

01NAEX - Lecture 06

2^k Factorial Design

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2^k Factorial Design

Last lesson: Introduction to 2^k Factorial Design

- ▶ 2^k factorial design is widely used in industrial experimentation, especially in the early stages of experimental work, when many factors are likely to be investigated.
- ▶ Special case of the general factorial design with k factors, all at 2 levels, usually called low (-) and high (+).
- ▶ Factors are fixed, the design is completely randomized, and the usual normality assumptions are satisfied.

Today's lesson:

- ▶ Introduction to FrF2 package in R.
- ▶ Daniel Plot: Normal and Half-Normal Plot Effects.
- ▶ Pareto plot: Lenth's method.

2^k Factorial Design - Recap

General 2^k Factorial Design has

- ▶ k main effects
- ▶ $\binom{k}{2}$ two-factor interactions (first order interaction effects)
- ▶ $\binom{k}{3}$ three-factor interactions (second order interaction effects)
- ▶ \vdots
- ▶ 1 k -factor interaction

Unreplicated (single replicated) 2^k Factorial Design:

- ▶ If the factors are spaced too closely, it increases the chances that the noise will overwhelm the signal in the data.
- ▶ More aggressive spacing is usually best.
- ▶ Lack of replication causes potential problems in statistical testing:
 - ▶ With no replication, fitting the full model results in zero degrees of freedom for error.
 - ▶ Pooling high-order interactions to estimate error.
 - ▶ Normal probability plotting of effects (Daniels, 1959)

2³ Factorial Design in R - FrF2 package (Plasma Etch Rate Experiment)

Back to Plasma Etch Rate Experiment from the last lesson

```
> FrF2(2^3, 3, replications = 2, randomize = FALSE,  
      factor.names = c("Gap", "Flow", "Power"))
```

creating full factorial with 8 runs ...

run.no	run.no.std.rp	Gap	Flow	Power
1	1.1	-1	-1	-1
2	2.1	1	-1	-1
3	3.1	-1	1	-1
4	4.1	1	1	-1
5	5.1	-1	-1	1
6	6.1	1	-1	1
7	7.1	-1	1	1
8	8.1	1	1	1
9	1.2	-1	-1	-1
10	2.2	1	-1	-1
11	3.2	-1	1	-1
12	4.2	1	1	-1
13	5.2	-1	-1	1
14	6.2	1	-1	1
15	7.2	-1	1	1
16	8.2	1	1	1

```
class=design, type= full factorial
```

2³ Factorial Design in R - FrF2 package (Plasma Etch Rate Experiment)

Care about settings: randomize = TRUE or randomize = FALSE

```
> FrF2(2^3, 3, replications = 2, randomize = T,  
      factor.names = c("Gap", "Flow", "Power"))
```

creating full factorial with 8 runs ...

run.no	run.no.std.rp	Gap	Flow	Power
1	7.1	-1	1	1
2	6.1	1	-1	1
3	3.1	-1	1	-1
4	5.1	-1	-1	1
5	4.1	1	1	-1
6	8.1	1	1	1
7	2.1	1	-1	-1
8	1.1	-1	-1	-1
9	3.2	-1	1	-1
10	2.2	1	-1	-1
11	6.2	1	-1	1
12	5.2	-1	-1	1
13	8.2	1	1	1
14	7.2	-1	1	1
15	4.2	1	1	-1
16	1.2	-1	-1	-1

```
class=design, type= full factorial
```

2^k Factorial Design in R - FrF2 package

Back to Plasma Etch Rate Experiment

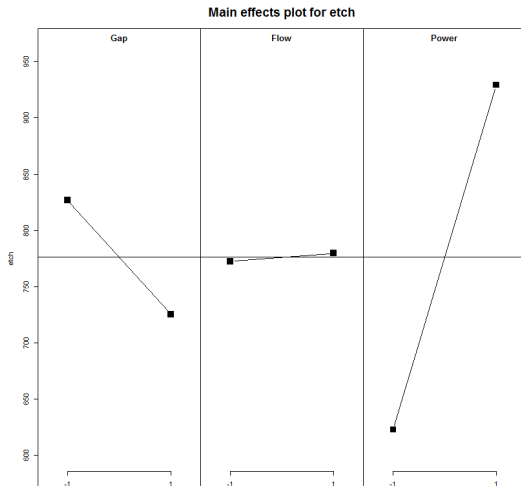
```
k = 3
plan <- FrF2(2^k, k, replications = 2, randomize = FALSE,
factor.names = c("Gap", "Flow", "Power"))
plan <- add.response(plan, etch)
> plan
```

Gap	Flow	Power	etch	Gap	Flow	Power	etch		
1	-1	-1	-1	550	9	-1	-1	-1	604
2	1	-1	-1	669	10	1	-1	-1	650
3	-1	1	-1	633	11	-1	1	-1	601
4	1	1	-1	642	12	1	1	-1	635
5	-1	-1	1	1037	13	-1	-1	1	1052
6	1	-1	1	749	14	1	-1	1	868
7	-1	1	1	1075	15	-1	1	1	1063
8	1	1	1	729	16	1	1	1	860

Analysis - same as in the last lesson.

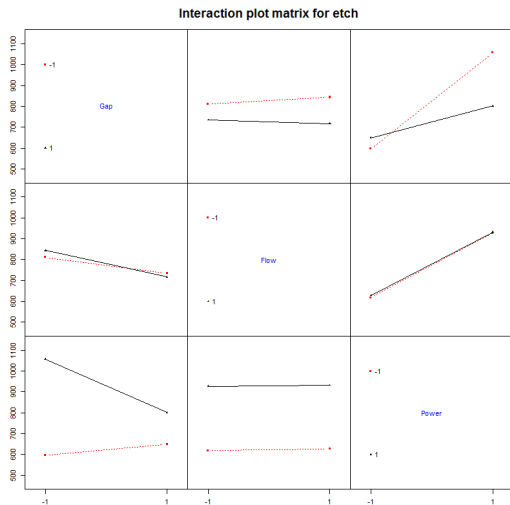
2^k Factorial Design in R - FrF2 package

MEPlot(plan)



2^k Factorial Design in R - FrF2 package

IAPlot(plan)



Unreplicated 2^k Factorial Designs

The 2^4 factorial design was used to investigate the effects of four factors on the filtration rate of a resin for a chemical process plant. The factors are:

A: temperature,

B: pressure,

C: concentration of chemical formaldehyde,

D: stirring rate.

Run Number	Factor				Run Label	Filtration Rate (gal/h)
	A	B	C	D		
1	–	–	–	–	(1)	45
2	+	–	–	–	<i>a</i>	71
3	–	+	–	–	<i>b</i>	48
4	+	+	–	–	<i>ab</i>	65
5	–	–	+	–	<i>c</i>	68
6	+	–	+	–	<i>ac</i>	60
7	–	+	+	–	<i>bc</i>	80
8	+	+	+	–	<i>abc</i>	65
9	–	–	–	+	<i>d</i>	43
10	+	–	–	+	<i>ad</i>	100
11	–	+	–	+	<i>bd</i>	45
12	+	+	–	+	<i>abd</i>	104
13	–	–	+	+	<i>cd</i>	75
14	+	–	+	+	<i>acd</i>	86
15	–	+	+	+	<i>bcd</i>	70
16	+	+	+	+	<i>abcd</i>	96

$2^4 = 16$ runs were made in random order.

Unreplicated 2^k Factorial Designs

Pilot Plant Filtration Rate Experiment - FrF2 package

```
> rate=FrF2(2^4, 4, replications = 1,  
            randomize = FALSE, factor.names = c("A", "B", "C", "D"))  
> Filtration = rate_experiment$Rate  
> rate      = add.response(rate, Filtration)  
> summary(rate)
```

Experimental design of type full factorial 16 runs

Factor settings: Responses:

A B C D [1] Filtration

1 -1 -1 -1 -1

2 1 1 1 1

The design itself:

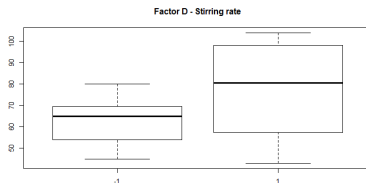
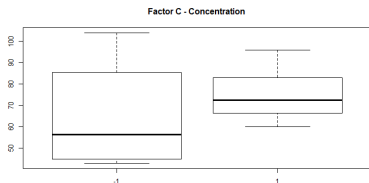
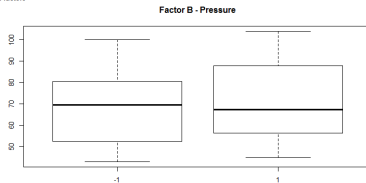
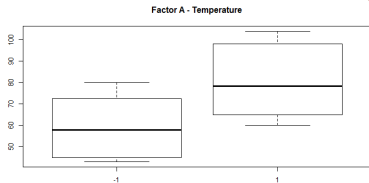
	A	B	C	D	Filtration	A	B	C	D	Filtration
1	-1	-1	-1	-1	45	-1	-1	-1	1	43
2	1	-1	-1	-1	71	1	-1	-1	1	100
3	-1	1	-1	-1	48	-1	1	-1	1	45
4	1	1	-1	-1	65	1	1	-1	1	104
5	-1	-1	1	-1	68	-1	-1	1	1	75
6	1	-1	1	-1	60	1	-1	1	1	86
7	-1	1	1	-1	80	-1	1	1	1	70
8	1	1	1	-1	65	1	1	1	1	96

Pilot Plant Filtration Rate Experiment

Box plot

```
boxplot(Filtration ~ A , main = "Factor A - Temperature")
```

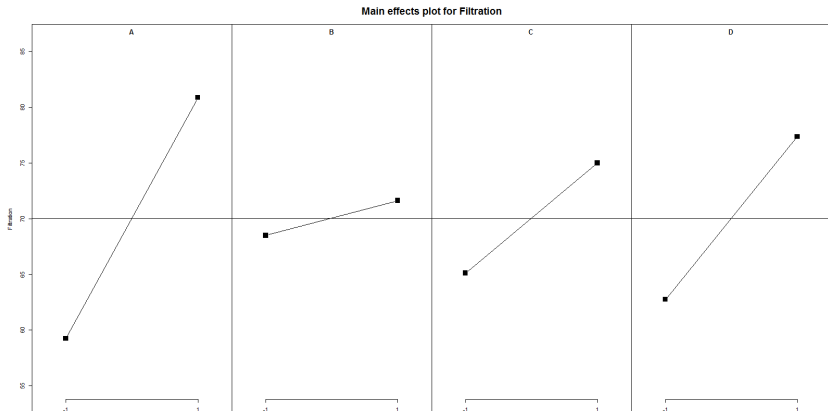
Box plots of all factors



Pilot Plant Filtration Rate Experiment

Main Effects plot for response variable

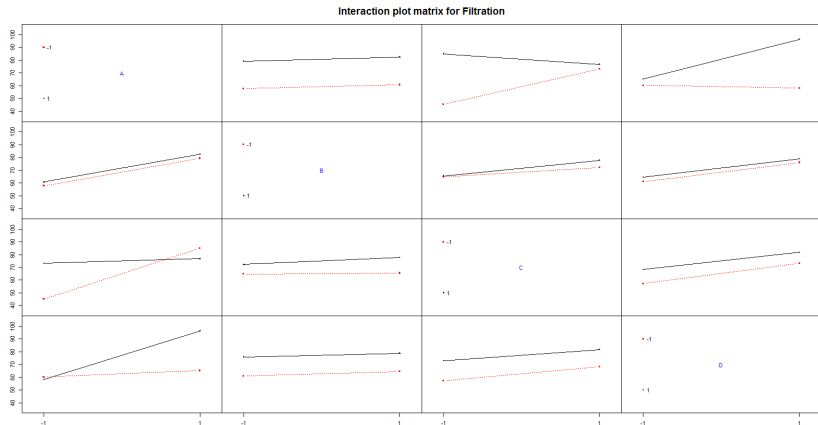
MEPlot(plan)



Pilot Plant Filtration Rate Experiment

Interaction Plot matrix for response variable

IAPlot(plan)



Pilot Plant Filtration Rate Experiment

ANOVA table - model with all factors and interactions

```
anova(aov(Filtration~A*B*C*D, data=rate))
```

Analysis of Variance Table, Response: Filtration

		Df	Sum Sq	Mean Sq	F value	Pr(>F)
A		1	1870.56	1870.56		
B		1	39.06	39.06		
C		1	390.06	390.06		
D		1	855.56	855.56		
A:B		1	0.06	0.06		
A:C		1	1314.06	1314.06		
B:C		1	22.56	22.56		
A:D		1	1105.56	1105.56		
B:D		1	0.56	0.56		
C:D		1	5.06	5.06		
A:B:C		1	14.06	14.06		
A:B:D		1	68.06	68.06		
A:C:D		1	10.56	10.56		
B:C:D		1	27.56	27.56		
A:B:C:D		1	7.56	7.56		
Residuals		0	0.00			

Warning message: ANOVA F-tests are unreliable

Pilot Plant Filtration Rate Experiment: LM - all factors and interactions

```
> summary(lm(Filtration~A*B*C*D, data=rate))
```

ALL 16 residuals are 0: no residual degrees of freedom!

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	70.0625	NA	NA	NA
A1	10.8125	NA	NA	NA
B1	1.5625	NA	NA	NA
C1	4.9375	NA	NA	NA
D1	7.3125	NA	NA	NA
A1:B1	0.0625	NA	NA	NA
A1:C1	-9.0625	NA	NA	NA
B1:C1	1.1875	NA	NA	NA
A1:D1	8.3125	NA	NA	NA
B1:D1	-0.1875	NA	NA	NA
C1:D1	-0.5625	NA	NA	NA
A1:B1:C1	0.9375	NA	NA	NA
A1:B1:D1	2.0625	NA	NA	NA
A1:C1:D1	-0.8125	NA	NA	NA
B1:C1:D1	-1.3125	NA	NA	NA
A1:B1:C1:D1	0.6875	NA	NA	NA

Residual standard error: NaN on 0 degrees of freedom

Multiple R-squared: 1, Adjusted R-squared: NaN

F-statistic: NaN on 15 and 0 DF, p-value: NA

Analysis of significant effects

We can't use usual way because:

- ▶ ANOVA gives SS but no error estimate!
- ▶ LM gives effect estimates but no error estimate!
- ▶ We try to adjust a 2^k dimensional model to 2^k observations, leaving 0 dimensions (degrees of freedom) for the error estimate.

We don't know σ and we can't estimate it. We know that

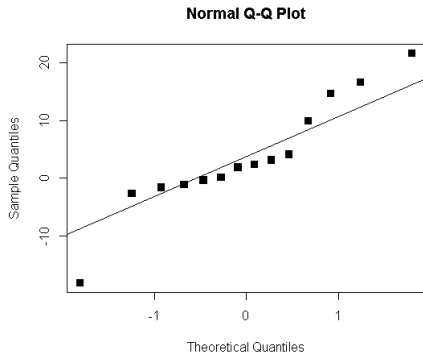
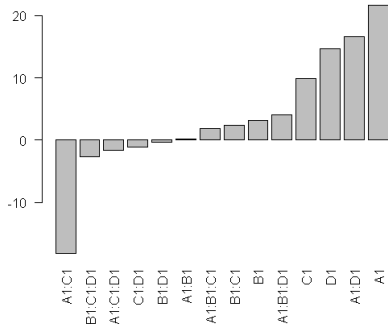
- ▶ the effects estimates are normally distributed with mean 0 and common standard deviation σ .
- ▶ the effect estimates are independent.

Lets make *QQ*-plot and look for the outliers - suspected significant effects.

Analysis of significant effects

Make QQ plot from Linear Model coefficients (without intercept) manually:

```
> model0 = lm(2*Filtration~A*B*C*D, data=rate)
> barplot(sort(model0$coeff[2:(2^4-1)]), las = 2)
> qqnorm(model0$coeff[2:(2^4-1)], cex = 1.3, pch = 15)
> qqline(model0$coeff[2:(2^4-1)], cex = 1.3, pch = 15)
```



Analysis of significant effects

In unreplicated 2^k factorial design with high-order interactions we can use R build-in functions:

- ▶ Daniel normal probability plot;
- ▶ The Half-Normal Plot Effects;
- ▶ Lenth's method - Pareto plot.

Required packages in R: FrF2, qualityTools

Daniel plot

Daniel plot is the normal probability plot of the estimates of the effects.

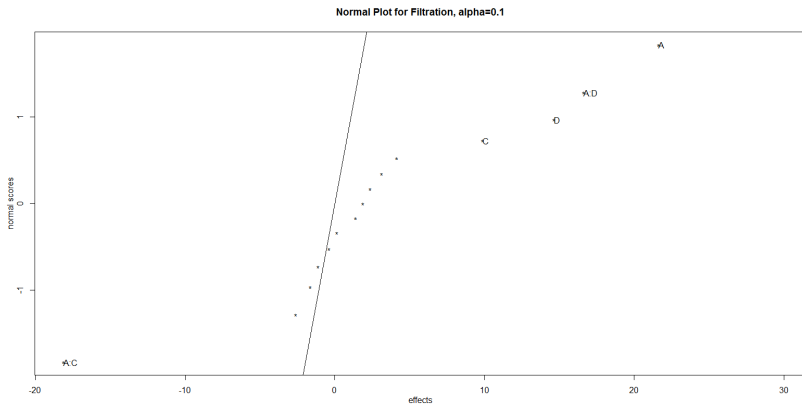
The effects that are negligible are normally distributed, with mean zero and variance σ^2 and will tend to fall along a straight line on this plot, whereas significant effects will have nonzero means and will not lie along straight line.

1. **Negligible effects:** lie along the qqline.
2. **Important effects:** lie far from the qqline.

Pilot Plant Filtration Rate Experiment

Daniel Plot (Classical effects qqplot) with $\alpha = 0.1$, qqline, and only significant factors

```
qqplot(DanielPlot(rate, alpha=0.1)$x, DanielPlot(rate)$y)  
qqline(DanielPlot(rate, alpha=0.1)$y)
```



The Half-Normal Plot Effects

Alternative plot to the Normal probability plot:

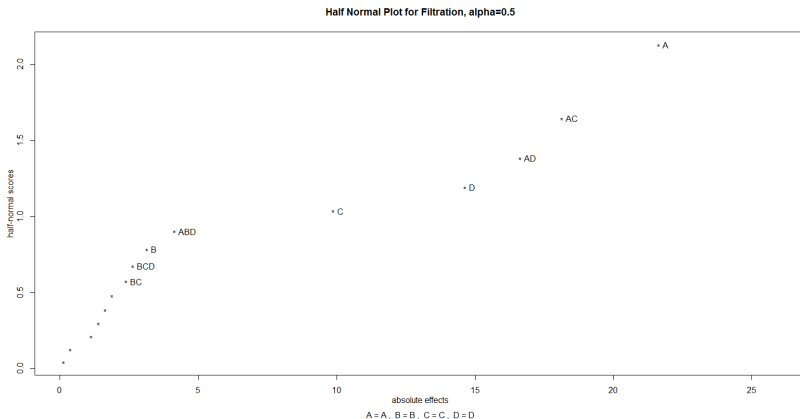
It is a plot of absolute value of the effects estimates against their cumulative normal probabilities.

The straight line on the half-normal plot always passes through the origin and should pass close to the fiftieth percentile data value. Better for interpretation with a few effects estimates (for example 8-run design).

Pilot Plant Filtration Rate Experiment

Half normal plot of effects

```
DanielPlot(rate, code=TRUE, alpha=0.1, half=TRUE)
```



Lenth's method - Pareto plot

Method proposed by Lenth (1989), sometime called Pareto plot has good power to detect significant effects.

Suppose we have m contrast of interest c_1, c_2, \dots, c_m . For 2^k unreplicated factor design $m = 2^k - 1$. Lenth's method estimates the variance of a contrast from the smallest contrast estimate.

$$s_0 = 1.5 \text{median}(|c_j|) \quad PSE = 1.5 \text{median}(|c_j|, |c_j| < 2.5s_0)$$

PSE is called pseudostandard error and it should be reasonable estimator of the contrast variance. An individual contrast is compared to the **margin of error**:

$$ME = t_{0.025,d} PSE, \quad d = \frac{m}{3}$$

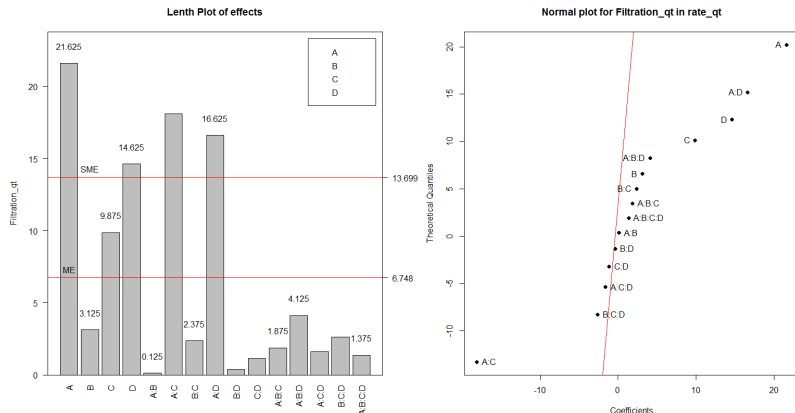
or to the **simultaneous margin of error**

$$SME = t_{\gamma,d} PSE, \quad \gamma = 1 - \frac{(1 + 0.95^{\frac{1}{m}})}{2}$$

Pilot Plant Filtration Rate Experiment - qualityTools library

```
rate_pareto = fracDesign(k = 4, replicates = 1)
response(rate_pareto) = rate$Filtration[rate_pareto[,1]]
paretoPlot(rate_pareto) normalPlot(rate_pareto)
```

$PSE = 1.5 \times |1.75| = 2.625$, $ME = 2.571 \times 2.625 = 6.748$, $SME = 5.219 \times 2.625 = 13.699$



Pilot Plant Filtration Rate Experiment

ANOVA table - model without factor B

By having dropped B totally, we obtain a 2^3 design with 2 replicates per cell.

```
> anova(aov(Filtration~A*C*D, data=rate))
```

Analysis of Variance Table

Response: Filtration

Df	Sum Sq	Mean Sq	F value	Pr(>F)
A	1870.56	1870.56	83.3677	1.667e-05 ***
C	390.06	390.06	17.3844	0.0031244 **
D	855.56	855.56	38.1309	0.0002666 ***
A:C	1314.06	1314.06	58.5655	6.001e-05 ***
A:D	1105.56	1105.56	49.2730	0.0001105 ***
C:D	5.06	5.06	0.2256	0.6474830
A:C:D	10.56	10.56	0.4708	0.5120321
Residuals	8	179.50	22.44	

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

>

Pilot Plant Filtration Rate Experiment

ANOVA table - final model

```
> anova(aov(Filtration~A*C+A*D, data=rate))
```

Analysis of Variance Table

Response: Filtration

Df	Sum Sq	Mean Sq	F value	Pr(>F)
A	1	1870.56	1870.56	95.865 1.928e-06 ***
C	1	390.06	390.06	19.990 0.001195 **
D	1	855.56	855.56	43.847 5.915e-05 ***
A:C	1	1314.06	1314.06	67.345 9.414e-06 ***
A:D	1	1105.56	1105.56	56.659 1.999e-05 ***
Residuals	10	195.12	19.51	

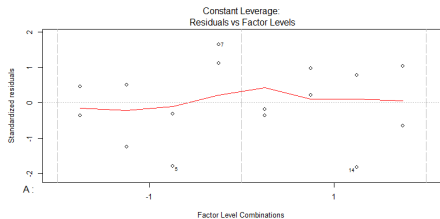
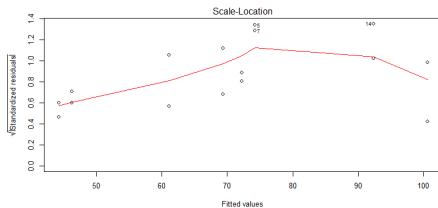
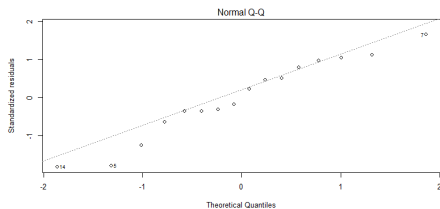
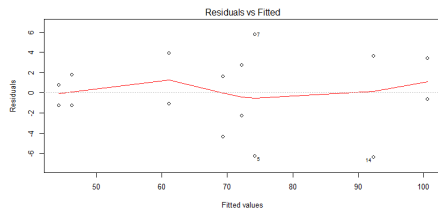
--

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

>

Pilot Plant Filtration Rate Experiment

Model validation



+ perform all relevant statistical hypothesis testing.

Pilot Plant Filtration Rate Experiment

Another approach - omitting the highest interaction

```
>anova(aov(Filtration~(.)^3, data=rate))
Analysis of Variance Table, Response: Filtration
Df  Sum Sq Mean Sq  F value    Pr(>F)
A      1 1870.56  1870.56  247.3471 0.04042 *
B      1   39.06   39.06   5.1653 0.26388
C      1  390.06  390.06  51.5785 0.08808 .
D      1  855.56  855.56 113.1322 0.05968 .
A:B    1    0.06    0.06   0.0083 0.94228
A:C    1 1314.06 1314.06 173.7603 0.04820 *
A:D    1 1105.56 1105.56 146.1901 0.05253 .
B:C    1   22.56   22.56   2.9835 0.33410
B:D    1    0.56    0.56   0.0744 0.83050
C:D    1    5.06    5.06   0.6694 0.56345
A:B:C  1   14.06   14.06   1.8595 0.40282
A:B:D  1   68.06   68.06   9.0000 0.20483
A:C:D  1   10.56   10.56   1.3967 0.44707
B:C:D  1   27.56   27.56   3.6446 0.30718
Residuals 1    7.56    7.56
```

Continue only with significant variables.

Pilot Plant Filtration Rate Experiment - Regression analysis

Preparation for contour plots (Swap factors for numerics)

```
> rate$A.num <- 2*(as.numeric(rate$A)-1.5)
> rate$C.num <- 2*(as.numeric(rate$C)-1.5)
> rate$D.num <- 2*(as.numeric(rate$D)-1.5)
> rate.lm      <- lm(Filtration~A.num*C.num+A.num*D.num,data=rate)
> summary(rate.lm)
```

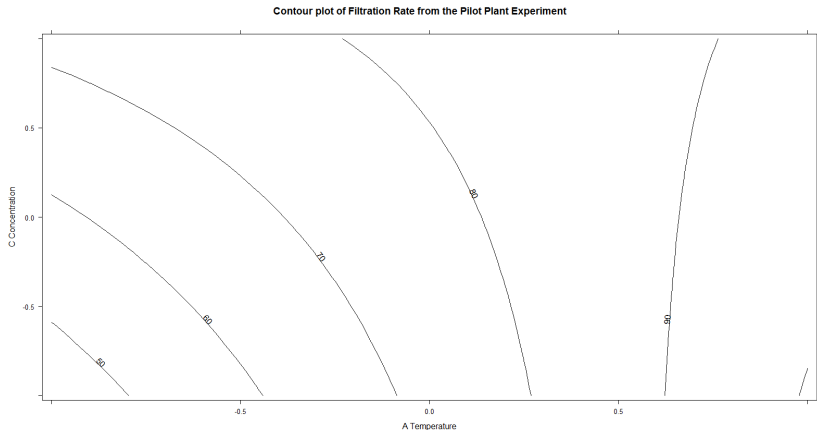
```
Coefficients: Estimate Std. Error t value Pr(>|t|)
(Intercept)    70.062      1.104   63.444 2.30e-14 ***
A.num           10.812      1.104    9.791 1.93e-06 ***
C.num           4.938      1.104    4.471 0.0012 **
D.num           7.313      1.104    6.622 5.92e-05 ***
A.num:C.num     -9.063      1.104   -8.206 9.41e-06 ***
A.num:D.num      8.312      1.104    7.527 2.00e-05 ***
Residual standard error: 4.417 on 10 degrees of freedom
Multiple R-squared: 0.966, Adjusted R-squared: 0.9489
F-statistic: 56.74 on 5 and 10 DF, p-value: 5.14e-07
```

```
> tmp      <- list(A.num=seq(-1,1,by=.05),C.num=seq(-1,1,by=0.05),
D.num=seq(-1,1,by=0.05),data=rate)
> new.data <- expand.grid(tmp)
> new.data$fit<- predict(rate.lm,new.data)
```

Pilot Plant Filtration Rate Experiment

Countour plot: A(temperature) and C(concentration) interaction only

```
contourplot (fit~A.num*C.num,new.data,xlab="A Temperature",  
ylab="C Concentration",  
main="Contour plot of Filtration Rate from the Pilot Plant Experiment ")
```



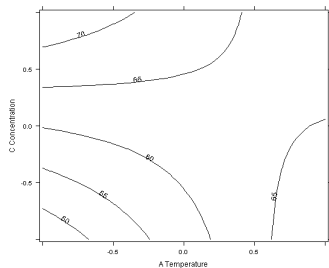
No specification in the code means last value of D, i.e. $D = 1$.

Pilot Plant Filtration Rate Experiment

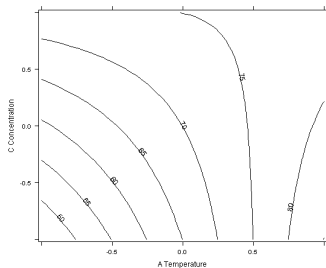
Countour plot: A(temperature) and C(concentration) for different D's.

```
contourplot (fit~A.num*C.num,new.data[new.data$D.num == -1,])
```

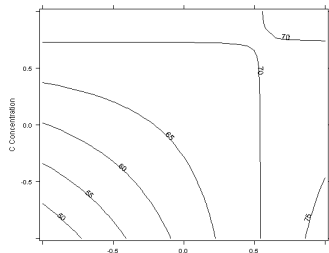
Contour plot of Filtration Rate from the Pilot Plant Experiment: D = -1



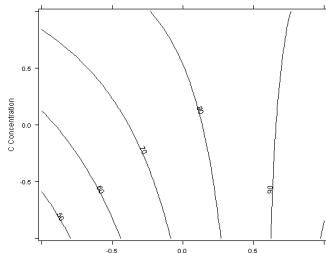
Contour plot of Filtration Rate from the Pilot Plant Experiment: D = 0



Contour plot of Filtration Rate from the Pilot Plant Experiment: D = -0.5



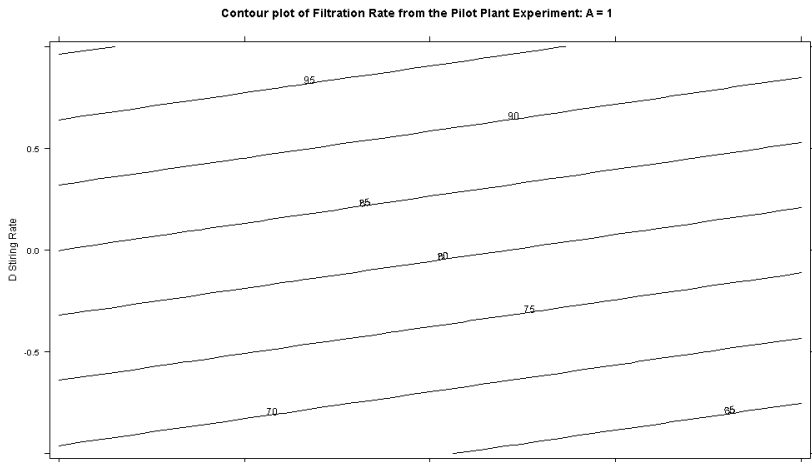
Contour plot of Filtration Rate from the Pilot Plant Experiment: D = 1



Pilot Plant Filtration Rate Experiment

Contour plot: C(concentration) and D(stirring rate) interaction only

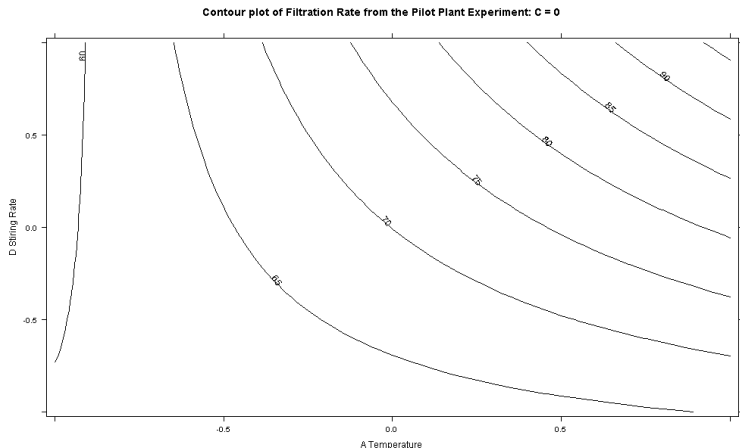
```
> contourplot(fit~C.num*D.num,new.data[new.data$A.num == 1,],  
  xlab="C Concentration",ylab="D Stiring Rate",  
  main = "Contour plot of Filtration Rate from the Pilot Plant Experiment  
  A = 1")
```



Pilot Plant Filtration Rate Experiment

Countour plot: A(temperature) and D(stirring rate) interaction only

```
contourplot(fit~A.num*D.num,new.data[new.data$C.num == 0,],  
xlab="A Temperature",ylab="D Stiring Rate",  
main="Contour plot of Filtration Rate from the Pilot Plant Experiment  
:C = 0")
```

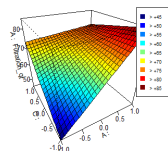


Pilot Plant Filtration Rate Experiment - qualityTools library

Countour and 3D plot

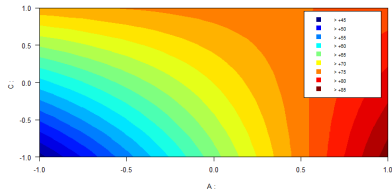
```
wirePlot(A, C, Filtration_qt, data = rate_qt)
contourPlot(A, C, Filtration_qt, data = rate_qt)
```

Response Surface for Filtration_qt

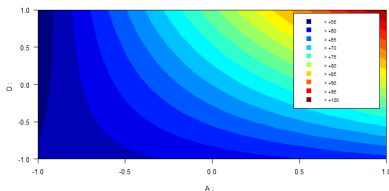


$$\text{Filtration_qt} = A * C + A * D$$

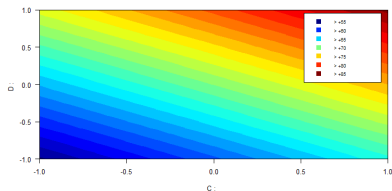
Filled Contour for Filtration_qt



Filled Contour for Filtration_qt



Filled Contour for Filtration_qt



Missing factor is equal to 0 in all three cases!

Today Exercise

Solve problems 6.31 and 6.32 from the chapter 6,
D. C. Montgomery DAE - 8. edition.

6.31. An experiment was conducted on a chemical process that produces a polymer. The four factors studied were temperature (*A*), catalyst concentration (*B*), time (*C*), and pressure (*D*). Two responses, molecular weight and viscosity, were observed. The design matrix and response data are shown in Table P6.8.

- (a) Consider only the molecular weight response. Plot the effect estimates on a normal probability scale. What effects appear important?
- (b) Use an analysis of variance to confirm the results from part (a). Is there indication of curvature?
- (c) Write down a regression model to predict molecular weight as a function of the important variables.
- (d) Analyze the residuals and comment on model adequacy.
- (e) Repeat parts (a)–(d) using the viscosity response.

6.32. Continuation of Problem 6.31. Use the regression models for molecular weight and viscosity to answer the following questions.

- (a) Construct a response surface contour plot for molecular weight. In what direction would you adjust the process variables to increase molecular weight?
- (b) Construct a response surface contour plot for viscosity. In what direction would you adjust the process variables to decrease viscosity?
- (c) What operating conditions would you recommend if it was necessary to produce a product with molecular weight between 2400 and 2500 and the lowest possible viscosity?

Don't use center points, skip the question about curvature term (will be covered in the next lesson).

Today Exercise

■ TABLE P6.8

The 2⁴ Experiment for Problem 6.31

Run Number	Actual Run Order	<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	Molecular Weight	Viscosity	Factor Levels		
								Low (–)	High (+)	
1	18	–	–	–	–	2400	1400	<i>A</i> (°C)	100	120
2	9	+	–	–	–	2410	1500	<i>B</i> (%)	4	8
3	13	–	+	–	–	2315	1520	<i>C</i> (min)	20	30
4	8	+	+	–	–	2510	1630	<i>D</i> (psi)	60	75
5	3	–	–	+	–	2615	1380			
6	11	+	–	+	–	2625	1525			
7	14	–	+	+	–	2400	1500			
8	17	+	+	+	–	2750	1620			
9	6	–	–	–	+	2400	1400			
10	7	+	–	–	+	2390	1525			
11	2	–	+	–	+	2300	1500			
12	10	+	+	–	+	2520	1500			
13	4	–	–	+	+	2625	1420			
14	19	+	–	+	+	2630	1490			
15	15	–	+	+	+	2500	1500			
16	20	+	+	+	+	2710	1600			
17	1	0	0	0	0	2515	1500			
18	5	0	0	0	0	2500	1460			
19	16	0	0	0	0	2400	1525			
20	12	0	0	0	0	2475	1500			