

# Stat 5309 Lab 6

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## 1

```
bit_sizes <- c("1/16", "1/8")
speeds <- c("40", "90")
treatments <- expand.grid(bit_size=rep(bit_sizes, 4), speed=speeds)
treatments
```

```
##   bit_size speed
## 1    1/16    40
## 2    1/8    40
## 3    1/16    40
## 4    1/8    40
## 5    1/16    40
## 6    1/8    40
## 7    1/16    40
## 8    1/8    40
## 9    1/16    90
## 10   1/8    90
## 11   1/16    90
## 12   1/8    90
## 13   1/16    90
## 14   1/8    90
## 15   1/16    90
## 16   1/8    90
```

```
circuit_data <- data.frame(treatments,
                           vibration = c(18.2, 27.2,
                                         18.9, 24.0,
                                         12.9, 22.4,
                                         14.4, 22.5,
                                         15.9, 41.0,
                                         14.5, 43.9,
                                         15.1, 36.3,
                                         14.2, 39.9
                                         )
                           )
circuit_data %>% kable()
```

bit_size	speed	vibration
1/16	40	18.2
1/8	40	27.2
1/16	40	18.9
1/8	40	24.0
1/16	40	12.9
1/8	40	22.4
1/16	40	14.4

bit_size	speed	vibration
1/8	40	22.5
1/16	90	15.9
1/8	90	41.0
1/16	90	14.5
1/8	90	43.9
1/16	90	15.1
1/8	90	36.3
1/16	90	14.2
1/8	90	39.9

**a**

analyze the data from this experiment.

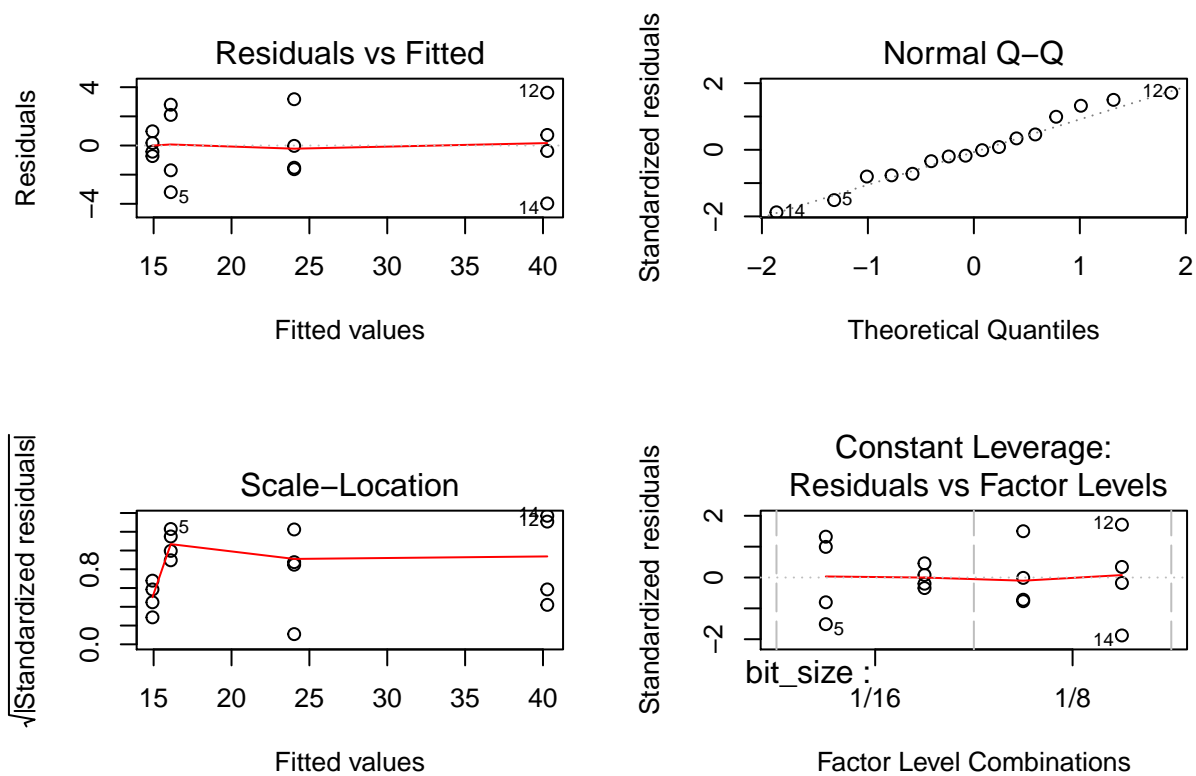
```
vibration_model <- aov(formula = vibration~bit_size*speed, data=circuit_data)
summary(vibration_model)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## bit_size      1 1107.2  1107.2   185.25 1.17e-08 ***
## speed         1  227.3   227.3    38.02 4.83e-05 ***
## bit_size:speed 1  303.6   303.6    50.80 1.20e-05 ***
## Residuals    12   71.7     6.0
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

**b**

construct a normal probability plot of the residuals, and plot the residuals versus the predicted vibration level. Interpret these plots.

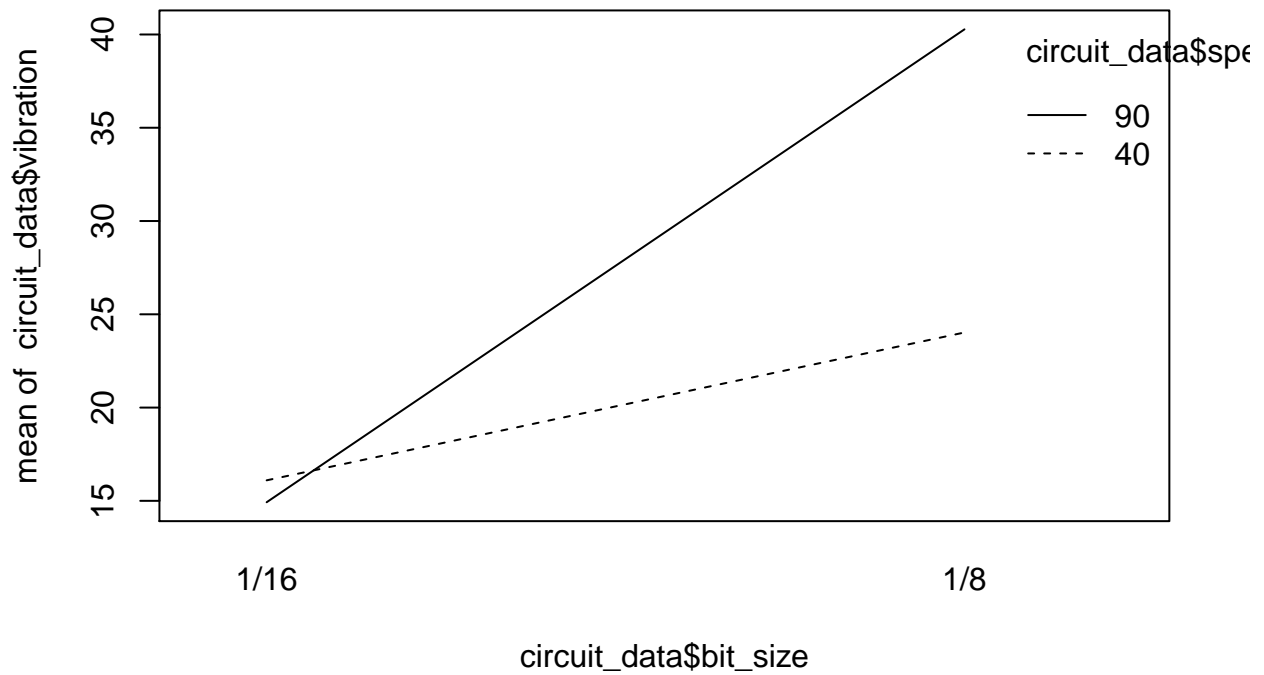
```
par( mfrow = c(2,2) )
plot(vibration_model)
```



c

Draw the AB interaction plot. What levels of bit size and speed would you recommend for routine operation?

```
interaction.plot(x.factor = circuit_data$bit_size,
               trace.factor = circuit_data$speed,
               response = circuit_data$vibration)
```



2

```
cutting_speeds <- c("-", "+")
tool_geometries <- c("-", "+")
cutting_angles <- c("-", "+")
machine_trts <- expand.grid(cutting_speed=rep(cutting_speeds,3),
                           tool_geometry=tool_geometries,
                           cutting_angle=cutting_angles)
machine_trts
```

```
##      cutting_speed tool_geometry cutting_angle
## 1              -              -              -
## 2              +              -              -
## 3              -              -              -
## 4              +              -              -
## 5              -              -              -
## 6              +              -              -
## 7              -              +              -
## 8              +              +              -
## 9              -              +              -
## 10             +              +              -
## 11             -              +              -
## 12             +              +              -
## 13             -              -              +
```

```
## 14      +      -      +
## 15      -      -      +
## 16      +      -      +
## 17      -      -      +
## 18      +      -      +
## 19      -      +      +
## 20      +      +      +
## 21      -      +      +
## 22      +      +      +
## 23      -      +      +
## 24      +      +      +
```

```
machine_data <- data.frame(machine_trts,
                           lifetime = c(22,32,
                                         31,42,
                                         25,29,
                                         35,55,
                                         34,47,
                                         50,46,
                                         44,40,
                                         45,37,
                                         38,36,
                                         60,39,
                                         50,41,
                                         54,47
                                         )
                           )
machine_data %>% kable()
```

cutting_speed	tool_geometry	cutting_angle	lifetime
-	-	-	22
+	-	-	32
-	-	-	31
+	-	-	42
-	-	-	25
+	-	-	29
-	+	-	35
+	+	-	55
-	+	-	34
+	+	-	47
-	+	-	50
+	+	-	46
-	-	+	44
+	-	+	40
-	-	+	45
+	-	+	37
-	-	+	38
+	-	+	36
-	+	+	60
+	+	+	39
-	+	+	50
+	+	+	41
-	+	+	54
+	+	+	47

a

Estimate the factor effects. Which effect appears to be large?

```
lifetime_model <- aov(lifetime~cutting_speed*tool_geometry*cutting_angle,data=machine_data)
lifetime_model$coefficients
```

```
##                (Intercept)
##                26.000000
##          cutting_speed+
##                8.333333
##          tool_geometry+
##                13.666667
##          cutting_angle+
##                16.333333
## cutting_speed+:tool_geometry+
##                1.333333
## cutting_speed+:cutting_angle+
##               -13.000000
## tool_geometry+:cutting_angle+
##               -1.333333
## cutting_speed+:tool_geometry+:cutting_angle+
##               -9.000000
```

b

Use the analysis of variance to confirm your conclusions for part a.

```
anova(lifetime_model)
```

```
## Analysis of Variance Table
##
## Response: lifetime
##              Df Sum Sq Mean Sq F value
## cutting_speed    1   0.38    0.38  0.0129
## tool_geometry     1 782.04   782.04 26.8129
## cutting_angle     1 287.04   287.04  9.8414
## cutting_speed:tool_geometry    1  15.04    15.04  0.5157
## cutting_speed:cutting_angle    1 459.38   459.38 15.7500
## tool_geometry:cutting_angle    1  51.04    51.04  1.7500
## cutting_speed:tool_geometry:cutting_angle  1  30.37    30.37  1.0414
## Residuals       16 466.67    29.17
##              Pr(>F)
## cutting_speed    0.911132
## tool_geometry    9.154e-05 ***
## cutting_angle    0.006365 **
## cutting_speed:tool_geometry    0.483031
## cutting_speed:cutting_angle    0.001102 **
## tool_geometry:cutting_angle    0.204473
## cutting_speed:tool_geometry:cutting_angle  0.322673
## Residuals
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

c

Write down a regression model for predicting tool life (in hours) based on the results of this experiment.

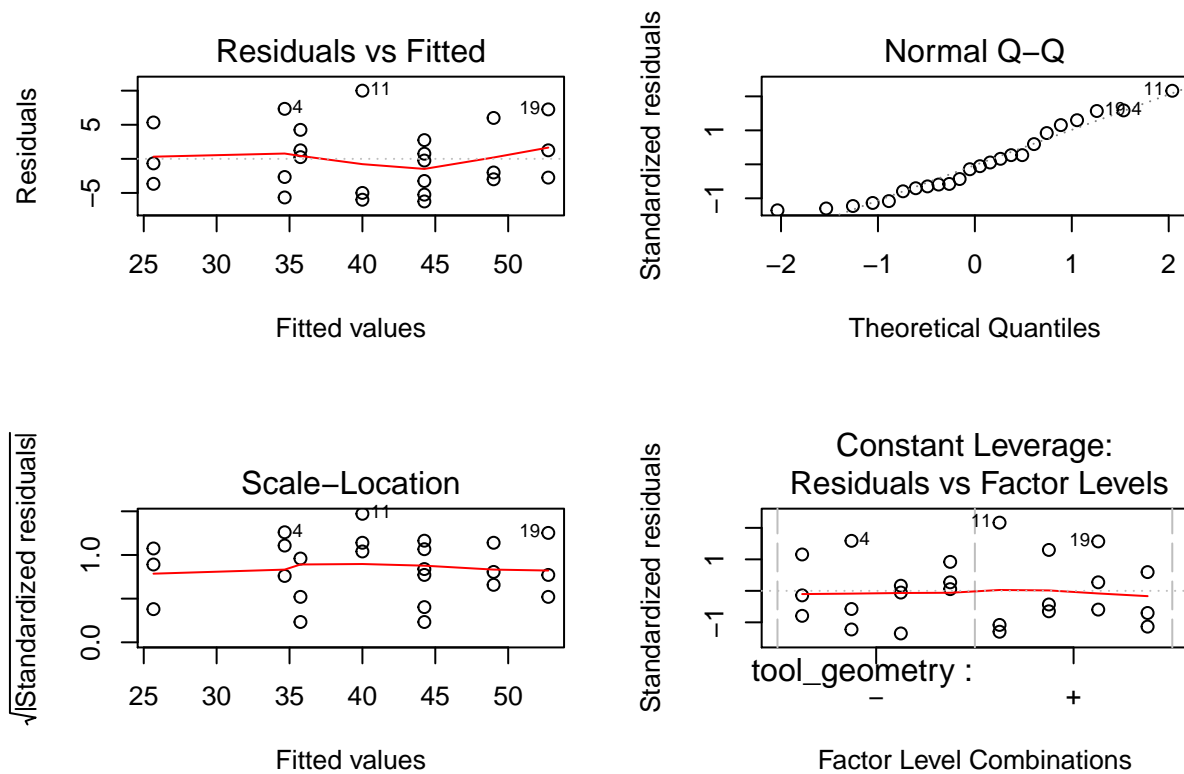
```
reduced_lifetime_model <- lm(lifetime~tool_geometry*cutting_angle+cutting_speed:cutting_angle,data=mach
summary(reduced_lifetime_model)
```

```
##
## Call:
## lm(formula = lifetime ~ tool_geometry * cutting_angle + cutting_speed:cutting_angle,
##     data = machine_data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -6.2500 -3.3542 -0.4583  3.1250 10.0000
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      25.667      2.667   9.624 1.6e-08 ***
## tool_geometry+      14.333      3.079   4.655 0.000197 ***
## cutting_angle+      18.583      3.772   4.927 0.000109 ***
## tool_geometry+:cutting_angle+ -5.833      4.355  -1.339 0.197091
## cutting_angle-:cutting_speed+  9.000      3.079   2.923 0.009090 **
## cutting_angle+:cutting_speed+ -8.500      3.079  -2.760 0.012888 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.334 on 18 degrees of freedom
## Multiple R-squared:  0.7552, Adjusted R-squared:  0.6872
## F-statistic: 11.11 on 5 and 18 DF,  p-value: 5.361e-05
```

d

Analyze the residuals. Are there any obvious problems?

```
par( mfrow = c(2,2) )
plot(reduced_lifetime_model)
```

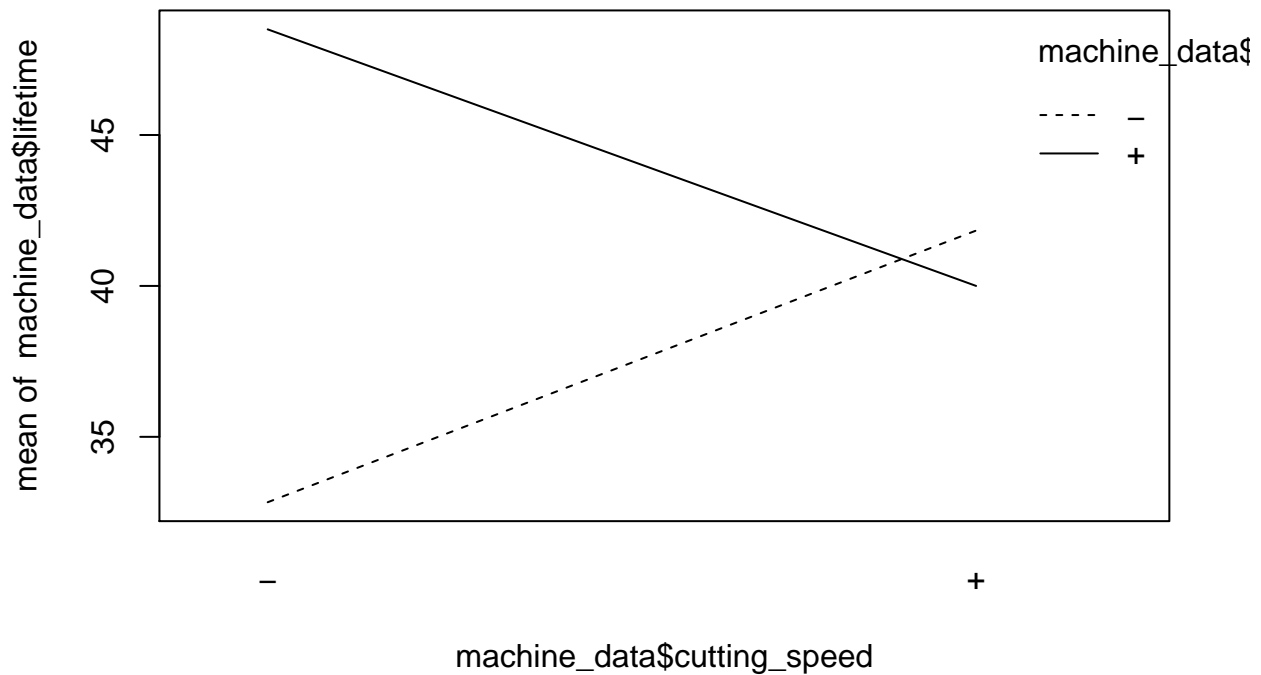


e

based on an analysis of main effect and interaction plots, what levels of A, B, and C would you recommend using?

```
interaction.plot(x.factor = machine_data$cutting_speed,
                 trace.factor = machine_data$cutting_angle,
                 response = machine_data$lifetime)
```





low cutting speed and high cutting angle is associated with the longest lifetime.

### 3

An experiment was performed to improve the yield of a chemical process. Four factors were selected, and two replicates of a completely randomized experiment were run. The results are shown in the following table.

```
lowhigh <- c("-", "+")
treatments <- expand.grid(factor1 = rep(lowhigh, 2),
                          factor2 = lowhigh,
                          factor3 = lowhigh,
                          factor4 = lowhigh
                        )
chem_data <- cbind(treatments,
                  yield = c(90, 74, 93, 78,
                           81, 83, 85, 80,
                           77, 81, 78, 80,
                           88, 73, 82, 70,
                           98, 72, 98, 76,
                           87, 85, 83, 86,
                           99, 79, 90, 75,
                           87, 80, 84, 80
                          )
                )
chem_data %>% kable()
```

factor1	factor2	factor3	factor4	yield
-	-	-	-	90
+	-	-	-	74
-	-	-	-	93
+	-	-	-	78
-	+	-	-	81
+	+	-	-	83
-	+	-	-	85
+	+	-	-	80
-	-	+	-	77
+	-	+	-	81
-	-	+	-	78
+	-	+	-	80
-	+	+	-	88
+	+	+	-	73
-	+	+	-	82
+	+	+	-	70
-	-	-	+	98
+	-	-	+	72
-	-	-	+	98
+	-	-	+	76
-	+	-	+	87
+	+	-	+	85
-	+	-	+	83
+	+	-	+	86
-	-	+	+	99
+	-	+	+	79
-	-	+	+	90
+	-	+	+	75
-	+	+	+	87
+	+	+	+	80
-	+	+	+	84
+	+	+	+	80

**a**

Estimate the factor effects.

```
chem_model <- aov(formula = yield ~ factor1*factor2*factor3*factor4,
                  data=chem_data)

chem_model$coefficients %>% kable()
```

	x
(Intercept)	91.5
factor1+	-15.5
factor2+	-8.5
factor3+	-14.0
factor4+	6.5
factor1+:factor2+	14.0
factor1+:factor3+	18.5
factor2+:factor3+	16.0

	x
factor1+:factor4+	-8.5
factor2+:factor4+	-4.5
factor3+:factor4+	10.5
factor1+:factor2+:factor3+	-30.5
factor1+:factor2+:factor4+	10.5
factor1+:factor3+:factor4+	-12.0
factor2+:factor3+:factor4+	-12.0
factor1+:factor2+:factor3+:factor4+	18.0

**b**

Prepare an analysis of variance table and determine which factors are important in explaining yield.

```
summary(chem_model)
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## factor1      1  684.5   684.5   92.814  4.6e-08 ***
## factor2      1   18.0    18.0    2.441  0.137785
## factor3      1   66.1    66.1    8.966  0.008580 **
## factor4      1  136.1   136.1   18.458  0.000555 ***
## factor1:factor2  1  144.5   144.5   19.593  0.000423 ***
## factor1:factor3  1    6.1     6.1    0.831  0.375652
## factor2:factor3  1    1.1     1.1    0.153  0.701269
## factor1:factor4  1   45.1    45.1    6.119  0.024964 *
## factor2:factor4  1    1.1     1.1    0.153  0.701269
## factor3:factor4  1   18.0    18.0    2.441  0.137785
## factor1:factor2:factor3  1  231.1   231.1   31.339  4.0e-05 ***
## factor1:factor2:factor4  1  190.1   190.1   25.780  0.000112 ***
## factor1:factor3:factor4  1    4.5     4.5    0.610  0.446129
## factor2:factor3:factor4  1    4.5     4.5    0.610  0.446129
## factor1:factor2:factor3:factor4  1   40.5    40.5    5.492  0.032355 *
## Residuals      16  118.0     7.4
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

**c**

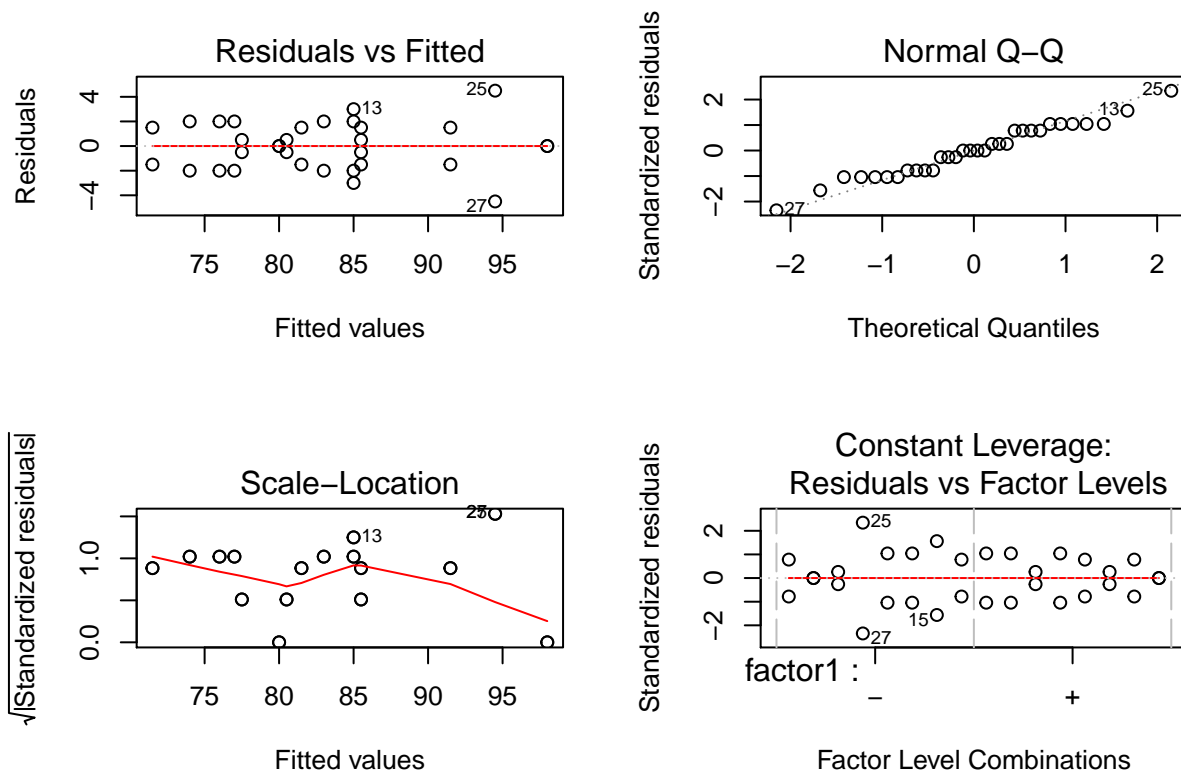
Write down a regression model for predicting yield, assuming that all four factors were varied over the range from -1 to +1.

$$yield = 91.5 - 15.5f_1 - 8.5f_2 - 14.0f_3 + 6.5f_4 + 14.0f_1f_2 - 30.5f_1f_2f_3 + 10.5f_1f_2f_4$$

**d**

Plot the residuals versus the predicted yield and on a normal probability scale. Does the residual analysis appear satisfactory?

```
par( mfrow = c(2,2) )
plot(chem_model)
```



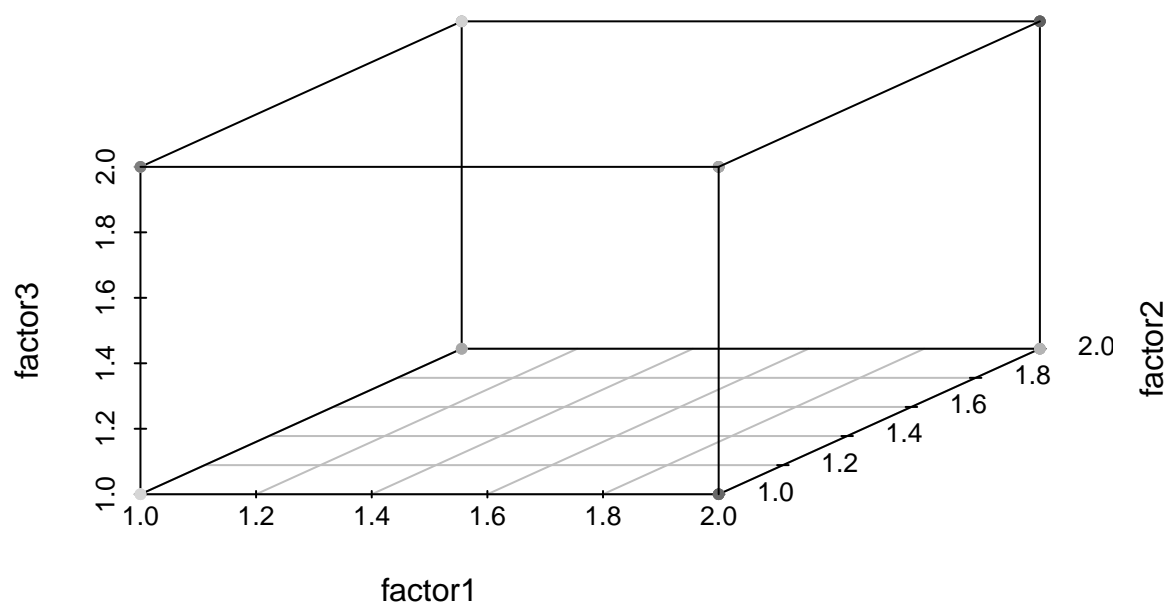
e

Two three-factor interactions, ABC and ABD, apparently have large effects. Draw a cube plot in the factors A, B, and C with the average yields shown at each corner. Repeat using the factors A, B, and D. Do these two plots aid in a data interpretation? Where would you recommend that the process be run with respect to the four variables?

```
colorpalette <- gray.colors(length(chem_data$yield))
colorpalette
```

```
## [1] "#4D4D4D" "#575757" "#606060" "#696969" "#707070" "#777777" "#7E7E7E"
## [8] "#848484" "#8A8A8A" "#8F8F8F" "#949494" "#999999" "#9E9E9E" "#A3A3A3"
## [15] "#A8A8A8" "#ACACAC" "#B0B0B0" "#B4B4B4" "#B8B8B8" "#BCBCBC" "#C0C0C0"
## [22] "#C4C4C4" "#C8C8C8" "#CBCBCB" "#CFCFCF" "#D2D2D2" "#D6D6D6" "#D9D9D9"
## [29] "#DCDCDC" "#DFDFDF" "#E2E2E2" "#E6E6E6"
```

```
colors <- colorpalette[rank(chem_data$yield)]
scatterplot3d(chem_data[1:3], pch=20, color = colors)
```



```
scatterplot3d(chem_data[c(1,2,4)],pch=20,color = colors)
```

