture", ylab = "Residuals", cex = 1, pch = 20, main = "Residuals vs. Aperture")
abline(h = 0, col = "black", lty = 3)
```

Residuals vs. Aperture



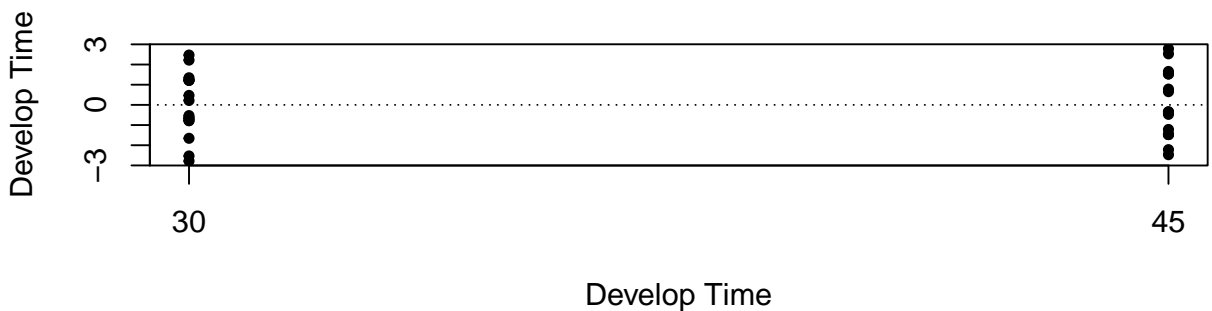
```
stripchart(residuals(semicon.aov) ~ semicon$B.Exposure.Time, vertical = TRUE,
  jitter = 0, xlab = "Exposure Time", ylab = "Residuals", cex = 1, pch = 20,
  main = "Residuals vs. Exposure Time")
abline(h = 0, col = "black", lty = 3)
```

Residuals vs. Exposure Time



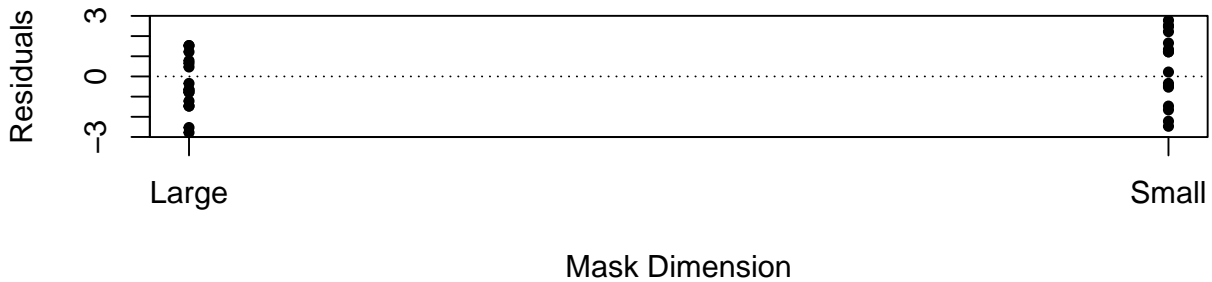
```
stripchart(residuals(semicon.aov) ~ semicon$C.Develop.Time, vertical = TRUE,
  jitter = 0, xlab = "Develop Time", ylab = "Develop Time", cex = 1, pch = 20,
  main = "Residuals vs. Develop Time")
abline(h = 0, col = "black", lty = 3)
```

Residuals vs. Develop Time



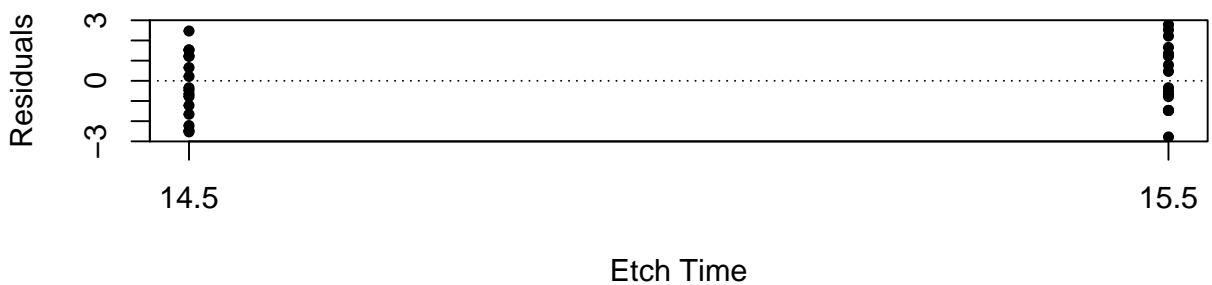
```
stripchart(residuals(semicon.aov) ~ semicon$D.Mask.Dimension, vertical = TRUE,
  jitter = 0, xlab = "Mask Dimension", ylab = "Residuals", cex = 1, pch = 20,
  main = "Residuals vs. Mask Dimension")
abline(h = 0, col = "black", lty = 3)
```

Residuals vs. Mask Dimension



```
stripchart(residuals(semicon.aov) ~ semicon$E.Etch.Time, vertical = TRUE, jitter = 0,
  xlab = "Etch Time", ylab = "Residuals", cex = 1, pch = 20, main = "Residuals vs. Etch Time")
abline(h = 0, col = "black", lty = 3)
```

Residuals vs. Etch Time

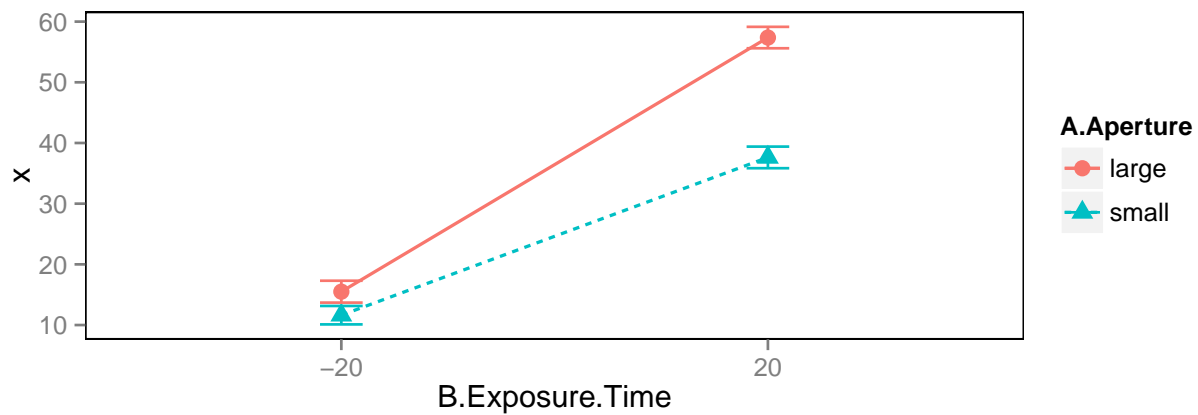


The residual versus predicted plot for exposure time show very slight amount of inequality of variance. By observing all five plots no significant problem is visible.

(f)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
semicon <- read.csv("6.26.csv")
semicon[, 1:5] <- lapply(semicon[, 1:5], factor)

library(ggplot2)
df <- with(semicon, aggregate(Yield, list(A.Aperture = A.Aperture, B.Exposure.Time = B.Exposure.Time),
  mean))
df$se <- with(semicon, aggregate(Yield, list(A.Aperture = A.Aperture, B.Exposure.Time = B.Exposure.Time),
  function(x) sd(x)/sqrt(10)))[, 3]
opar <- theme_update(panel.grid.major = element_blank(), panel.grid.minor = element_blank(),
  panel.background = element_rect(colour = "black"))
gp <- ggplot(df, aes(x = B.Exposure.Time, y = x, colour = A.Aperture, group = A.Aperture))
gp + geom_line(aes(linetype = A.Aperture), size = 0.6) + geom_point(aes(shape = A.Aperture),
  size = 3) + geom_errorbar(aes(ymax = x + se, ymin = x - se), width = 0.1)
```



From the interaction plot, we find that Factor A doesn't have much effect when B is at low level. When B is at higher level, Factor A has very large effect.

(g)

For getting higher yield, we need to run B and A at a higher level by keeping C at higher level.

9 Exercise 6.27

(a)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
semicon1 <- read.csv("6.27.csv")
semicon1a.aov <- aov(Yield ~ A.Aperture + B.Exposure.Time + C.Develop.Time +
  I(B.Exposure.Time^2) + A.Aperture * B.Exposure.Time, data = semicon1)
summary(semicon1a.aov)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## A.Aperture      1     992      992    122.6 4.1e-12 ***
## B.Exposure.Time  1    9214     9214   1138.1 < 2e-16 ***
## C.Develop.Time   1     751      751    92.7 1.1e-10 ***
## I(B.Exposure.Time^2) 1    6114     6114   755.2 < 2e-16 ***
## A.Aperture:B.Exposure.Time 1     504      504    62.3 8.3e-09 ***
## Residuals      30      243         8
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

From the ANOVA table, we find that the factor A, B, C, interaction AB and the curvature all have significant effects.

(b)

The possible next step would be adding axial points and try to fit a second-order model.

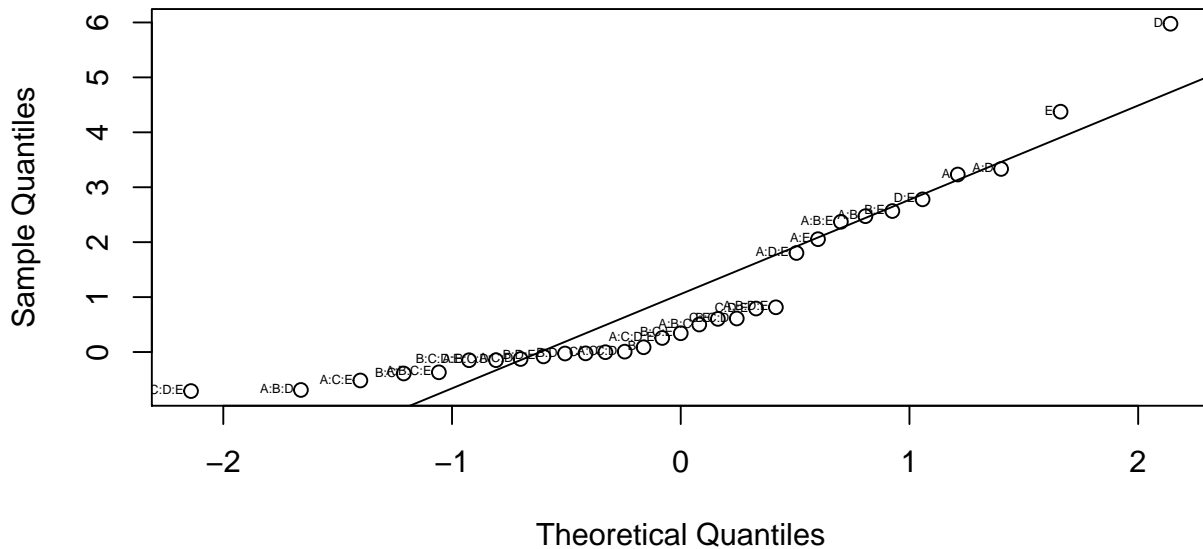
10 Exercise 6.39

(a)

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
exp <- read.csv("6.39.csv")
# exp[,1:5] <- lapply(exp[,1:5],factor)
mydata.lm = lm(y ~ A * B * C * D * E, data = exp)
n = 1
k = 5
effects = coefficients(mydata.lm)[-c(1)] * 2
SS = effects^2 * n * 2^(k - 2)
```

```
percentage = SS/sum(SS) * 100
tem = qqnorm(effects)
qqline(effects)
text(tem$x, tem$y, names(effects), pos = 2, offset = 0.2, cex = 0.45)
```

Normal Q-Q Plot



```
cbind(effects, SS, percentage)
```

##	effects	SS	percentage
## A	3.231875	8.356e+01	9.161e+00
## B	0.086875	6.038e-02	6.619e-03
## C	-0.024375	4.753e-03	5.211e-04
## D	5.976875	2.858e+02	3.133e+01
## E	4.375625	1.532e+02	1.679e+01
## A:B	2.473125	4.893e+01	5.364e+00
## A:C	-0.003125	7.813e-05	8.565e-06
## B:C	-0.390625	1.221e+00	1.338e-01
## A:D	3.333125	8.888e+01	9.744e+00
## B:D	-0.026875	5.778e-03	6.335e-04
## C:D	0.006875	3.781e-04	4.145e-05
## A:E	2.054375	3.376e+01	3.701e+00
## B:E	2.566875	5.271e+01	5.779e+00
## C:E	0.603125	2.910e+00	3.190e-01
## D:E	2.779375	6.180e+01	6.775e+00
## A:B:C	0.500625	2.005e+00	2.198e-01
## A:B:D	-0.690625	3.816e+00	4.183e-01
## A:C:D	-0.126875	1.288e-01	1.412e-02
## B:C:D	0.610625	2.983e+00	3.270e-01
## A:B:E	2.370625	4.496e+01	4.929e+00
## A:C:E	-0.518125	2.148e+00	2.354e-01
## B:C:E	0.341875	9.350e-01	1.025e-01
## A:D:E	1.803125	2.601e+01	2.851e+00
## B:D:E	-0.079375	5.040e-02	5.526e-03
## C:D:E	0.791875	5.017e+00	5.500e-01
## A:B:C:D	-0.148125	1.755e-01	1.924e-02
## A:B:C:E	-0.369375	1.092e+00	1.197e-01
## A:B:D:E	0.814375	5.306e+00	5.817e-01
## A:C:D:E	0.255625	5.228e-01	5.731e-02

```
## B:C:D:E -0.149375 1.785e-01 1.957e-02
## A:B:C:D:E -0.710625 4.040e+00 4.429e-01
```

From the normality effect plot, we find that factor D, E and interaction ABD, ACE and CDE have higher effect. No single factor have individual higher effect.

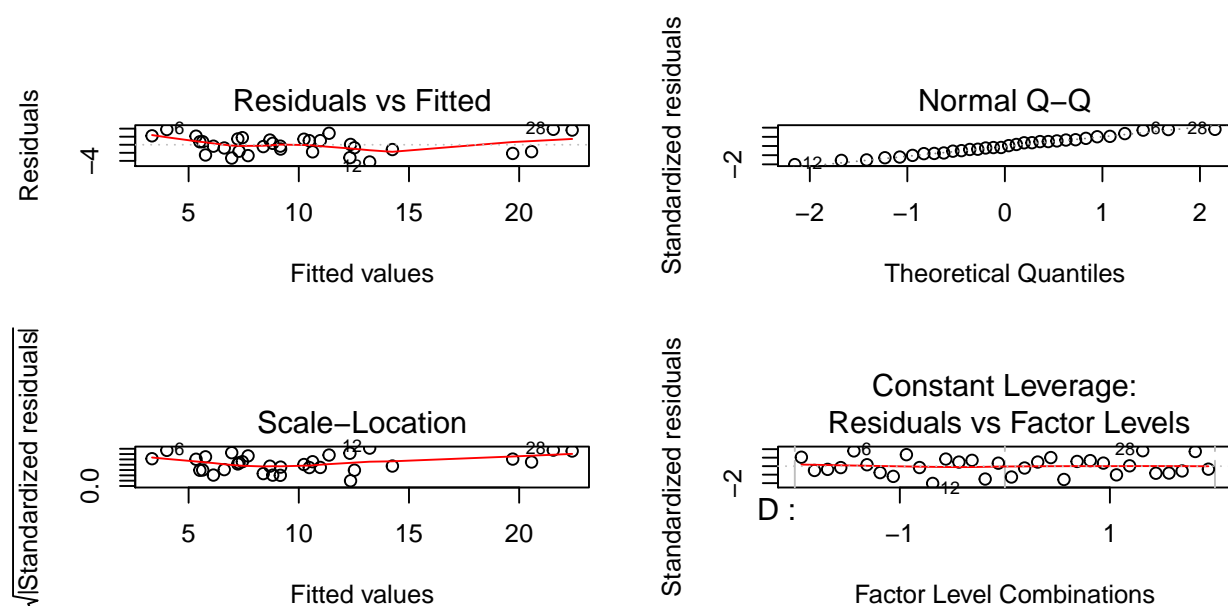
The tentative ANOVA table is shown below:

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
exp <- read.csv("6.39.csv")
exp[, 1:5] <- lapply(exp[, 1:5], factor)
mydata.aov = aov(y ~ D + E + A * B * D + C * D * E + A * C * E, data = exp)
summary(mydata.aov)
```

##		Df	Sum Sq	Mean Sq	F value	Pr(>F)
## D		1	285.8	285.8	30.12	6.2e-05 ***
## E		1	153.2	153.2	16.14	0.0011 **
## A		1	83.6	83.6	8.81	0.0096 **
## B		1	0.1	0.1	0.01	0.9375
## C		1	0.0	0.0	0.00	0.9824
## A:B		1	48.9	48.9	5.16	0.0383 *
## D:A		1	88.9	88.9	9.37	0.0079 **
## D:B		1	0.0	0.0	0.00	0.9806
## D:C		1	0.0	0.0	0.00	0.9950
## E:C		1	2.9	2.9	0.31	0.5879
## D:E		1	61.8	61.8	6.51	0.0221 *
## A:C		1	0.0	0.0	0.00	0.9977
## E:A		1	33.8	33.8	3.56	0.0787 .
## D:A:B		1	3.8	3.8	0.40	0.5355
## D:E:C		1	5.0	5.0	0.53	0.4783
## E:A:C		1	2.1	2.1	0.23	0.6411
## Residuals		15	142.3	9.5		
## ---						
## Signif. codes:						0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(b)

```
par(mfrow = c(2, 2))
plot(mydata.aov)
```



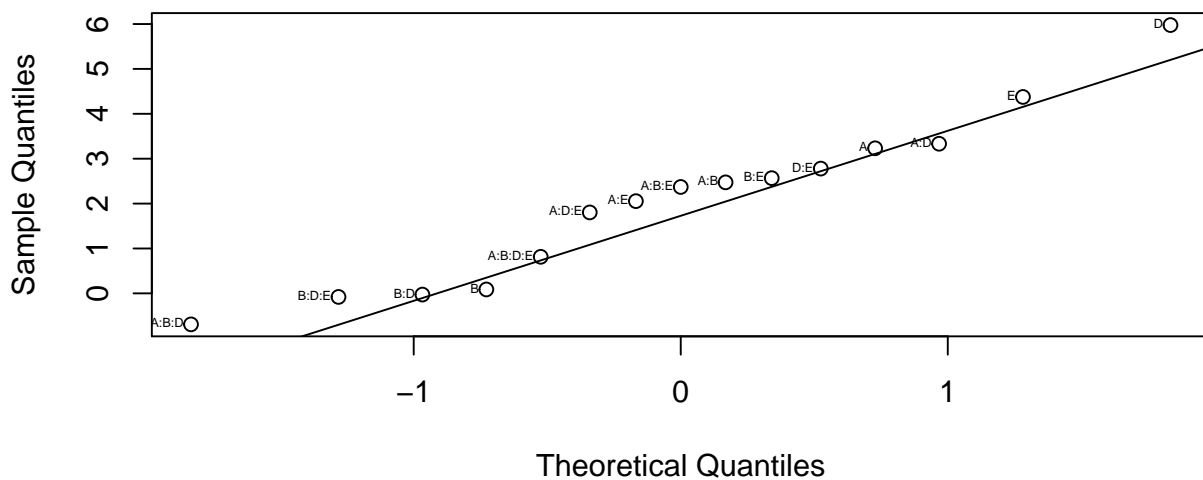
From the normality plot and residual versus predicted plot, we find that observations are 28 and 32 are outliers. This is slight indication of model inadequacy.

(c)

From the ANOVA table in part (a), we see that factor C is less important. By dropping this factor we perform the factorial design again.

```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
exp_1 <- read.csv("6.39_1.csv")
# exp[,1:5] <- lapply(exp[,1:5],factor)
mydata.lm = lm(y ~ A * B * D * E, data = exp_1)
n = 1
k = 4
effects = coefficients(mydata.lm)[-c(1)] * 2
SS = effects^2 * n * 2^(k - 2)
percentage = SS/sum(SS) * 100
tem = qqnorm(effects)
qqline(effects)
text(tem$x, tem$y, names(effects), pos = 2, offset = 0.2, cex = 0.45)
```

Normal Q-Q Plot



```
cbind(effects, SS, percentage)
```

##	effects	SS	percentage
## A	3.23188	4.178e+01	9.401e+00
## B	0.08688	3.019e-02	6.793e-03
## D	5.97687	1.429e+02	3.215e+01
## E	4.37562	7.658e+01	1.723e+01
## A:B	2.47312	2.447e+01	5.505e+00
## A:D	3.33313	4.444e+01	1.000e+01
## B:D	-0.02688	2.889e-03	6.501e-04
## A:E	2.05437	1.688e+01	3.799e+00
## B:E	2.56688	2.636e+01	5.931e+00
## D:E	2.77938	3.090e+01	6.953e+00
## A:B:D	-0.69062	1.908e+00	4.293e-01
## A:B:E	2.37063	2.248e+01	5.058e+00
## A:D:E	1.80312	1.301e+01	2.926e+00
## B:D:E	-0.07937	2.520e-02	5.671e-03
## A:B:D:E	0.81437	2.653e+00	5.969e-01

```

### ANOVA Table
mydata.aov2 = aov(y ~ D + E + A * B * D, data = exp)
summary(mydata.aov2)

##           Df Sum Sq Mean Sq F value    Pr(>F)
## D           1  285.8   285.8    26.51 3.2e-05 ***
## E           1  153.2   153.2    14.21 0.0010 ***
## A           1   83.6    83.6     7.75 0.0105 *
## B           1    0.1     0.1     0.01 0.9410
## A:B         1   48.9    48.9     4.54 0.0441 *
## D:A         1   88.9    88.9     8.24 0.0086 **
## D:B         1    0.0     0.0     0.00 0.9817
## D:A:B       1    3.8     3.8     0.35 0.5577
## Residuals   23  248.0    10.8
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

(d)

The settings of the active factors will be consisted of A, D and E to find the value of y maximum.