

TMA4267 Linear statistical models

25. march 2025

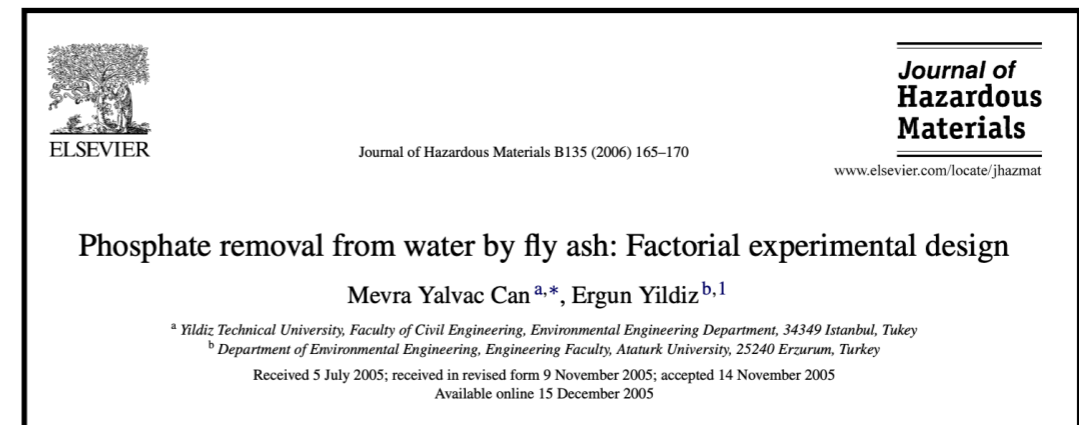
Thea Bjørnland

Last Thursday and today

Three scientific publications (2004, 2005, 2012)
illustrating the use of 2-level factorial experiments,
and motivating the theory that we cover in TMA4267



2^3 duplicated ($n = 16$)



2^3 ($n = 8$)

Repetition

Example: 2^3 full factorial design

$$y = \beta_0 + \beta_1x_1 + \beta_2x_2 + \beta_3x_3 + \beta_{12}x_1x_2 + \beta_{13}x_1x_3 + \beta_{23}x_2x_3 + \beta_{123}x_1x_2x_3 + \varepsilon$$

The length of β is 8, if we do one run we have 8 observations.
To test for significance of effects we discussed 'sacrificing'
some interactions, for example:

$$y = \beta_0 + \beta_1x_1 + \beta_2x_2 + \beta_3x_3 + \beta_{12}x_1x_2 + \beta_{123}x_1x_2x_3 + \varepsilon$$

If the experiment has to be **blocked**, we also took advantage of interactions, for example using AB and AC as blocking factors.
Then, no main effect is confounded by the block effect, but
all of the two-factor interactions are (AB*AC = BC)

Today: fractional factorial designs



Process Biochemistry 40 (2005) 779–788

PROCESS
BIOCHEMISTRY

www.elsevier.com/locate/procbio

Biosorption of chromium using factorial experimental design

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Assume now that we can only afford to do 4 experiments, which?

Tmp	Conc	pH	Tmp:Conc	Tmp:pH	Conc:pH	Tmp:Conc:pH
1	1	1	1	1	1	1
1	1	-1	1	-1	-1	-1
1	-1	1	-1	1	-1	-1
1	-1	-1	-1	-1	1	1
-1	1	1	-1	-1	1	-1
-1	1	-1	-1	1	-1	1
-1	-1	1	1	-1	-1	1
-1	-1	-1	1	1	1	-1

Example 3

Statistics
in Medicine

Research Article

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(wileyonlinelibrary.com) DOI: 10.1002/sim.5526

Application of fractional factorial designs to study drug combinations

Jessica Jaynes,^a Xianting Ding,^b Hongquan Xu,^{a*†}
Weng Kee Wong^c and Chih-Ming Ho^b

<https://doi.org/10.1002/sim.5526>

2^{6-1} fractional factorial design

Example 3: Application of fractional factorial designs to study drug combinations

- Response ('readout'): the percentage of GFP-positive* cells after combinatorial drug treatments
- Factors: 6 drugs for HSV-1 (low and high dosage)
- 2^{6-1} fractional factorial design
- Defining relation: ABCDE = F

Table II. Factors and levels for the initial two-level antiviral drug experiment.		
Factor	Levels (ng/mL)	
	Low (−1)	High (+1)
<i>A</i> = interferon alpha	3.12	50
<i>B</i> = interferon beta	3.12	50
<i>C</i> = interferon gamma	3.12	50
<i>D</i> = ribavirin	1560	2.5e4
<i>E</i> = acyclovir	312	5e3
<i>F</i> = tumor necrosis factor alpha	0.31	5

*GPF-positive means cells carry green fluorescent protein (GFP) gene, which occurs when infected with HSV-1 virus (Herpes simplex virus type 1)

Example 3: Application of fractional factorial designs to study drug combinations

2.4. Analysis and results

As explained in Section 2.2, our design can estimate all six main effects, all 15 two-factor interactions, and 10 pairs of aliased three-factor interactions, assuming that four-factor and higher interactions are negligible.

Effect aliasing is a consequence of using a fractional factorial design. A related concept is **resolution**, which captures the amount of aliasing. This half-fraction design has resolution VI, which allows the estimation of all main effects and two-factor interactions under the assumption that fourth-order and higher interactions are negligible. In general, the higher the resolution of a fractional factorial design, the less restrictive is the assumption regarding which interactions are negligible to obtain a unique interpretation of the data.

Resolution in fractions of 2^p experiments.

Definition. A design is said to be of resolution R if no p -factor effect is aliased with an effect containing less than $R - p$ factors.

Table I. Design and data for the initial two-level experiment: a 2^{6-1} design.

Run	Factor						readout
	<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	<i>F</i>	
1	−1	−1	−1	−1	−1	−1	31.6
2	−1	−1	−1	−1	1	1	32.6
3	−1	−1	−1	1	−1	1	13.4
4	−1	−1	−1	1	1	−1	13.2
5	−1	−1	1	−1	−1	1	27.5
6	−1	−1	1	−1	1	−1	32.5
7	−1	−1	1	1	−1	−1	11.6
8	−1	−1	1	1	1	1	20.8
9	−1	1	−1	−1	−1	1	37.2
10	−1	1	−1	−1	1	−1	51.6
11	−1	1	−1	1	−1	−1	14.1
12	−1	1	−1	1	1	1	19.9
13	−1	1	1	−1	−1	−1	27.3
14	−1	1	1	−1	1	1	40.2
15	−1	1	1	1	−1	1	19.3
16	−1	1	1	1	1	−1	23.3
17	1	−1	−1	−1	−1	1	31.2
18	1	−1	−1	−1	1	−1	32.6
19	1	−1	−1	1	−1	−1	14.2
20	1	−1	−1	1	1	1	22.4
21	1	−1	1	−1	−1	−1	32.7
22	1	−1	1	−1	1	1	41.0
23	1	−1	1	1	−1	1	20.1
24	1	−1	1	1	1	−1	18.7
25	1	1	−1	−1	−1	−1	29.6
26	1	1	−1	−1	1	1	42.3
27	1	1	−1	1	−1	1	18.5
28	1	1	−1	1	1	−1	20.0
29	1	1	1	−1	−1	1	30.9
30	1	1	1	−1	1	−1	34.3
31	1	1	1	1	−1	−1	19.4
32	1	1	1	1	1	1	23.4
33	0	0	0	0	0	0	16.8
34	0	0	0	0	0	0	17.5
35	0	0	0	0	0	0	16.2

NB: 3 'center runs', not in our curriculum,
but useful for estimating variance

Example 3: Application of fractional factorial designs to study drug combinations

2.4. Analysis and results

As explained in Section 2.2, our design can estimate all six main effects, all 15 two-factor interactions, and 10 pairs of aliased three-factor interactions, assuming that four-factor and higher interactions are negligible.

Table III. Estimates for the initial two-level experiment.			
Effect	Estimates	Sum sq.	% Sum sq.
<i>A</i>	0.017	0.009	1
<i>B</i>	0.03	0.029	3.1
<i>C</i>	0.008	0.002	0.2
<i>D</i>	−0.141	0.636	68
<i>E</i>	0.046	0.068	7.3
<i>F</i>	0.024	0.018	1.9
<i>AB</i>	−0.022	0.015	1.6
<i>AC</i>	0.005	0.001	0.1
<i>AD</i>	0.019	0.011	1.2
<i>AE</i>	−0.009	0.002	0.3
<i>AF</i>	0.005	0.001	0.1
<i>BC</i>	−0.009	0.003	0.3
<i>BD</i>	0.008	0.002	0.2
<i>BE</i>	0.008	0.002	0.2
<i>BF</i>	−0.008	0.002	0.2
<i>CD</i>	0.024	0.018	1.9
<i>CE</i>	0.002	0	0
<i>CF</i>	0.003	0	0
<i>DE</i>	0.001	0	0
<i>DF</i>	0.014	0.006	0.7
<i>EF</i>	−0.001	0	0
<i>ABC + DEF</i>	−0.002	0	0
<i>ABD + CEF</i>	0.002	0	0
<i>ABE + CDF</i>	−0.006	0.001	0.1
<i>ABF + CDE</i>	−0.001	0	0
<i>ACD + BEF</i>	−0.017	0.009	0.9
<i>ACE + BDF</i>	−0.015	0.007	0.8
<i>ACF + BDE</i>	−0.012	0.004	0.5
<i>ADE + BCF</i>	−0.004	0	0
<i>ADF + BCE</i>	−0.009	0.002	0.2
<i>AEF + BCD</i>	0.014	0.007	0.7
Residuals	—	0.077	8.3
Total	—	0.935	100

Example 3: Application of fractional factorial designs to study drug combinations

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<i>DF</i>	0.014	0.006	0.7
<i>EF</i>	−0.001	0	0

Table III suggests that the effects of drugs *D* and *E* are the largest. The linear effect of drug *D* is the most significant with an estimate of three times the estimate of the next most significant drug, *E*, showing that drug *D* is very significant and important relative to the other drugs. Together, drugs *D* and *E* account for 75.3% of the total sum of squares in the data. Overall, the six main effects contribute 81.5% of the sum of squares, the 15 two-factor interactions contribute 6.8%, the 10 pairs of three-factor interactions contribute 3.2%, and the residuals account for 8.3%. In this antiviral experiment, the main effects dominate the system, and drug *D* alone accounts for 68.0% of the total sum of squares within the system.

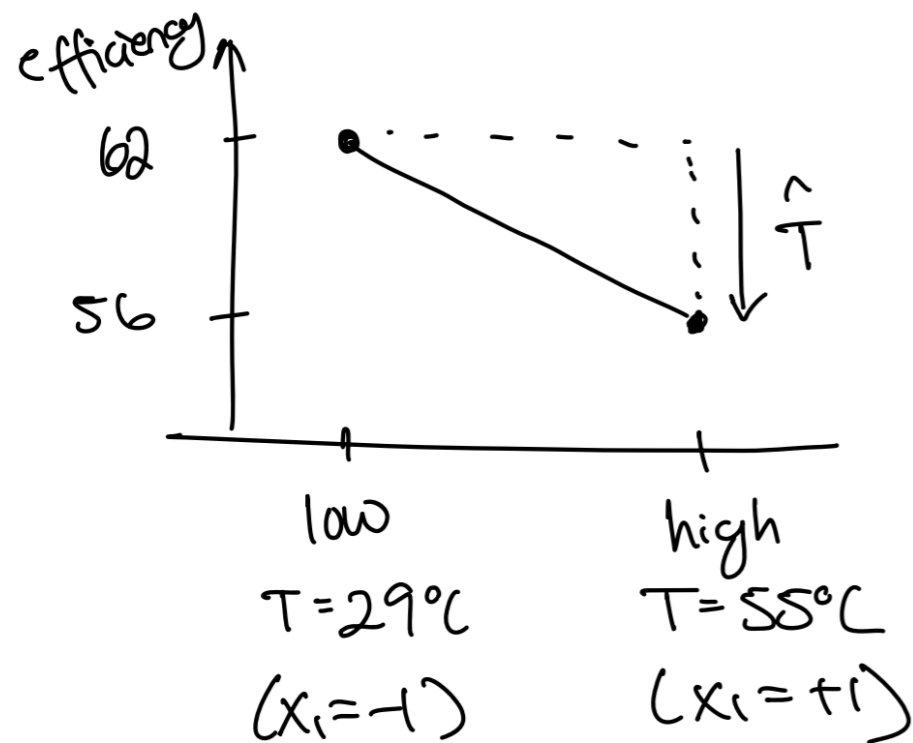
Example 3: Application of fractional factorial designs to study drug combinations

- "A follow-up experiment using a **blocked three-level fractional factorial design** indicates that tumor necrosis factor alpha has little effect and that HSV-1 infection can be suppressed effectively by using the right combination of the other five antiviral drugs"

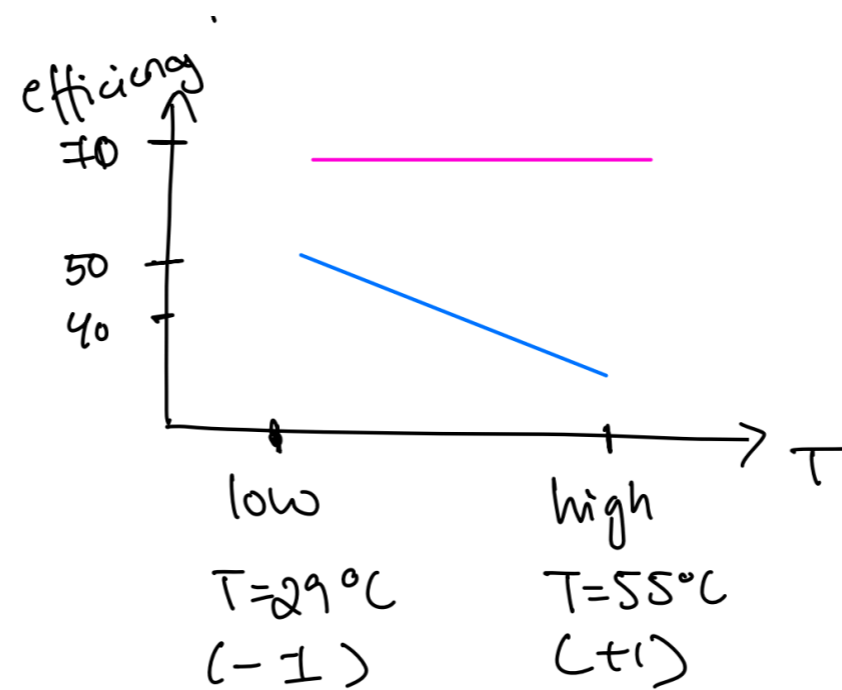
Presentation of results

Some R code for plotting

Main effects:



Interaction effects:



— C low level (10 mg/L)
— C high level (200 mg/L)

Presentation of results

Some R code for plotting

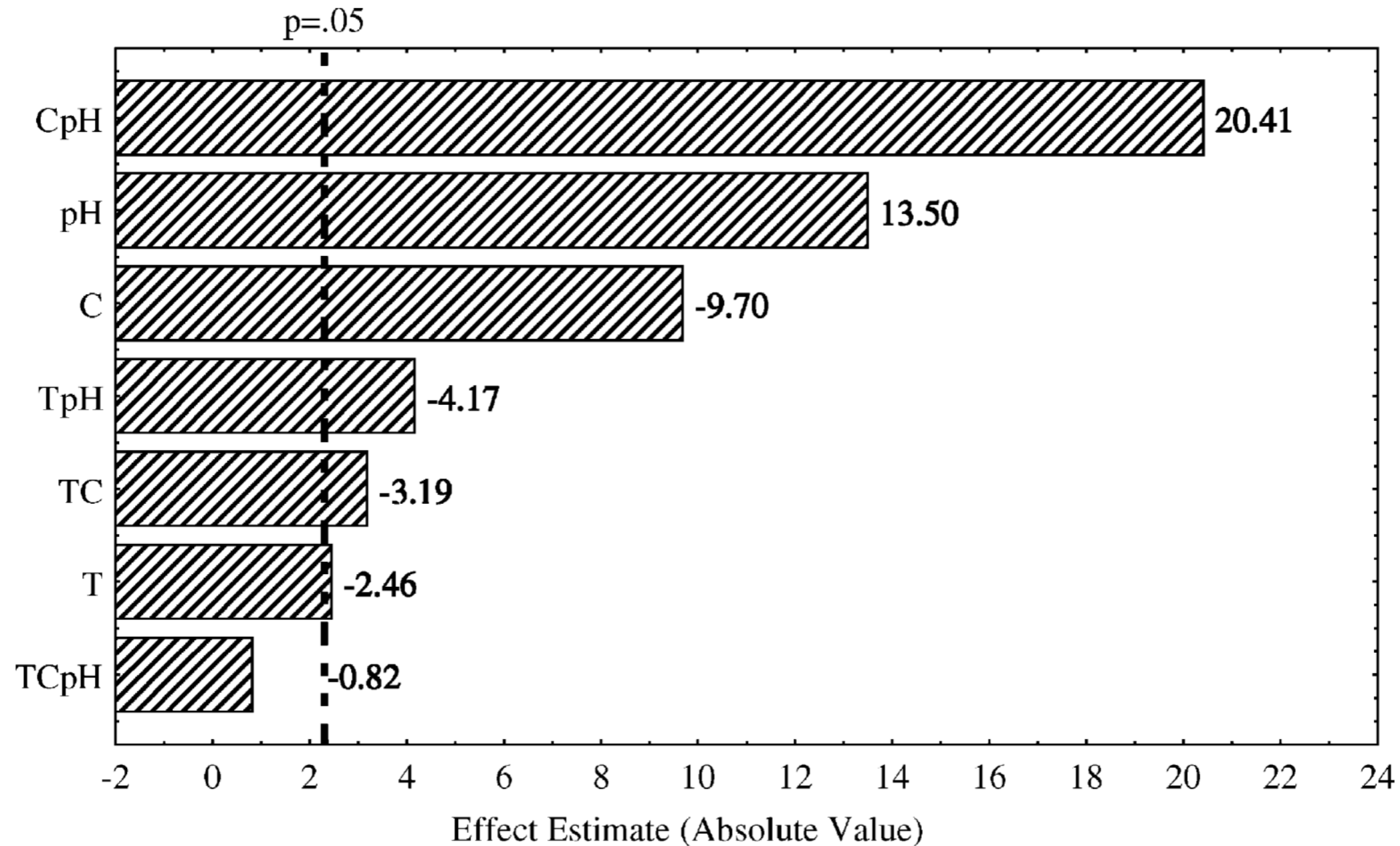


Fig. 1. Pareto chart of effects on the removal efficiency of Cr^{3+} .

NB: Thursday 27th

Use the lecture time (and room) to work on your project and ask questions.

Also use the exercise class.