

01NAEX - Lecture 06

2^k Factorial Design

Exploring Experimental Designs for Multiple Factors

Jiri Franc

Czech Technical University
Faculty of Nuclear Sciences and Physical Engineering
Department of Mathematics

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:

- ▶ pyDOE3 in Python, FrF2 package in R.
- ▶ Daniel Plot: Normal and Half-Normal Plot Effects.
- ▶ Pareto plot: Lenth's method.

General 2^k Factorials (Overview & Notation)

Design: k two-level factors $\Rightarrow N = 2^k$ runs.

Levels coding: $-1, +1$.

Regression Model (no replication):

$$y = \beta_0 + \sum_i \beta_i x_i + \sum_{i < j} \beta_{ij} x_i x_j + \cdots + \beta_{12\dots k} x_1 \cdots x_k + \varepsilon.$$

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)
- ▶ :
- ▶ 1 k -factor interaction

2^k Factorial Design - (Overview & Notation)

Effect estimates (orthogonal coding):

- ▶ Contrast for effect E : $C_E = \sum_{\text{runs}} s_r y_r$, where $s_r \in \{-1, +1\}$ is the column for E .
- ▶ Estimated effect: $\hat{E} = C_E / 2^{k-1}$.
- ▶ Sum of Squares: $\text{SS}_E = C_E^2 / 2^k$.

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 tiny effects (high-order interactions) to estimate error.
 - ▶ σ^2 can be estimated by Daniel/half-normal/Lenth.

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

Back to Plasma Etch Rate Experiment from the last lesson - R

```
> 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

2^3 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

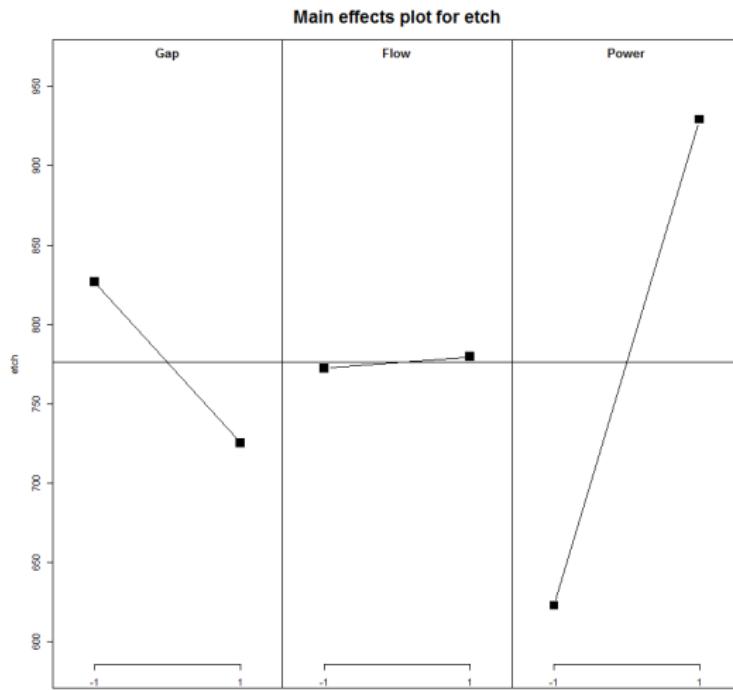
Back to Plasma Etch Rate Experiment - R

```
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.

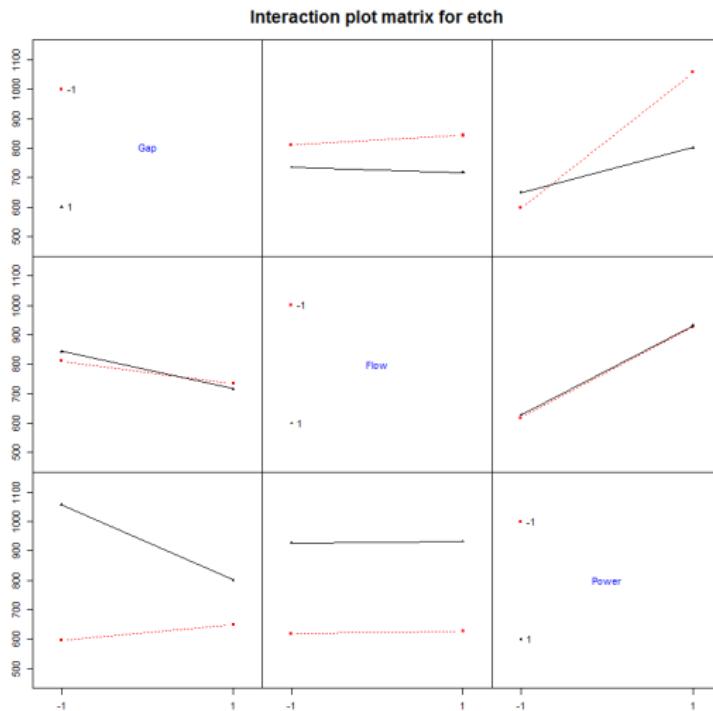
2^k Factorial Design in R - FrF2 package

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	+	-	-	-	a	71
3	-	+	-	-	b	48
4	+	+	-	-	ab	65
5	-	-	+	-	c	68
6	+	-	+	-	ac	60
7	-	+	+	-	bc	80
8	+	+	+	-	abc	65
9	-	-	-	+	d	43
10	+	-	-	+	ad	100
11	-	+	-	+	bd	45
12	+	+	-	+	abd	104
13	-	-	+	+	cd	75
14	+	-	+	+	acd	86
15	-	+	+	+	bcd	70
16	+	+	+	+	abcd	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		Responses:
1	-1	-1	-1	-1	[1] Filtration
2	1	1	1	1	

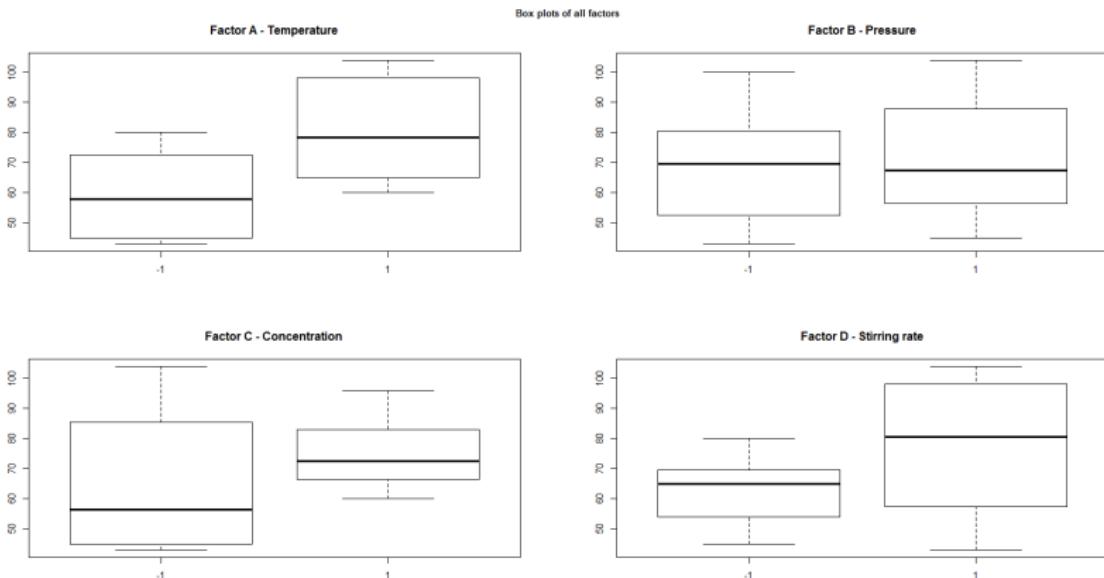
The design itself:

	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

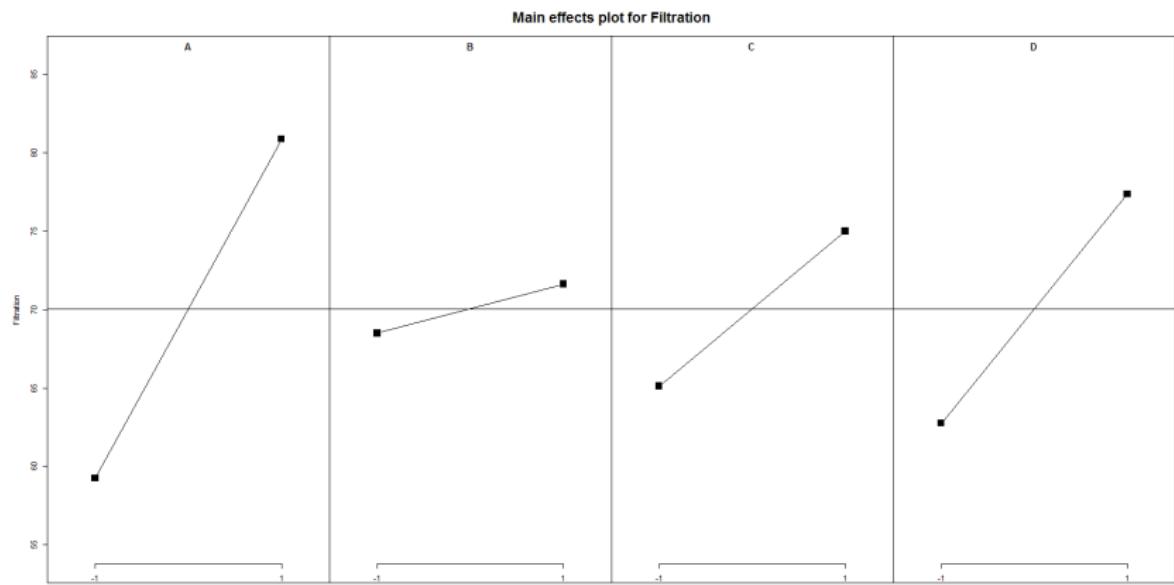
```
boxplot(Filtration ~ A , main = "Factor A - Temperature")
```



Pilot Plant Filtration Rate Experiment

Main Effects plot for response variable

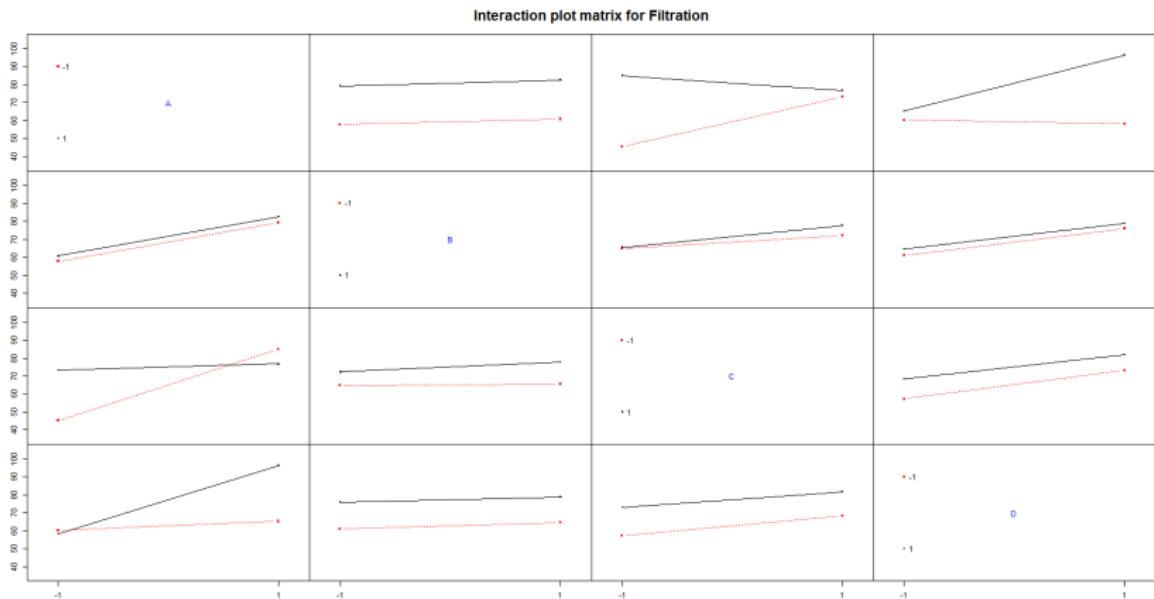
MEplot (plan)



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

Pilot Plant Filtration Rate Experiment

ANOVA table - model with all factors and interactions in Python

```
import statsmodels.api as sm
from statsmodels.formula.api import ols

# Define the model
model=ols('EtchRate ~ Gap*Flow*Power', data=design_df).fit()

# Perform ANOVA
anova_table = sm.stats.anova_lm(model, typ=2)
print(anova_table)
```

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

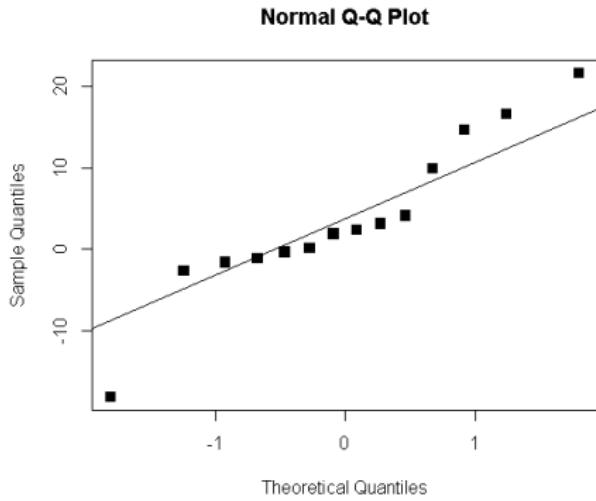
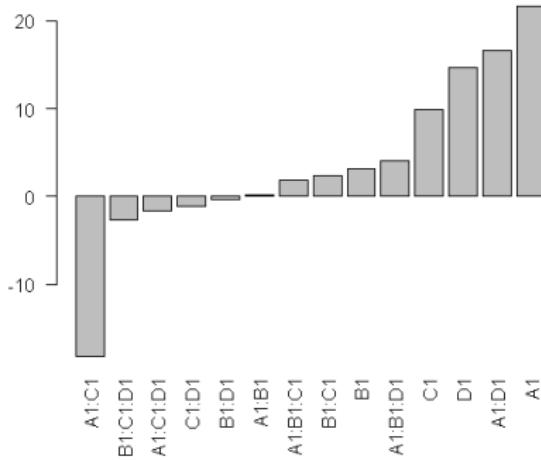
- ▶ the effects estimates are normally distributed with mean 0 and common standard deviation σ .
- ▶ the effect estimates are independent.

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)
```



Daniel plot

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

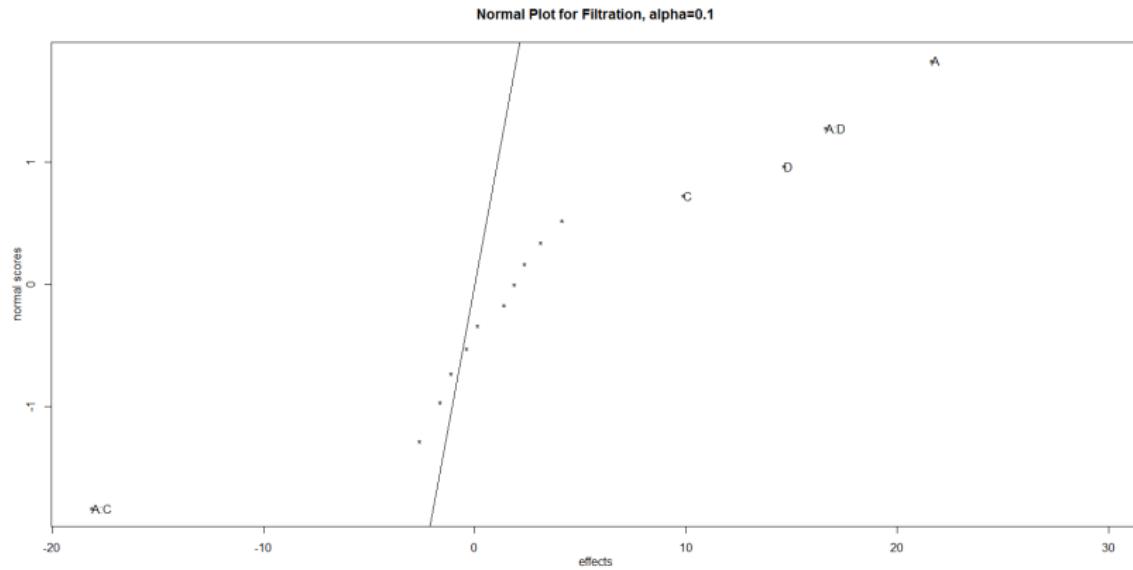
The effects that are negligible are normally distributed, with mean zero and variance σ^2 and will tend to fall along a straight line on this plot, whereas significant effects will have nonzero means and will not lie along straight line.

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:

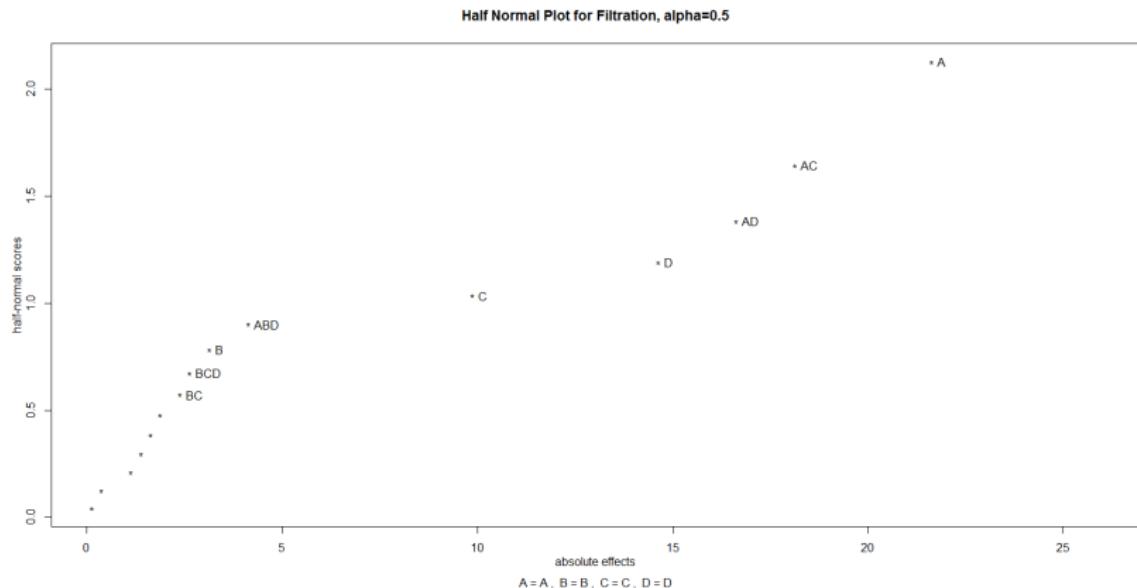
It is a plot of absolute value of the effects estimates against their cumulative normal probabilities.

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**:

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

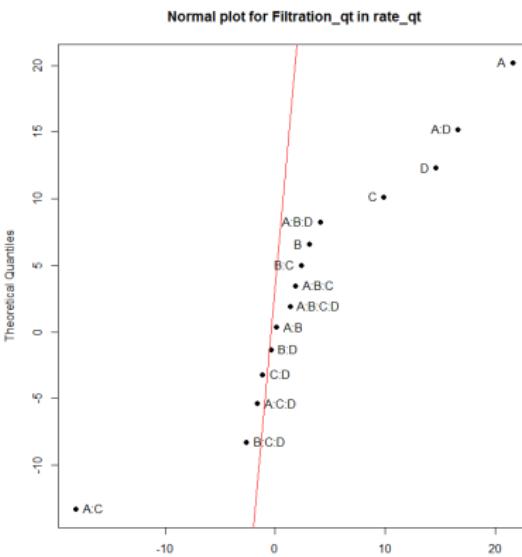
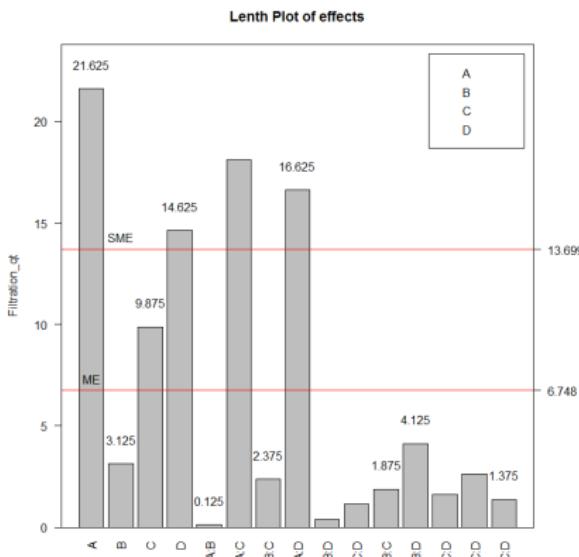
or to the **simultaneous margin of error**

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

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)
```

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



Daniel vs Half-Normal vs Lenth (Summary When No Replication)

Daniel (normal QQ): plot effect estimates \hat{E} vs normal quantiles. Active effects depart from the line.

Half-normal: plot $|\hat{E}|$ vs half-normal quantiles; no sign ambiguity.

Lenth PSE & Pareto: robust pseudo- σ from small effects,

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

then thresholds via

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

Build a Pareto of $|c_j|$ with ME/SME lines.

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	1	1870.56	1870.56	83.3677	1.667e-05	***
C	1	390.06	390.06	17.3844	0.0031244	**
D	1	855.56	855.56	38.1309	0.0002666	***
A:C	1	1314.06	1314.06	58.5655	6.001e-05	***
A:D	1	1105.56	1105.56	49.2730	0.0001105	***
C:D	1	5.06	5.06	0.2256	0.6474830	
A:C:D	1	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

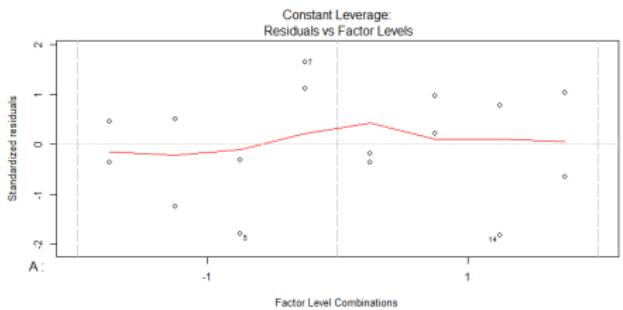
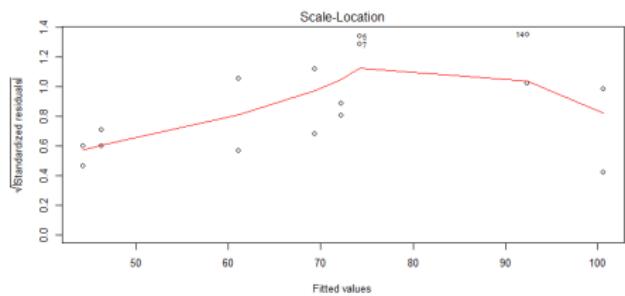
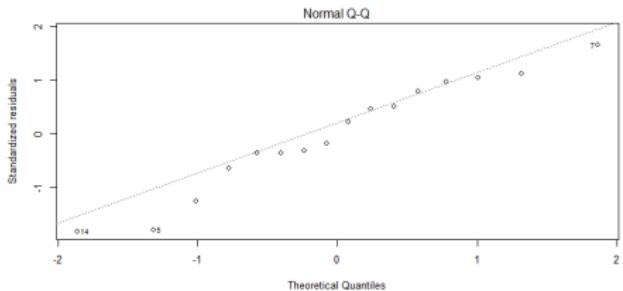
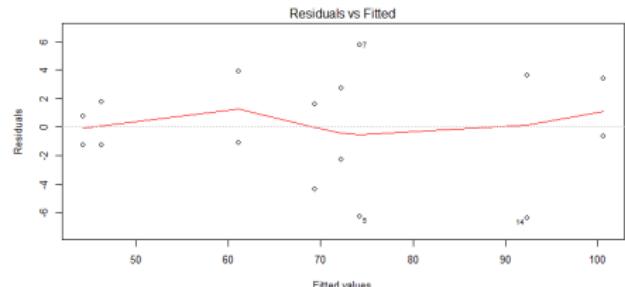
ANOVA table - final model

```
> 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)
```

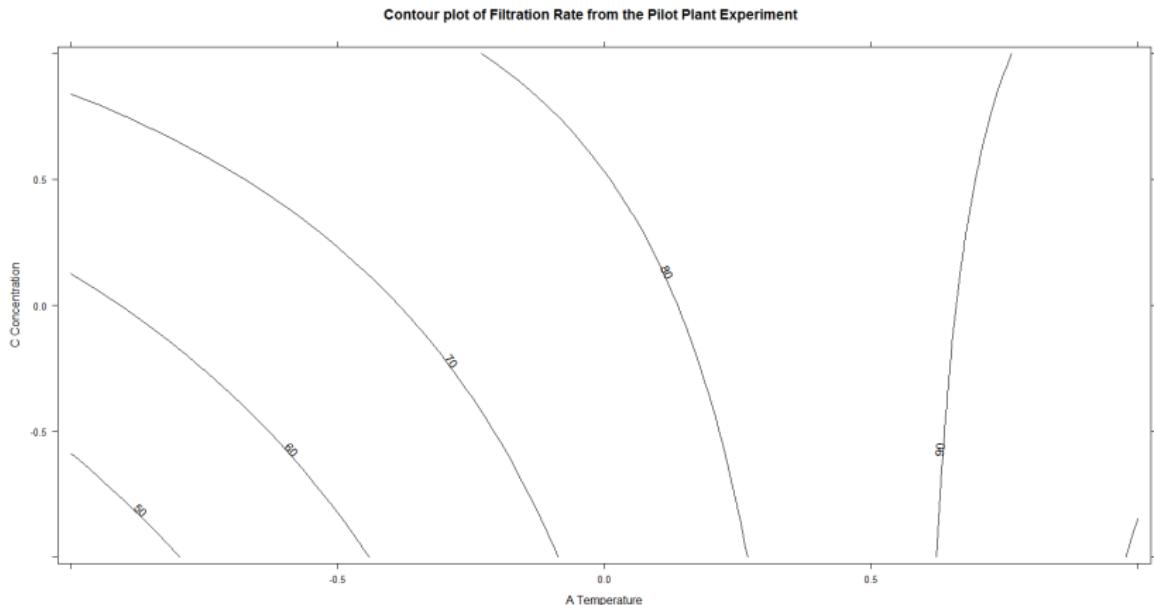
	Coefficients:	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	70.062	1.104	63.444	2.30e-14	***
A.num	10.812	1.104	9.791	1.93e-06	***
C.num	4.938	1.104	4.471	0.0012	**
D.num	7.313	1.104	6.622	5.92e-05	***
A.num:C.num	-9.063	1.104	-8.206	9.41e-06	***
A.num:D.num	8.312	1.104	7.527	2.00e-05	***
Residual 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

Contour 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")
```



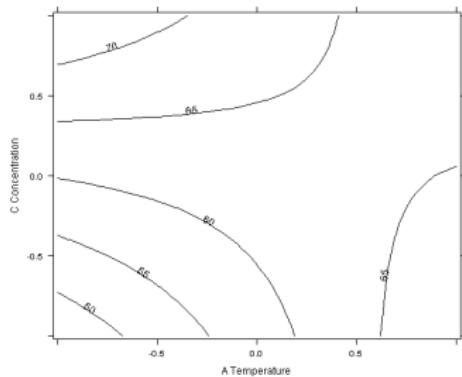
No specification in the code means last value of D, i.e. D = 1.

Pilot Plant Filtration Rate Experiment

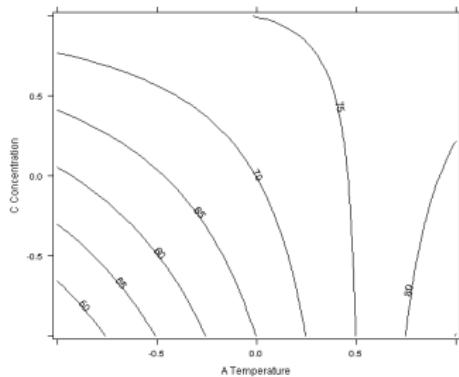
Contour plot: A(temperature) and C(concentration) for different D's.

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

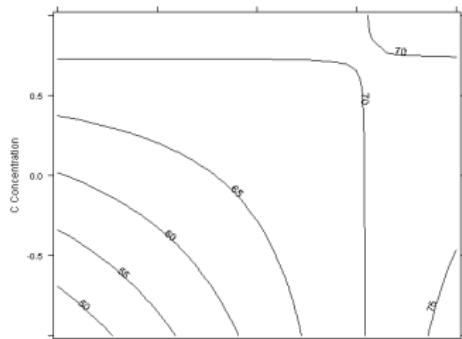
Contour plot of Filtration Rate from the Pilot Plant Experiment: D = -1



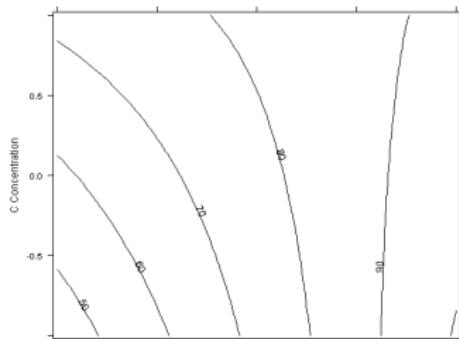
Contour plot of Filtration Rate from the Pilot Plant Experiment: D = 0



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



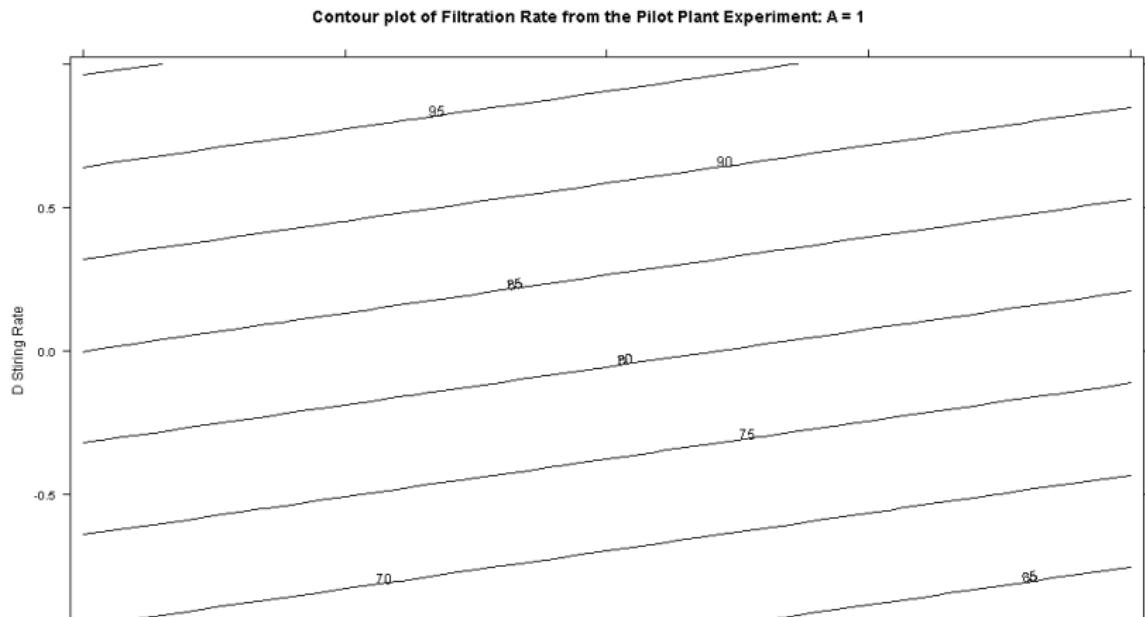
Contour plot of Filtration Rate from the Pilot Plant Experiment: D = 1



Pilot Plant Filtration Rate Experiment

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

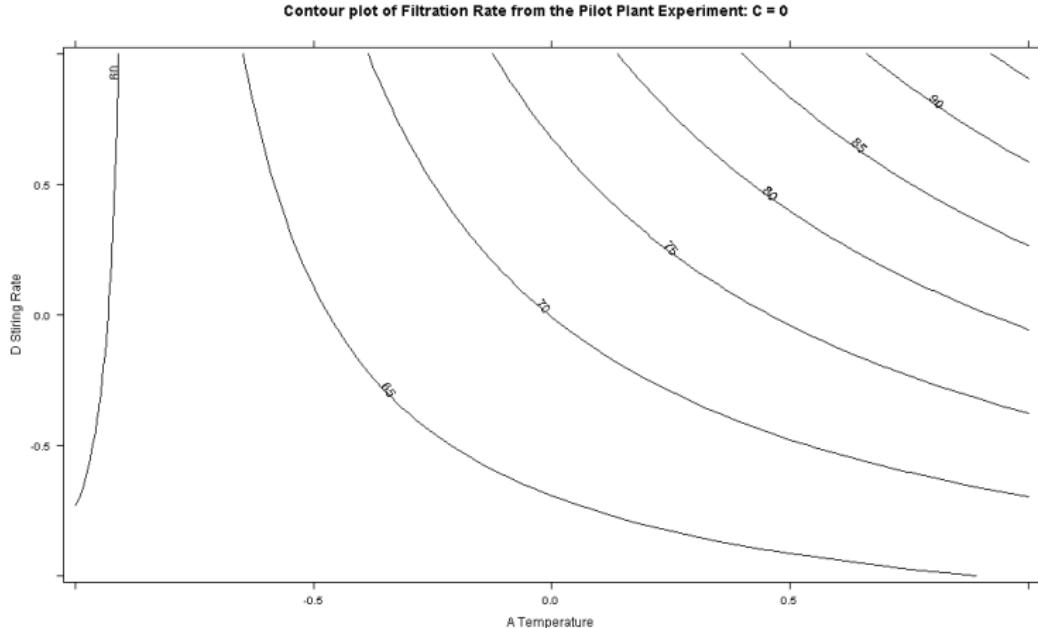
```
> contourplot(fit~C.num*D.num,new.data[new.data$A.num == 1,],  
  xlab="C Concentration",ylab="D Stirring Rate",  
  main = "Contour plot of Filtration Rate from the Pilot Plant Exper  
 A = 1")
```



Pilot Plant Filtration Rate Experiment

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

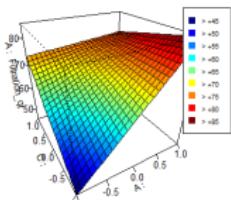
```
contourplot(fit~A.num*D.num,new.data[new.data$C.num == 0,],  
xlab="A Temperature",ylab="D Stirring Rate",  
main="Contour plot of Filtration Rate from the Pilot Plant Experiment  
:C = 0")
```



Pilot Plant Filtration Rate Experiment

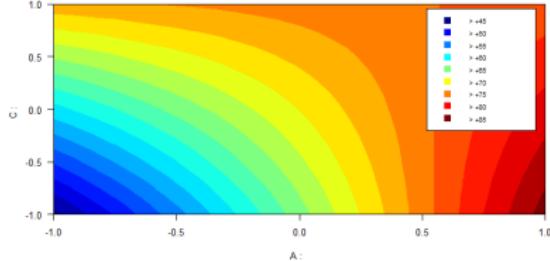
```
wirePlot(A, C, Filtration_qt, data = rate_qt)  
contourPlot(A, C, Filtration_qt, data = rate_qt)
```

Response Surface for Filtration_qt

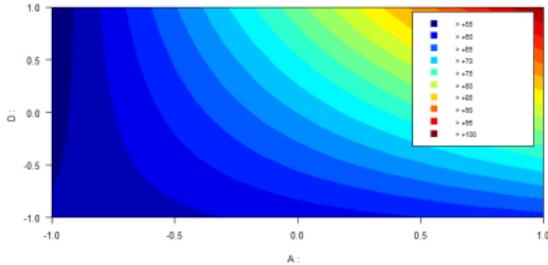


$$\text{Filtration}_\text{qt} \sim A * C + A * D$$

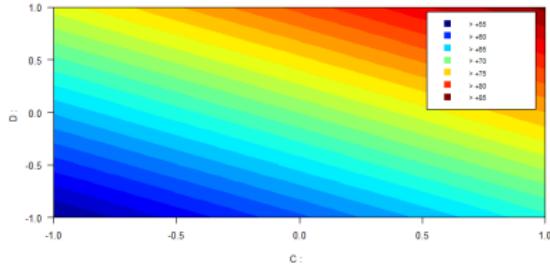
Filled Contour for Filtration_qt



Filled Contour for Filtration_qt



Filled Contour for Filtration_qt



Missing factor is equal to 0 in all three cases!

Today Exercise

Solve problems 6.28 + 6.29 and 6.31 + 6.32 from the chapter 6,
D. C. Montgomery DAoE - 8. edition.
Don't use center points and curvature terms
(will be covered in the next lesson).

Today Exercise

■ TABLE P6.8

The 2⁴ Experiment for Problem 6.31

Run Number	Actual Run Order					Molecular Weight	Viscosity	Factor Levels	
		A	B	C	D			Low (-)	High (+)
1	18	-	-	-	-	2400	1400	A (°C)	100
2	9	+	-	-	-	2410	1500	B (%)	4
3	13	-	+	-	-	2315	1520	C (min)	20
4	8	+	+	-	-	2510	1630	D (psi)	60
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		