

Biometrical Best Practices

in light of 2015 working experience with Afghan colleagues

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Presentation Roadmap

- 2015 Working Stats
- Alpha vs. RCBD
- REML vs. ANOVA
- Can and Can't in Alpha Design
- RCBD vs. DiGGer
- What is the Right Block Size?
- Alpha Implementation
- Layout Information
- Meta Data
- Data Validation and Integrity Check
- Combine Data for MET Analysis
- Excel files vs. Database



2015 Working Stats

Trials# per Location Trials% per Design





CRD, RCBD, and Alpha Designs

A method which partitions the total variation in the response into the components (sources of variation):

* Response = $\mu + \tau + \xi$ (CRD) * Response = $\mu + \tau + \pi + \xi$ (RCBD)

* Response = μ + τ + π + β + ξ (Alpha)







μ) grand mean

 τ) effect of treatments

 π) effect of replicates

tes β) effect of blocks

ξ) experimental error



REML vs. ANOVA







Can and Can't in Alpha Design

You Can

- Get more precision without increase any resources in the field (e.g. land, seeds, labors)
- Analysis your data using simple ANOVA and ignore block information!
- Generate it for free (e.g. ICARDA BioComuting online service, Agricolae R package from CIP, CIMMYT ALPHAGEN software, etc..)

You Can't

- Convert RCBD randomization into Alpha design simply by add an extra column to split your replications into hypothesis blocks!
- Lose precision comparing to RCBD (i.e. worst scenario when blocks within replicates are homogeneous)



RCBD vs. DiGGer

0	4	0
3	1	2
7	14	6
17	8	17
5	24	14
21	5	12
6	12	15
20	20	5
23	11	7
9	21	10
12	7	21
19	15	23
2	2	3
24	19	18
15	17	8
14	6	24
4	10	4
10	4	1
16	9	13
1	3	9
13	18	11
11	23	20
18	22	16
22	13	19
8	16	22

21	5	10
19	14	15
24	7	9
13	6	2
22	8	4
18	16	12
23	17	11
1	3	20
20	23	14
6	9	21
4	13	1
15	11	24
10	19	17
7	2	18
16	22	5
3	12	8
2	10	3
17	4	7
8	18	19
14	24	6
9	20	22
12	21	23
5	15	13
11	1	16



What is the Right Block Size?

- Do NOT ask ME that question! It is constrained by the physical arrangement of plots in the field, for example:
 - Total number of genotypes
 - Field homogeneity
 - Plot size
 - Field layout (i.e. rows & columns)



1 2 3

4 5 6



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Alpha Design Implementation (1)



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DA

Science for Better Livelihoods in Dry Areas

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Layout Information (1)





Layout Information (2)



Rep	Block	Col	Row	Plot
1	1	1	1	1
1	1	1	:	:
1	1	1	6	6
1	2	1	7	7
1	2	1	:	:
1	2	1	12	12
1	3	2	12	13
1	3	2	:	:
1	3	2	7	18
1	4	2	6	19
1	4	2	:	:
1	4	2	1	24





- Experiment name
- Description
- Coordinator name
- Coordinator institute
- Coordinator contact information
- Location
- Province (state)
- Country
- Latitude
- Longitude



- Crop
- List of entries / genotypes
- Season and cycle
- Experiment design
- Total number of entries
- Total number of plots
- Number of replications
- Block size (plots per block)
- Number of rows
- Number of columns



Data Validation and Integrity Check (1)

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Data Validation and Integrity Check (2)

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Use Pivot Table Functionality

Rep	Block	Plot	Geno	Count of Plot	Column Labels 💌			PivotTable Fields 🔹 👻
1	1	1	18	Row Labels 💌	1 2	3 Grand Tot	tal	Channe Galdete addete annat
1	1	2	10	1	2 1	. 1	4	Choose fields to add to report:
1	1	3	14	2	1 1	1	3	✓ Rep
1	1	4	22	3	1 1	1	3	Block
1	1	5	15	4	1 1	1	3	✓ Plot
1	1	6	6	5	1 1	1	3	Geno
1	2	1	24	6	1 1	1	3	MORE TABLES
1	2	2	19	7	1 1	1	3	MORE TABLES
1	2	3	2	8	1 1	1	3	
1	2	4	1	9	1 1	1	3	
1	2	5	21	10	1 1	1	3	
1	2	6	8	11	1	. 1	2	
1	3	1	12	12	1 1	1	3	
1	3	2	1	13	1 1	1	3	Drag fields between areas below:
1	3	3	3	14	1 1	1	3	
1	3	4	5	15	1 1	1	3	
1	3	5	4	16	1 1	1	3	Кер 👻
1	3	6	17	17	1 1	1	3	
1	4	1	13	18	1 1	1	3	
1	4	2	16	19	1 1	1	3	
1	4	3	20	20	1 1	1	3	\equiv ROWS Σ VALUES
1	4	4	7	21	1 1	1	3	Geno Count of Plot
1	4	5	9	22	1 1	1	3	
1	4	6	23	23	1 1	1	3	
2	1	1	13	24	1 1	1	3	
2	1	2	23	Grand Total	24 24	24	72	



Data Validation and Integrity Check (3)

	А	В	С	D	
1	Rep 💌	Block 💌	Entry 🖵	Yield 💌	
7	1	2	10	5.67	
14	2	1	10	11.23	
31	3	2	10	9.80	
39	4	2	10	11.93	
42					

1	Rep 💌	Block 💌	Entry 💌	Yield 📐
	1	1	7	10. 🛃
	1	1	4	10. <u>z</u> j
	1	1	5	12.
	1	1	1	10.
	1	1	9	10. 📉
	1	2	10	5.
	1	2	3	10.
	1	2	8	9.
	1	2	2	11.
	1	2	6	9.
	2	1	7	9.
	2	1	3	9.
	2	1	10	11.
	2	1	4	8.
	2	1	9	9.
	2	2	5	10.
	2	2	1	11.
	2	2	2	10.
	2	2	8	11.
	2	2	6	11

Sort	Smallest	to Largest		
↓ S <u>o</u> rt	Largest t	o Smallest		
ort	by Color			- • -
<u> </u>	ar Filter Fr	om "Yield"		
Filt	r by Colo	r		- • I
Nur	n per <u>F</u> ilter	rs		- • I
Sea	rcl			2
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	✓ 5.67			
	8.13			
	8.43			
	8.50			
	8.64			
	0.57			
	8.87			-
	0.07			×
		OK	Cano	el

Use Filter Functionality

Α	В	С	D	E	F	G	Н
Rep 💌	Block 💌	Entry 💌	Yield 🔻				
1	1	7	10. 2	<u>S</u> ort Smal	llest to Large	st	
1	1	4	10. Z	Sort Large	est to Smalle	st	
1	1	5	12.	Sort by C	olor		
1	1	1	10.		-		
1	1	9	10. 🏷	Clear Filte	er From "Yie	ld"	
1	2	10	5.	F <u>i</u> lter by C	Color		
1	2	3	10.	Number	Filters		•
1	2	8	9.	Search			0
1	2	2	11.	Jan			~
1	2	6	9.		.23		^
2	1	7	9.		.65		
2	1	3	9.	···· 🗹 11.	.67		
2	1	10	11.		.73		
2	1	4	8.		.82		
2	1	9	9.		.97		
2	2	5	10.	···· 🗹 12	.27		
2	2	1	11.	····· 🗹 83.	.1		~
		5	10.				
	2 <		11.		O	(Cancel
(2)			11.				.:
Ş		Z	10.03				
3	1	3	83.1				



Combine Data for MET Analysis

- Genotype name as a join key:
 - FLIP 96-15L (Ibla 1)
 - Ibla 1
 - **FLIP** 96-15L
 - **FLIP96-15L**
 - FLIP 1996-15L
 - Leading and trailing space
 - Letter case (i.e. FLIP, Flip, or flip)
 - Typo (e.g. FLOP ~ FLIP)
- Naming style (e.g. Yield ~ GY or BY)
- Units (e.g. T/ha, Kg/ha, or gr/m2)
- Abbreviation & acronym (e.g. DFLR and DF)





Excel Files vs. Database

DB or not *DB*, is it really your question!?

- Data integrity (independent copies causing duplicate and often outdated versions)
- Query and retrieve data offers a greater range of complexity (SQL)
 - * What was planted in this field during last **n** years?
 - * Who planted **x** genotype in this area and what was the performance?



- Better access control (multi-user, security privileges, web access/open access)
- Better for long-term storage vs. Excel work-in-progress module

DB Cons

DB Pros

Excel is really widely used and more user friendly







Thank you

