Screening Designs

BIOE 498/598 PJ

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Why do we use screening designs?

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- Too many factors waste resources
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- Too many factors waste resources
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- **Solution:** A *screening design* tests a large number of factors
- Only active factors are carried forward for optimization

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We don't worry about estimates of TWIs. We're selecting factors, not interactions.

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 - Con: Complex aliasing

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- Definitive Screening Designs
 - Hybrid screening/optimization design. We'll discuss later!

Don't rule out Fractional Factorial Designs.



						orrans						
		8	16	32	64	128	256	512	1024	2048	4096	
							only the MA design					
number of factors	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		III	IV	IV	- VI	IX	full				
	10		Ш	IV	IV	V	VI	Х	full			
	11		- 11	IV	IV	V	VI I	VII	XI	full		
	12		- 111	IV.	IV	IV.	VI	VI	VIII	XII	full	
	13		111	IV	IV	- IV	V	VI	VII	VIII	XIII	
	14			IV	IV	- IV	V	VI	VII	VIII	IX	
	15		Ξ	IV	IV	١٧	V	VI	VII	VIII	VIII	
	16			IV	IV	IV	V	VI	VI	VIII	VIII	
	17			111	IV	١V	V	VI	VI	VII	VIII	
	18			111	IV	IV	IV	VI	VI	VII	VIII	
	19			- 111	IV	IV	IV	V	VI	VII	VIII	
	20			111	IV	IV	IV	V	VI	VII	VIII	
	21			111	IV	IV	IV	V	VI	VII	VIII	
	22			111	IV	IV	IV	V	VI	VII	VIII	
	23			111	IV	IV	١V	V	VI	VII	VIII	
	24			111	IV	IV	IV	IV	VI	VI	VIII	
Res	olution	ı III up	to	31	127			factor	S.			
Resolution IV up to 32 64								160	factors.			
Rac	Resolution V up to number of fastors								22		65	
nes 0	-lut	. vi up	to nun				33		05			
Resolution vi up to number of factors: 24 34 4										48		
Firs	t desig	n is M	A up to	numb	er of f	actors						
				31	63	127	36	29	28	32	26	

number of runs

Gromping, 2014 J. Stat. Software

Workflow for Resolution III screens

- 1. Run the design
- 2. Fit the model with main effects. If you have DoF left over, add any TWIs that are **not** confounded with main effects.
- If the overall model fit is bad, or if you expected certain effects to be significant that were not, consider a second batch of runs with a mirror image design.
- 4. Drop any factors that are not important (practically or statistically).

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Plackett-Burman Designs

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- > Orthogonal designs, so main effects can be estimated independently
- Run sizes in multiples of 4
- Both PB designs and FF designs are Orthogonal Arrays
 PB = FF when N = 2^k
- PB designs have complex aliasing. Every ME is partially confounded with all TWIs.

Creating a PB design (up to 23 factors)

1. Start with the first run from the following table.

Runs	Factor Levels
12	++-++++-
20	++++++-+-+++-
24	+++++-+-+++++

- 2. Cycle the factor levels by one to get run #2. Repeat for 11, 19, or 23 runs.
- 3. Set the final run to all low (-).
- 4. If the number of factors k is less than the number of runs, select the first k columns.

Workflow for PB designs

- 1. Run the design.
- 2. Fit a model with main effects plus an effect for any unused column in the design.
- 3. Optional: Perform subset regression to identify factors that appear frequently in smaller models with good predictive power.
- 4. Drop any factors that are not important (practically or statistically).
- 5. If only a small number of factors remain, try refitting the small model.

Example PB design: Cast fatigue



This design includes 7 factors; however, effects are estimated for all columns. The last 4 "factors" are interactions with complex aliasing.

To replicate or not to replicate?

- Many screening designs are saturated there are no DoF to estimate confidence intervals for the parameters.
- The number of estimable factors is bounded by the rank of the model matrix. Replicates do not changes the rank.
- If you don't replicate the design, you can select factors based on the magnitude of the effects alone (half-normal plot).
 - Remember that half-normal plots work better as the number of factors grows.

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 - Remember that half-normal plots work better as the number of factors grows.
- Replicating a Resolution III Design
 - Consider a mirror-image instead. This will give clear main effects.
 - Check if you can afford a Resolution IV instead. This gives clear main effects and a confounding structure.
- Replicating a PB Design
 - Replicating the design will help you estimate the "pure error".
 - You can "move up" to a larger PB design to get extra runs. This won't estimate pure error, but you can add more confounded effects to the model to improve the estimates.