Alternative Fractional Factorial Designs

BIOE 498/598 PJ

Spring 2022

How low can we go?

The efficiency of fractional factorial designs offsets the exponential increase in runs for factorial designs.



How low can we go? (zoomed in)



					n	umber	of run	S			
		8	16	32	64	128	256	512	1024	2048	4096
							only the MA design				
	3	full									
	4	IV	full								
	5	- 111	V	full							
	6	- 111	IV	VI	full						
	7	- 111	IV	IV	VII	full					
	8		IV	IV	V	VIII	full				
	9		- 111	IV	IV	VI	IX	full			
s	10		- 111	IV	IV	V	VI	X	full		
p	11		- 111	IV	IV	V	VI	VII	XI	full	
aci	12		- 111	IV	IV	IV	VI	- VI	VIII	XII	full
Ť	13		- 111	IV	IV	IV	V	- VI	VII	VIII	XIII
ž	14		- 11	IV	IV	IV	V	- VI	VII	VIII	IX
ĕ.	15		- 11	IV	IV	IV	V	- VI	VII	VIII	VIII
5	16			IV	IV	IV	V	- VI	VI	VIII	VIII
5	17			- 11	IV	IV	V	- VI	VI	VII	VIII
	18			- 11	IV	IV	- IV	- VI	VI	VII	VIII
	19			- 11	IV	IV	IV	V	VI	VII	VIII
	20			11	N	N	- IV	V	VI	VII	VIII
	21			11	IV	N	- IV	V	VI	VII	VIII
	22			- 11	IV	IV	- IV	V	VI	VII	VIII
	23			- 11	IV	IV	- IV	V	VI	VII	VIII
	24			- 11	IV	IV	IV	IV	VI	VI	VIII
Res	olution	ı III up	to	31	63	127			factor	S.	
Resolution IV up to 32 64 8						80	160	factor	S.		
Res	Resolution V up to number of factors:								33	47	65
Resolution VI up to number of factors: 24 34						48					
Circ	+ docia	up	A	. numb	ar of f				-4		40
rirs	t desig	TISIW	H up to			actors			~~		
				- 31	63	127	35	29	28	32	26

Gromping, 2014

J. Stat. Software

Foldover Designs

Imagine a 2_{III}^{6-3} design with $D = AB, \quad E = AC, \quad F = BC$

$$I = ABD = ACE = BCF = DEF$$

= $BCDE = ACDF = ABEF$

After analysis, we find that both B and D are significant.

Since D = AB, the significance of D might be due to B and AB.

We can *augment* the design by doubling the runs *with D flipped*. This clears *D* and its interactions.

Run	Α	В	С	D	Ε	F
1	—	—	—	+	+	+
2	+	—	—	—	—	+
3	-	+	_	_	+	_
4	+	+	—	+	—	_
5	_	_	+	+	—	—
6	+	_	+	_	+	_
7	—	+	+	—	—	+
8	+	+	+	+	+	+

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Run	Α	В	С	D	Ε	F
1	—	—	—	+	+	+
2	+	—	—	—	—	+
3	—	+	—	—	+	—
4	+	+	—	+	—	—
5	—	—	+	+	—	—
6	+	_	+	_	+	_
7	_	+	+	_	_	+
8	+	+	+	+	+	+
9	_	_	_	_	+	+
10	+	_	—	+	_	+
11	_	+	—	+	+	_
12	+	+	—	—	_	_
13	_	_	+	—	_	_
14	+	_	+	+	+	_
15	_	+	+	+	_	+
16	+	+	+	_	+	+

If we combine a Resolution III design with its mirror image (all factors flipped), we have a Resolution $\rm IV$ design with all main effects clear.

If we add a blocking factor we can perform the experimental batches sequentially.

As with foldover designs, mirror image designs are only necessary if more than one main effect is significant.

Blocked Designs

Sometimes logistics force us to group runs into "blocks".

Examples

- Mice need to be housed in separate cages.
- The experimenter cannot do all runs in a single day/batch.
- Two experimenters need to split up the runs.
- A single container of reagent doesn't cover the experiment.

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 - Mice need to be housed in separate cages.
 - The experimenter cannot do all runs in a single day/batch.
 - Two experimenters need to split up the runs.
 - A single container of reagent doesn't cover the experiment.
- A "blocking factor" is added to these experiment to capture inter-block differences.
- Blocks are added to designs and analyzed like any other factor.
- Most block × factor interactions are ignored.

Blocking our Pilot Plant Experiment





Building our normal model

model <- lm(yield ~ T*K*C, data=pilot); show_effects(model)</pre>

##	(Intercept)	60.25
##	Т	9.75
##	К	5
##	C	-1.
##	T:K	3.5
##	T:C	2.
##	K:C	1.75
##	T:K:C	4.25

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##	T:K:C	4.25			
mod	del <- lm(yield	~ T*K*C	+ block,	<pre>data=pilot);</pre>	<pre>show_effects(model)</pre>
	/				
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