

STAT 522 — Assignment 5

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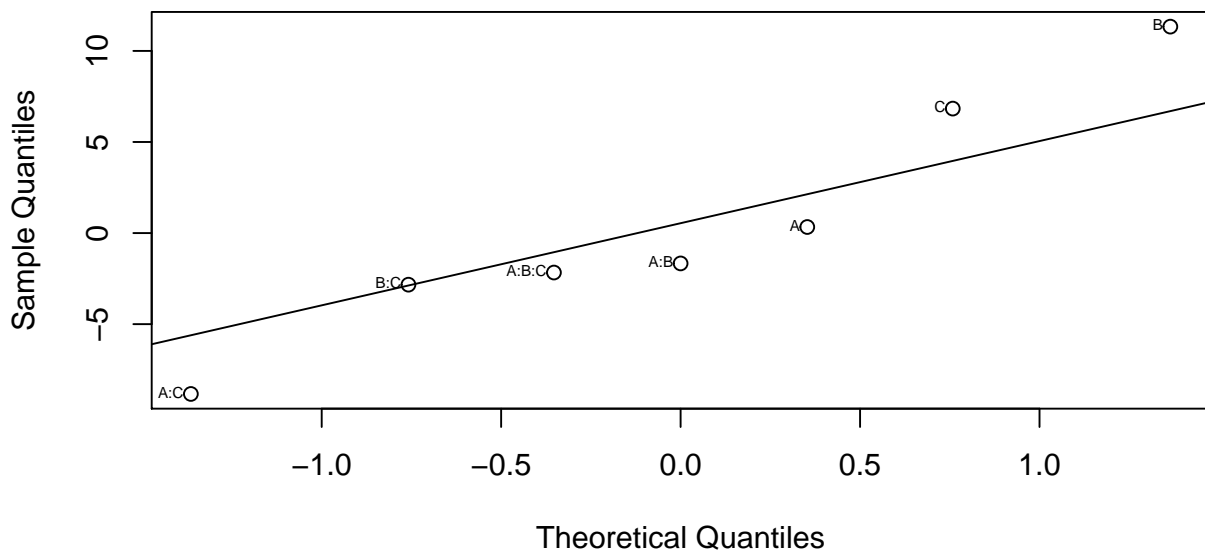
14th April 2014

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
```

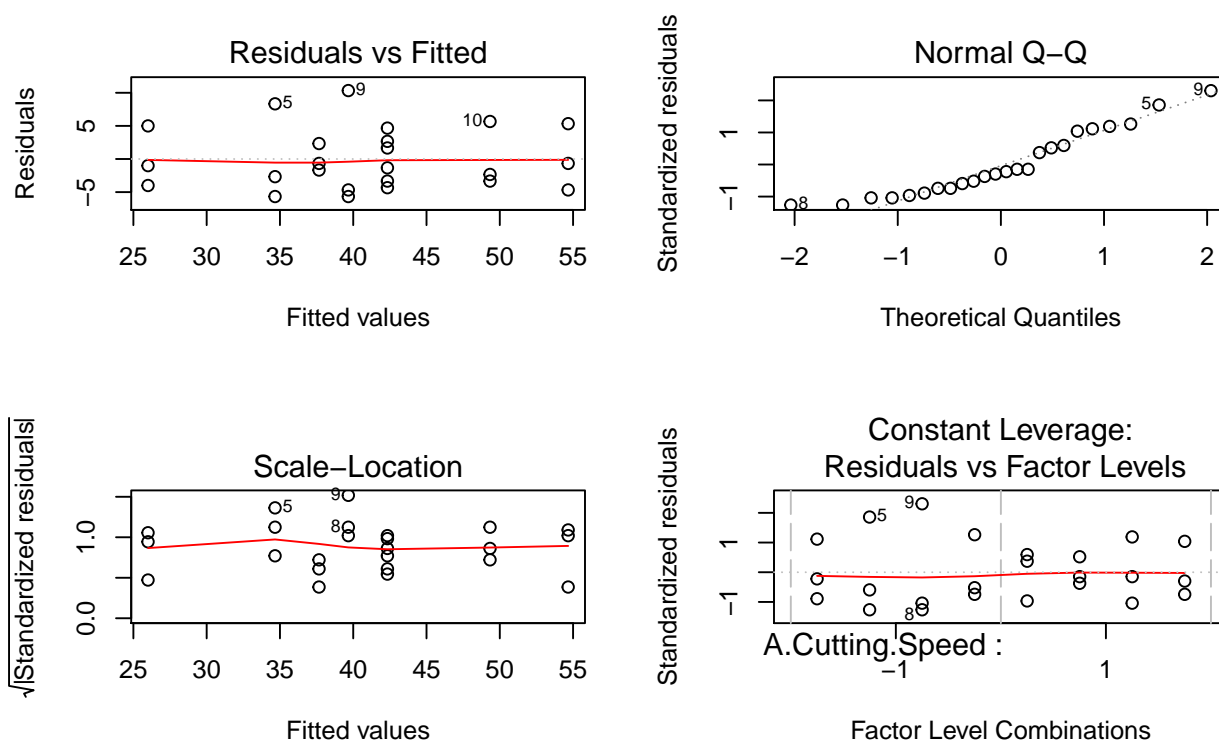
The regression model:

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

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

(d)

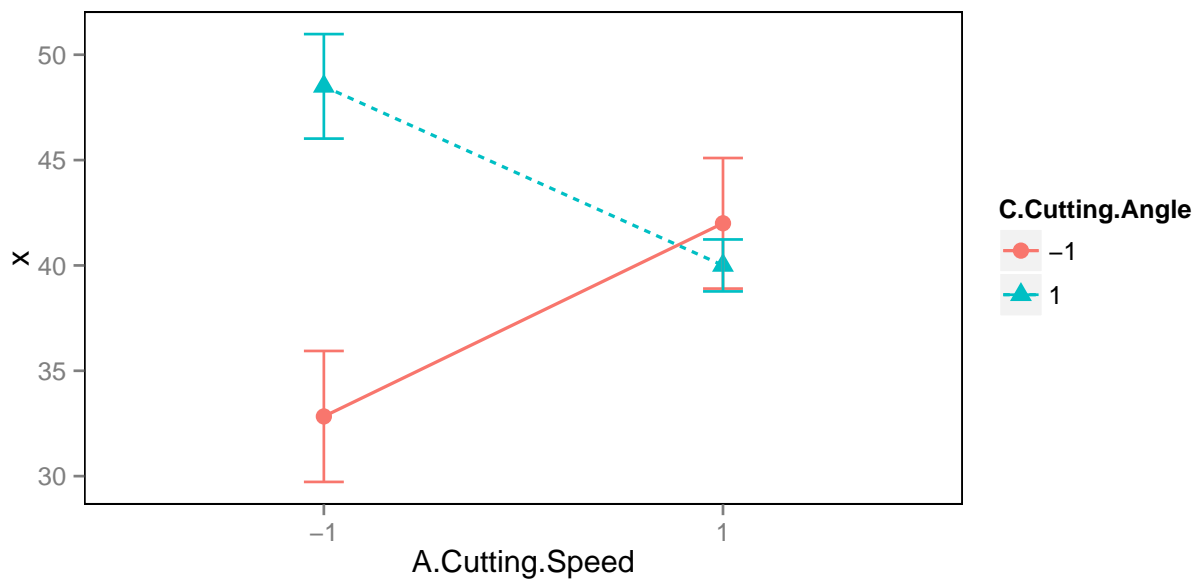
```
par(mfrow = c(2, 2))
plot(tool1.aov)
```



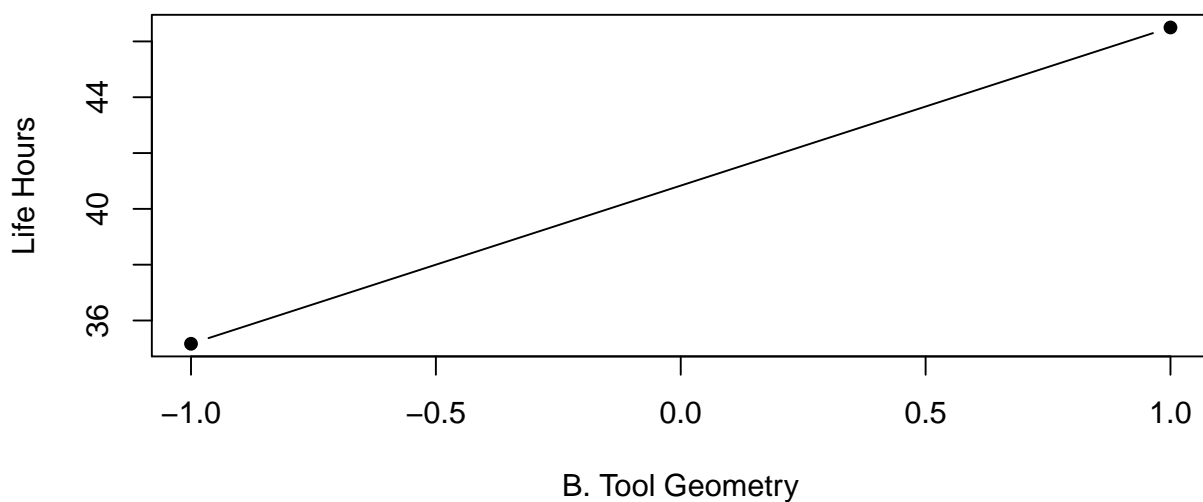
Nothing unusual is visible regarding the residual plots.

(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"))
```

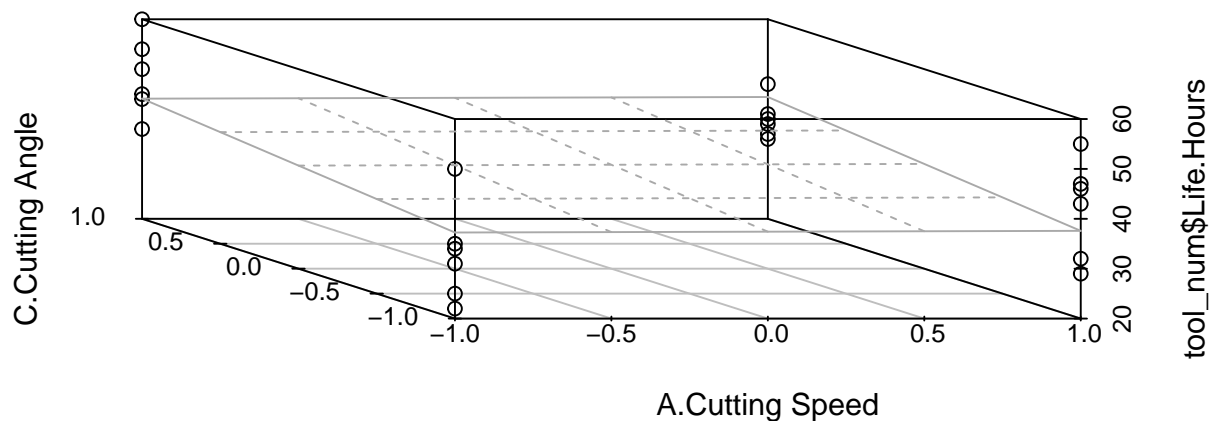


As B has positive effect, we can set B at a high level to increase the life hours. The interaction plot of AC indicates that life hours will be maximum at higher level of C and lower level of A.

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)

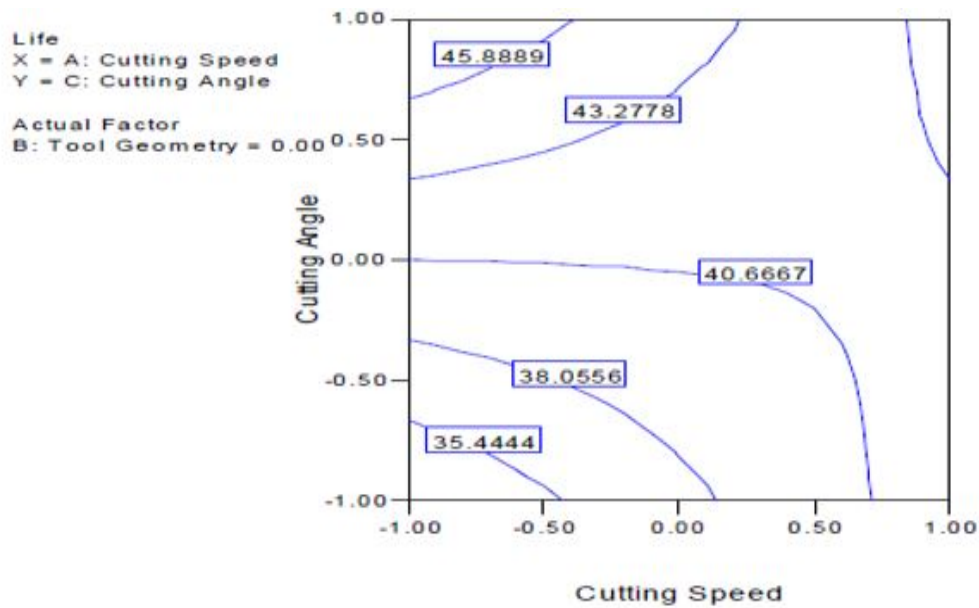
library(scatterplot3d)
s3d <- scatterplot3d(tool_num$A.Cutting.Speed, tool_num$C.Cutting.Angle, tool_num$Life.Hours,
  type = "p", angle = 135, scale.y = 1, xlab = "A.Cutting Speed", ylab = "C.Cutting Angle")
s3d$plane3d(tool_num.lm, lty.box = "solid", col = "darkgray")
```



```
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

c <- contourplot(fit ~ A.Cutting.Speed * C.Cutting.Angle, new.data, xlab = "A.Cutting Speed",
  ylab = "C.Cutting Angle")
# plot in minitab
```



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 (From ANOVA table)
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

## 95% Confidence Limit (by Using Effects value)
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
```

Variable	Effect	Conf. Int.
A	0.333	±4.395
B	11.333	±4.395
AC	-1.667	±4.395
C	6.833	±4.395
AC	-8.833	±4.395
BC	-2.883	±4.395
ABC	-2.167	±4.395

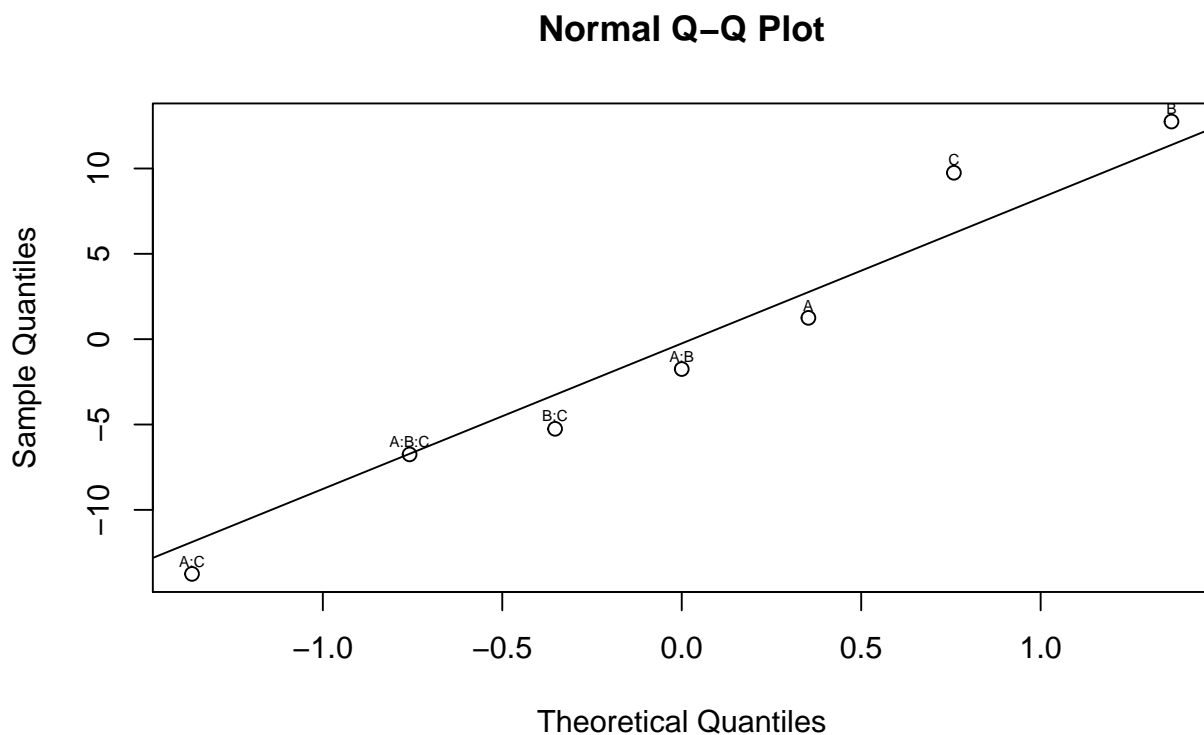
The standard error is 2.24.

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

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

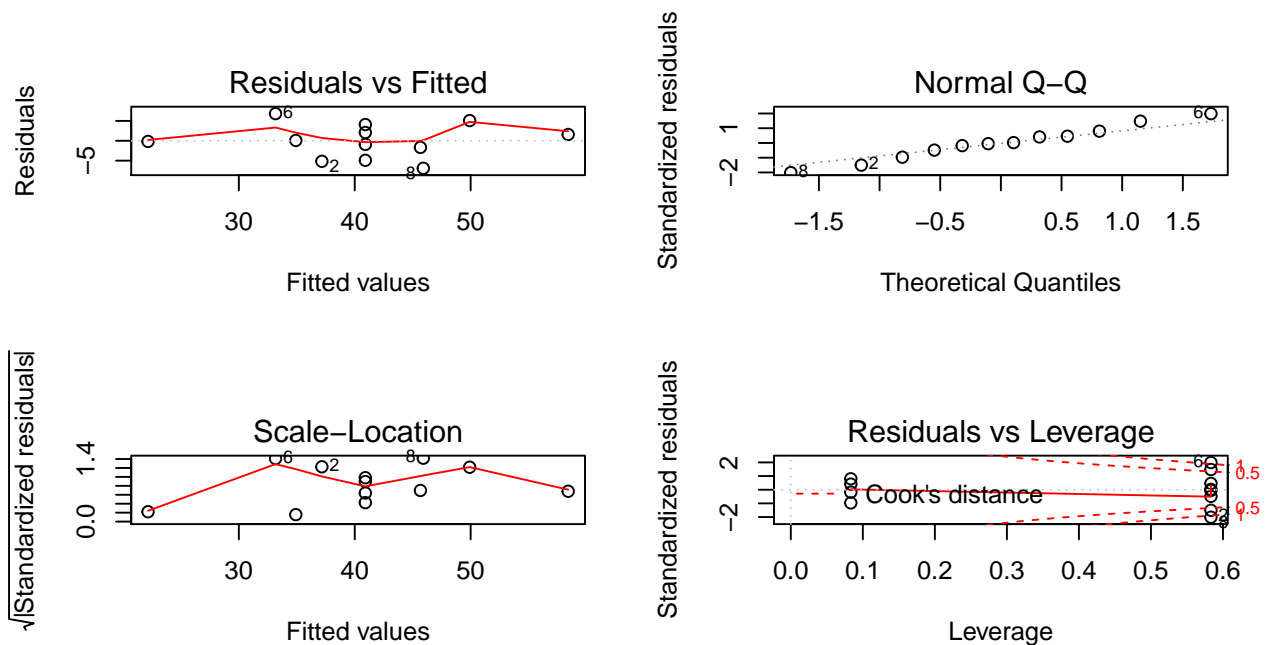
The regression model from 6.1 part (c):

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

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)

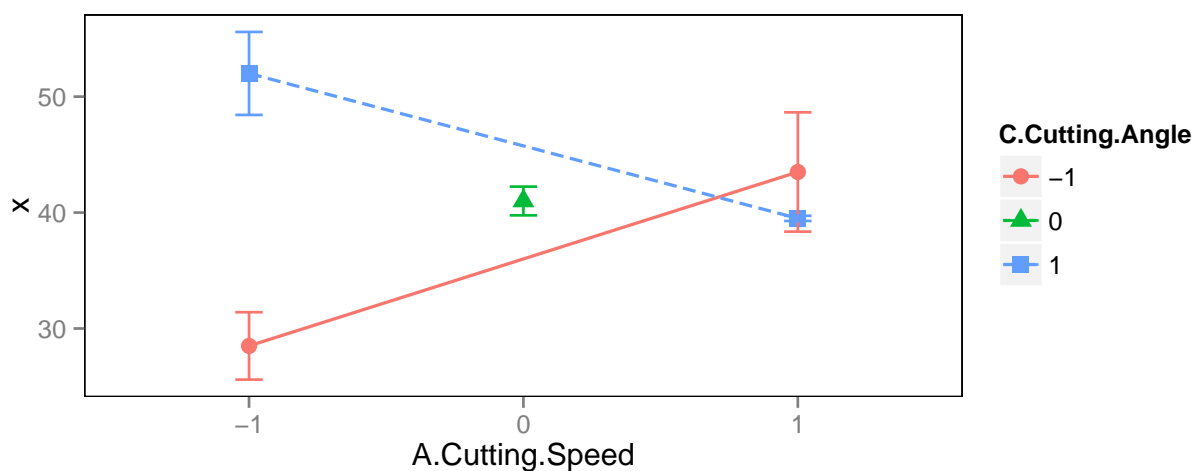
```
par(mfrow = c(2, 2))
plot(tool2a.aov)
```



Nothing unusual is 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("6.12b.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   -0.3173 0.8052      19.23
## B    0.5860 2.7472      65.62
## A:B   0.2815 0.6339      15.14
```

From the factor effect values, 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
```

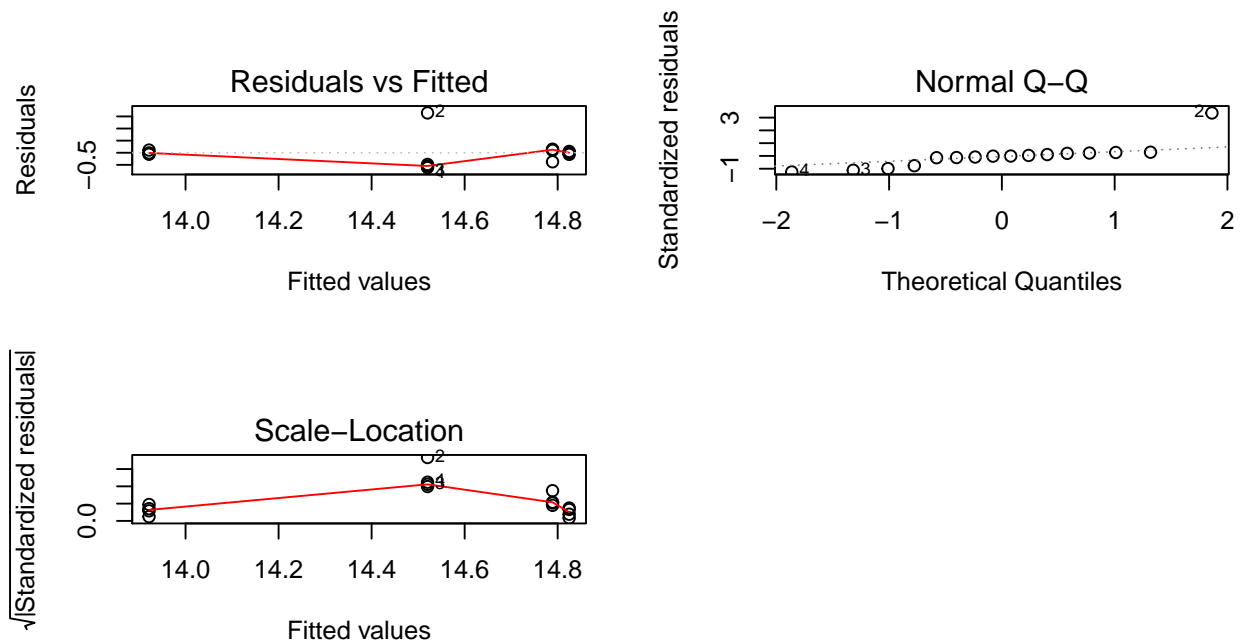
The regression equation:

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

(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 finds out any error, it should be corrected.

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

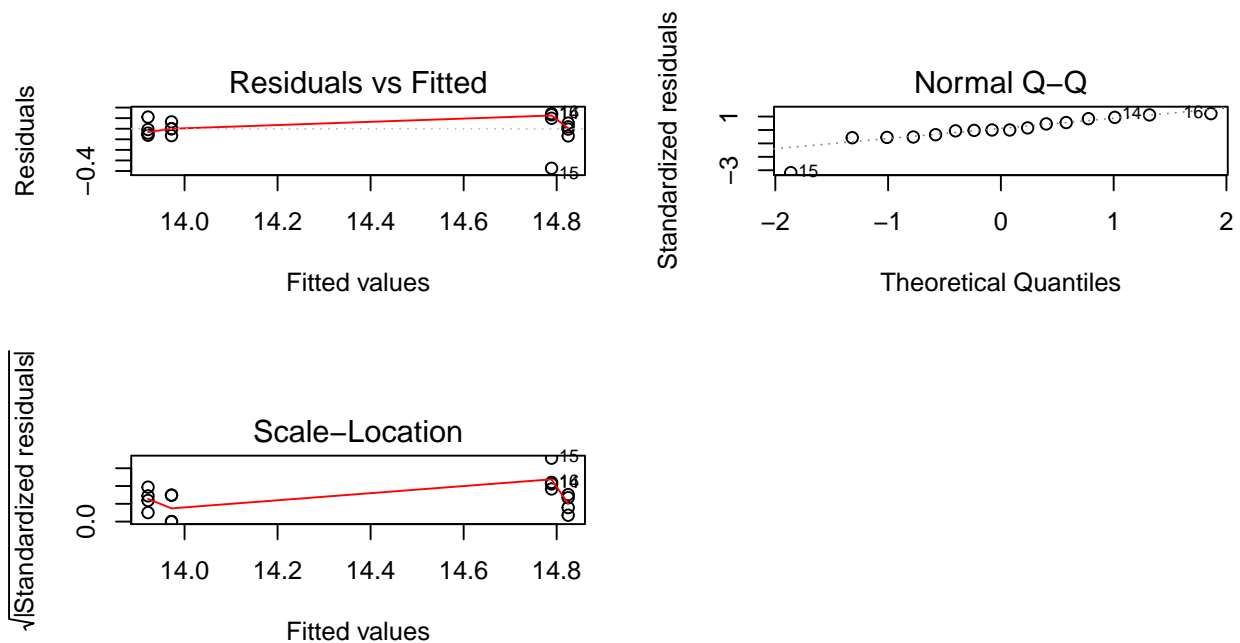
```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
circuit_1 <- read.csv("6.12_1.csv")
circuit_1.aov <- aov(Thickness ~ A.Flow.Rate * B.Dep.Time, data = circuit_1)
summary(circuit_1.aov)
```

	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)

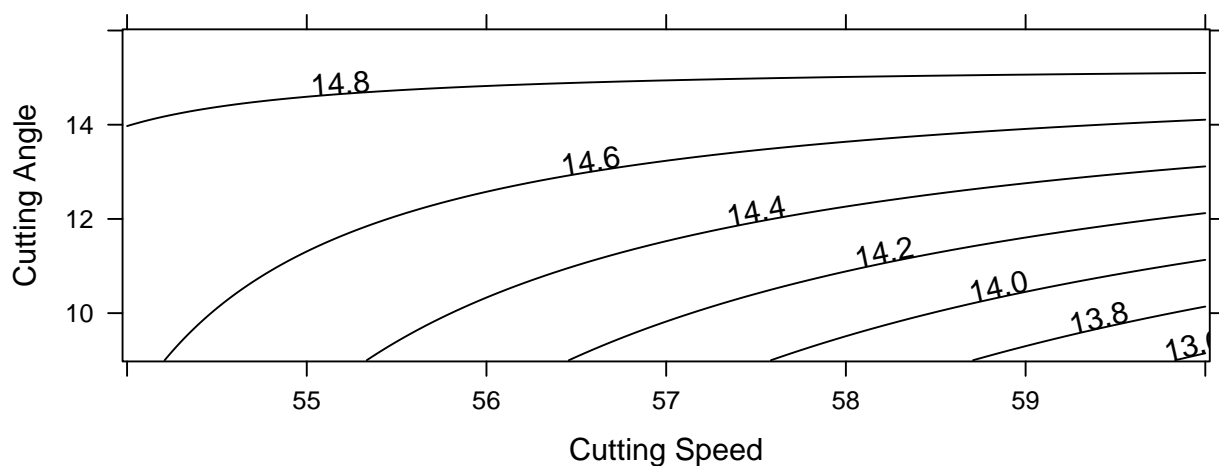
## hat values (leverages) are all = 0.25
## and there are no factor predictors; no plot no. 5
```



The significance of B is improved by changing the observation value of 2. Improvement is visible by checking the normality plot and residual versus predicted plot.

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")
```

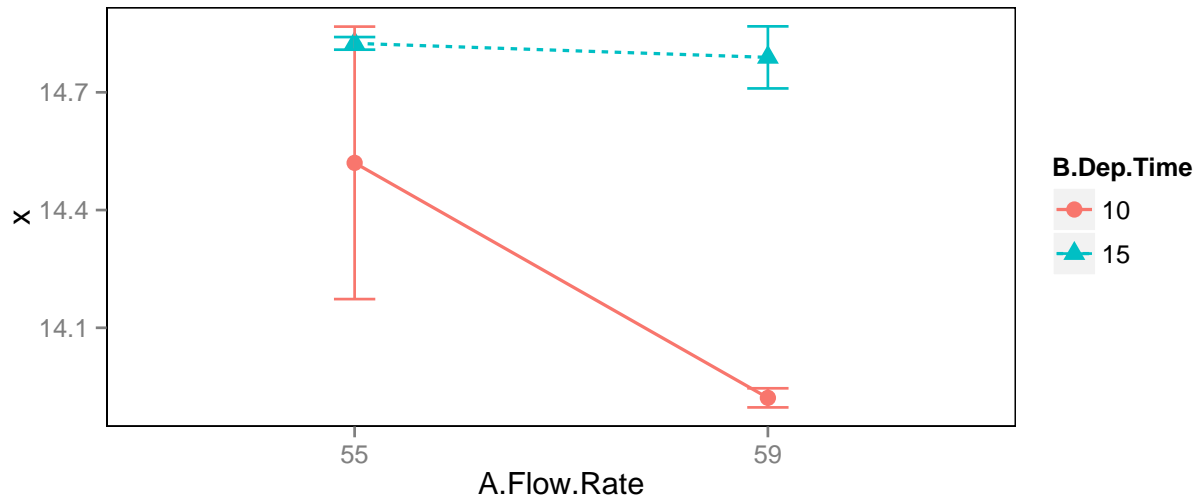


```
setwd("C:/Users/Subasish/Dropbox/A Spring 2014/Dr Novelo/HW/HW5")
circuit <- read.csv("6.12.csv")
circuit[, 1:2] <- lapply(circuit[, 1:2], factor)
library(ggplot2)
df <- with(circuit, aggregate(Thickness, list(B.Dep.Time = B.Dep.Time, A.Flow.Rate = A.Flow.Rate),
                                mean))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

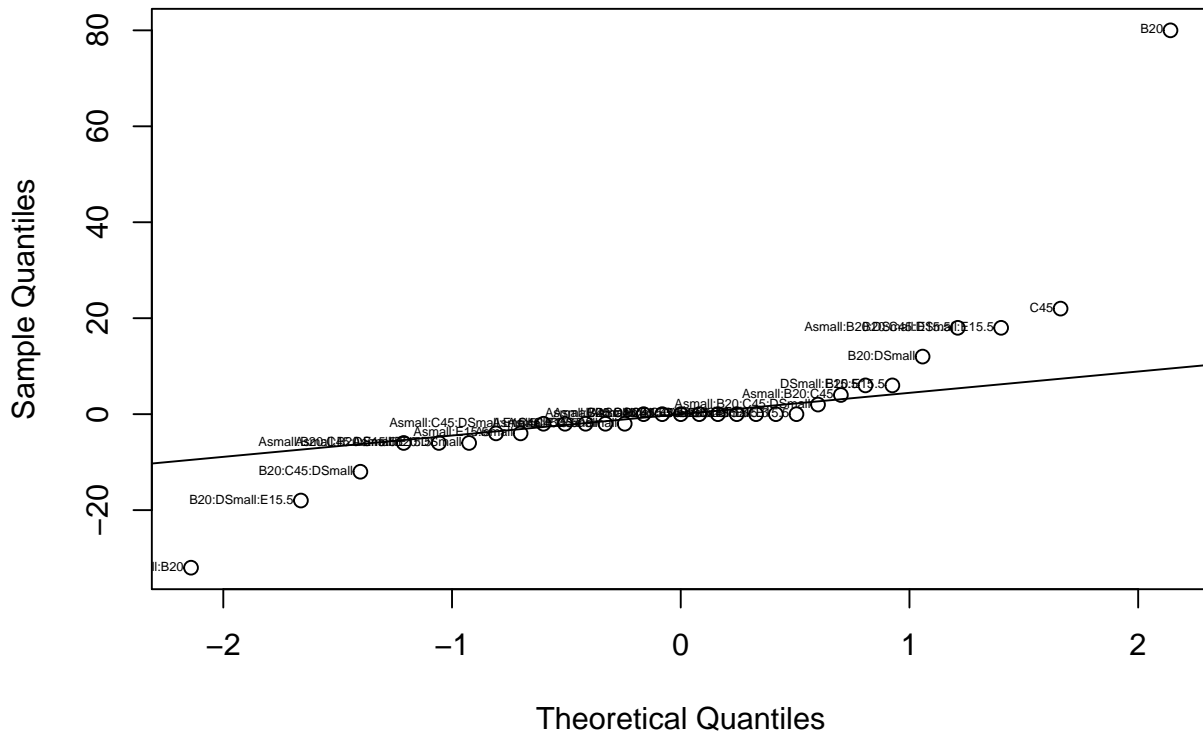
(a)

```

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):

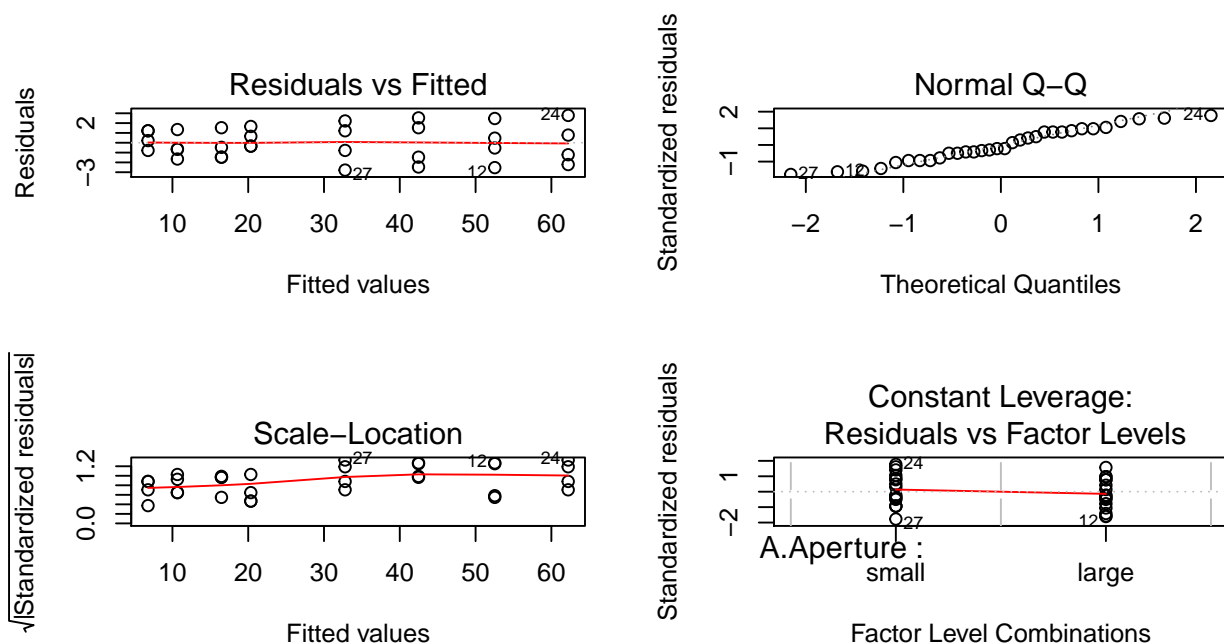
$$y = 1.5 + 0.650x_A + 0.645x_B$$

The regression equation (Aperture=Large):

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

(d)

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

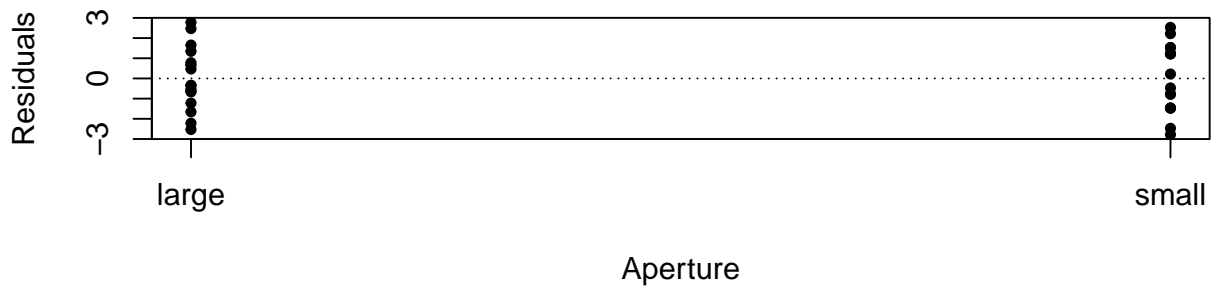


Nothing unusual is visible from the normality plot.

(e)

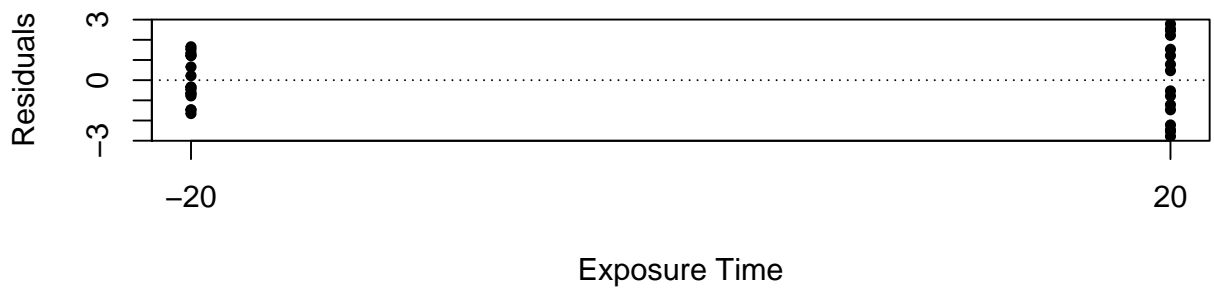
```
stripchart(residuals(semicon.aov) ~ semicon$A.Aperture, vertical = TRUE, jitter = 0,
           xlab = "Aperture", ylab = "Residuals", cex = 1, pch = 20, main = "Residuals vs. Aperture")
abline(h = 0, col = "black", lty = 3)
```

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



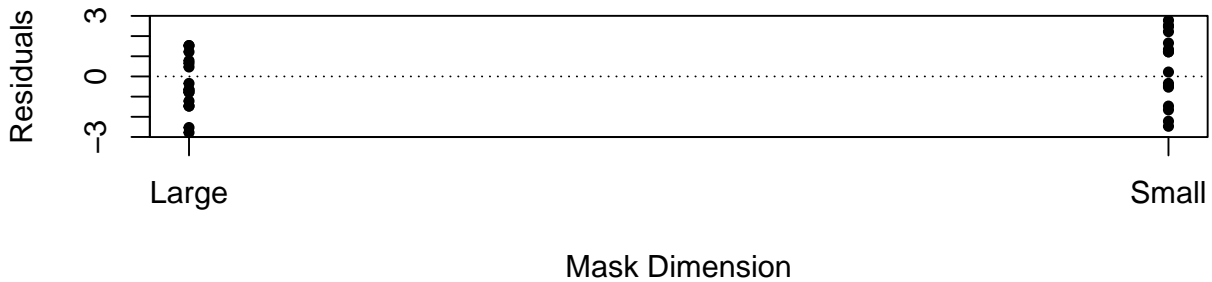
```
stripchart(residuals(semicon.aov) ~ semicon$C.Develop.Time, vertical = TRUE,
  jitter = 0, xlab = "Develop Time", ylab = "Develop Time", cex = 1, pch = 20,
  main = "Residuals vs. Develop Time")
abline(h = 0, col = "black", lty = 3)
```

Residuals vs. Develop Time



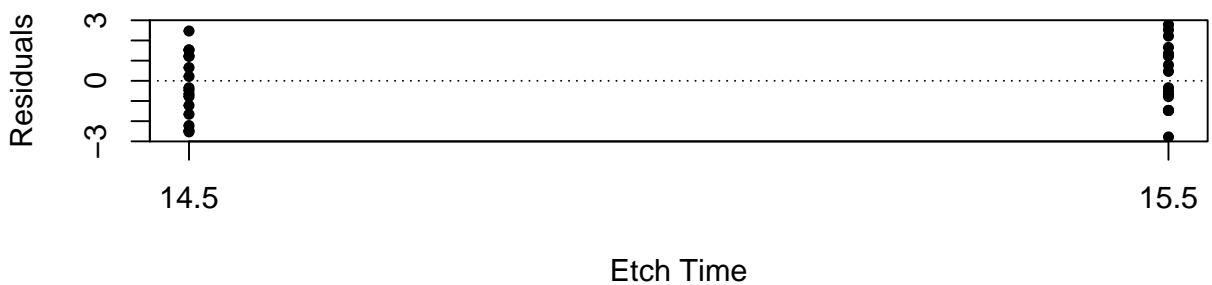
```
stripchart(residuals(semicon.aov) ~ semicon$D.Mask.Dimension, vertical = TRUE,
  jitter = 0, xlab = "Mask Dimension", ylab = "Residuals", cex = 1, pch = 20,
  main = "Residuals vs. Mask Dimension")
abline(h = 0, col = "black", lty = 3)
```

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

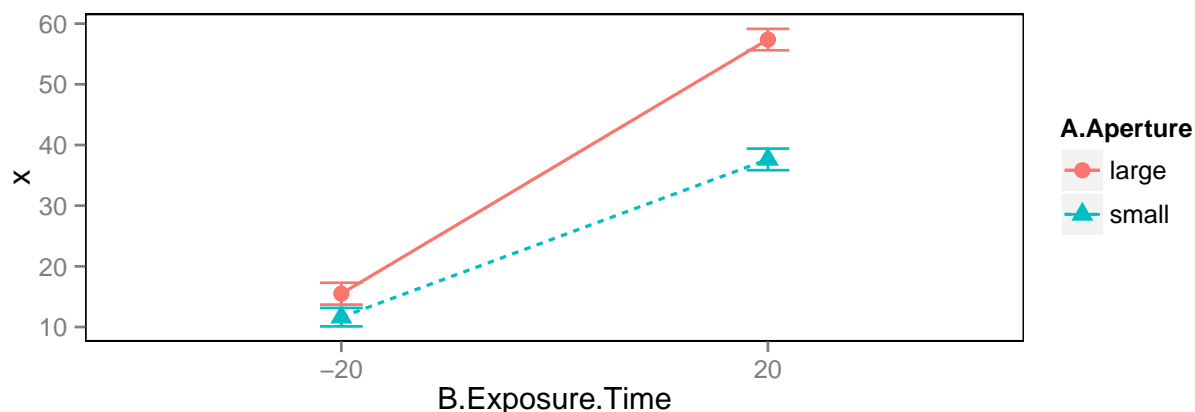


The residual versus predicted plot for exposure time show very slight amount of inequality of variance. By observing all five plots no significant problem is visible.

(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.

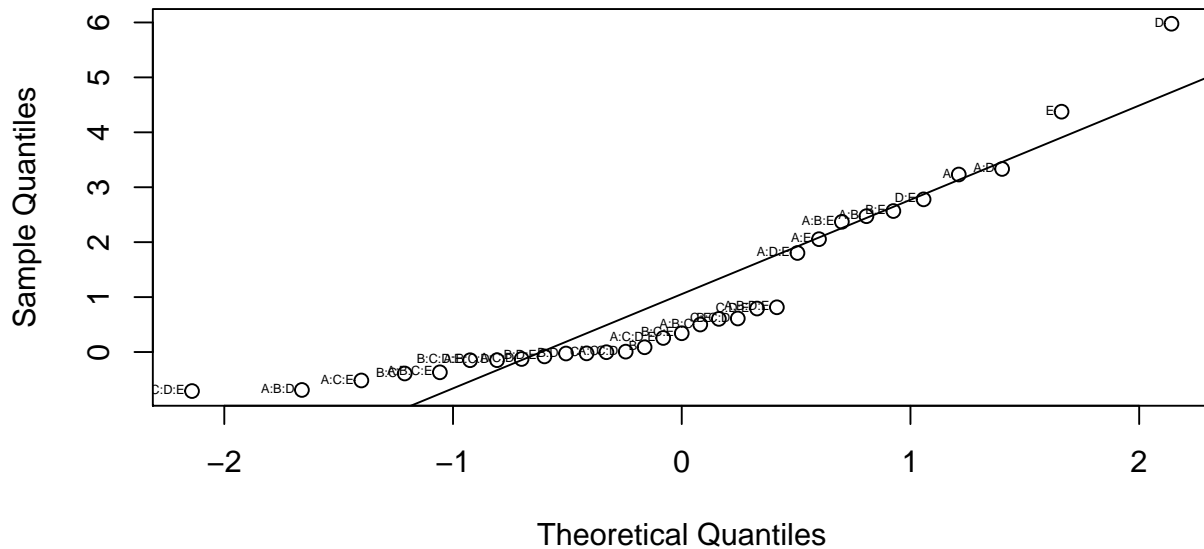
10 Exercise 6.39

(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.

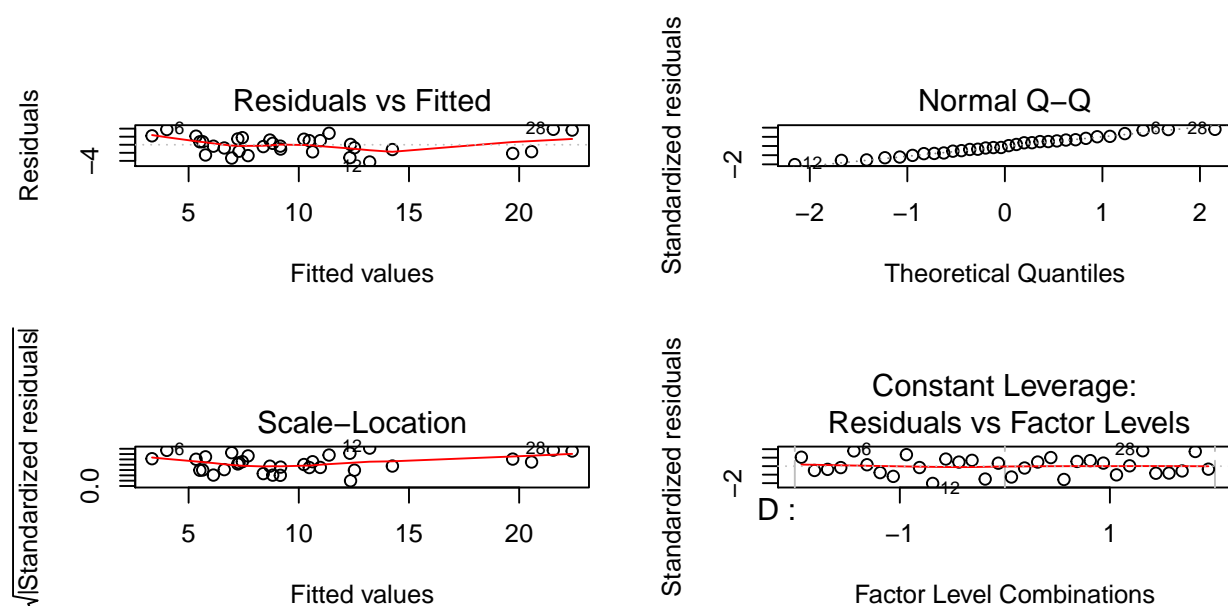
The tentative ANOVA table is shown below:

```
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)
```



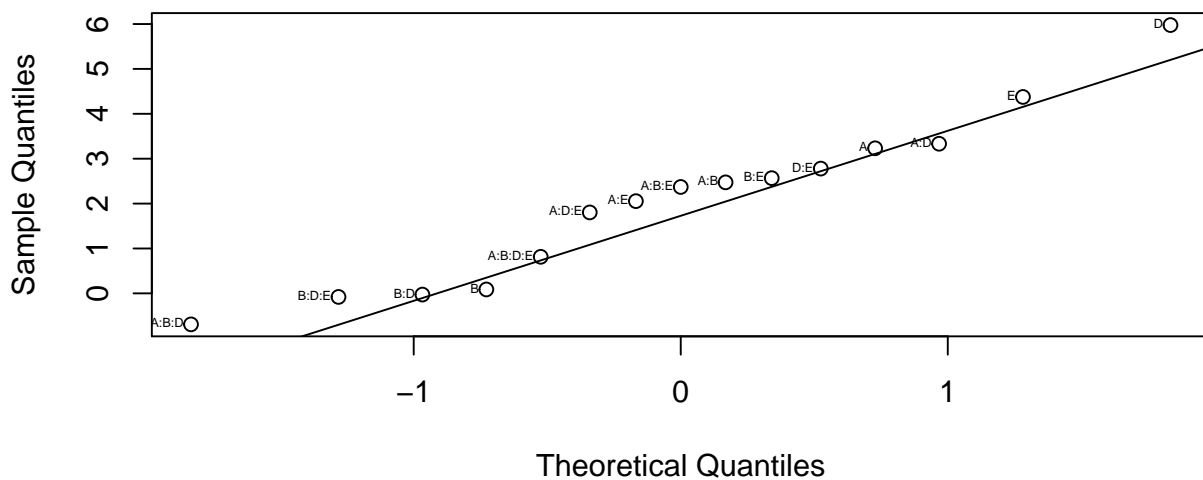
From the normality plot and residual versus predicted plot, we find that observations are 28 and 32 are outliers. This is slight indication of model inadequacy.

(c)

From the ANOVA table in part (a), we see that factor C is less significant. 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

```

By comparing the two ANOVA tables, we found that in the later design the significance of D and E are improved.

(d)

```

### ANOVA Table
aov_1 = aov(y ~ D + E + A, data = exp)
summary(aov_1)

##           Df Sum Sq Mean Sq F value    Pr(>F)
## D           1   286   285.8    20.5 1e-04 ***
## E           1   153   153.2    11.0 0.0025 **
## A           1    84    83.6     6.0 0.0208 *
## Residuals   28   390    13.9
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

aov_2 = aov(y ~ D + E + A + D * A, data = exp)
summary(aov_2)

##           Df Sum Sq Mean Sq F value    Pr(>F)
## D           1  285.8   285.8    25.65 2.6e-05 ***
## E           1  153.2   153.2    13.75 0.00095 ***
## A           1   83.6    83.6     7.50 0.01079 *
## D:A          1   88.9    88.9     7.98 0.00879 **
## Residuals   27  300.8    11.1
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

aov_3 = aov(y ~ D + E + A + D * A + A * B, data = exp)
summary(aov_3)

##           Df Sum Sq Mean Sq F value    Pr(>F)
## D           1  285.8   285.8    28.38 1.6e-05 ***
## E           1  153.2   153.2    15.21 0.00064 ***
## A           1   83.6    83.6     8.30 0.00803 **
## B           1    0.1     0.1     0.01 0.93890
## D:A          1   88.9    88.9     8.82 0.00648 **
## A:B          1   48.9    48.9     4.86 0.03694 *
## Residuals   25  251.8    10.1
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

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