# Important Statistic Tools in ARM



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### **Overview: Research Questions**

When we pose a research question, we want to know whether the outcome is from:

- the treatments (independent variables), or
- chance (meaning tested treatments are probably not effective)

based on experiment samples we collect.

## **Statistics**

- Inferential statistics are used to make generalizations from a sample to a population
- The reason for calculating an inferential statistic is to get a p value (p = probability)
- For crop research, Analysis of Variance is commonly used for initial statistical analysis to obtain a *p* value

# AOV Table for RCB Design

- Randomized Complete Block is a common design
- Consider key components of AOV table when planning and conducting trials:



# ARM Tools Can Assist With Important Trial Activities

### Planning

- Trial Layout
- Data Review

Statistical Analysis







#### Summary

- Assessment Data Summary
- AOV Means Table
- Factorial AOV Table
- Correlations
- Dose-Response Analysis

# Planning - Key Considerations

- At least **12 error DF** (degrees of freedom) for RCB: Error DF=(Trts-1) x (Reps-1)
  - 2 T x 13 R
    5 T x 4 R
  - 3 T x 7 R 7 T x 3 R
  - 4 T x 5 R 13 T x 2 R

Include enough replicates to statistically distinguish expected treatment differences
 Minimum alpha for statistical significance?

### Define in ARM Protocol and Trial Settings

#### Protocol Settings

General Design Treatme	ent Application Layout										
Randomized Complete Blo	ock (RCB)		•	Pov	wer and Ef	ficiency					
Factors: 1	Trea	atments	Merge Factor fields to	CV Lock	10.0 🚖 cat 🔲	Reps 4 🚔	Power 80 🚔	αSL <u>5%</u> ▼	% Mean Diff 10.0	V	
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					10	17				64	85
					12	24				92	120
					14	32				124	160

- Help plan experiments that successfully detect expected treatment differences
- Available in both protocols and trials so:
  - Protocol writers can more effectively plan experiments
  - Trialists can verify whether CV expectations are realistic based on local experience for specified crop(s)

Calculates "power" based on:

- Estimated <u>CV</u> of key assessment (e.g. yield)
- Number of <u>replicates</u>
- <u>Power</u> = Level of certainty to detect "real" treatment effects (80% or 90%)
- <u>Alpha</u> Significance Level (e.g. 5%, 10%)
- Mean Diff = estimated treatment effect, expressed as percentage of overall (grand) mean across treatments of key assessment



- "Lock at" to keep 3 columns constant
- Calculates table of possible values for "unlocked" columns (e.g. Rep or CV)
- Values entered by protocol writer are carried into trials created from protocol, conveying protocol expectations

# Compare effect of significance level on minimum replicates for CV=6% vs. 10%

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	10	16		7	6			
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1			- L	8	8			
				10	13			
				12	18			
				14	25			

# Consider impact of Replicates on precision to detect treatment differences



# ARM Tools Can Assist With Important Trial Activities

Planning

### Trial Layout

Data Review

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Trial Layout

Experimental site

- Is it uniform?
- Is there a known gradient?
- Use an effective Block (replicate)
  - 1. Placement: within rep more uniform than trial
  - 2. Shape: reduce distance between plots
- Want to decrease Error Sum of Squares by increasing Replicate Sum of Squares

Trial Layout Should Include Randomization Quality Review

Goal is to improve experiment precision:

- 1. Arrange replicates as squares, not strips
- 2. Equalize treatment distribution
  - a. Balance average distance from all other treatments
  - **b.** Balance "Edge effect" across treatments
- 3. Randomize all replicates

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# Arrange Replicates as Squares not Strips

### "Optimum" is smallest surface-to-area ratio

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	Rep Ler	ngth	103	77	51	25			
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	Trial W	/idth	50.5	67.5	101.5	203.5			
	Trial Le	ngth	415	311	207	103			

Jan. 2017

### **Equalize Treatment Distribution**

"Undesirable" layout of 7 treatments and 5 replicates in Randomized Complete Block:

- Trt. 6 in middle 3 columns of all reps
- Trt. 5 in right 2 cols for all but one plot



Example from Federer, "Experimental Design" 1955 18

Uses "Average Distance of Treatment" Comparison (ADTC)

- van Es and van Es, "Spatial Nature of Randomization and Its Effect on the Outcome of Field Experiments", Agronomy Journal, 85:420-428 (1993).
- Comparison between treatments 1 and 2 is taken from 5 plots for each treatment.
- Measure the plot-to-plot distance for each plot containing treatment 1 to the paired plot within replicate containing treatment 2, for a total of 5 distances.
- ADTC for the treatment pair 1-2 is the average of the 5 distances.

### Distances, Treatments 1-2

### Average distance = 3 plots = 24 feet

for 8 foot wide plots



### **Unequal Treatment Distribution**

- Average distance from 17.9 to 24.6
- Ranges from 11.9(T3,T6) to 34(T2,T5)
   Error variances for treatments may not be homogeneous

501 7	502 2	503 6	504 3	505 4	506 1	507 5	Treat	tment				B
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2		000	001	0.05	000	0.07	2	3	24.6	5.56	17	34
301 7	302 5	303 4	304 6	305 2	306 3	307	3	2	19.8	5.66	(11.9)	25.5
·	0	-	•	2			4	3	21.3	3.18	17	25.5
201 1	202	203	204 4	205 6	206	207	5	3	27	5.83	20.4	34
•	1	J	4	0	2	J	6	2	17.9	3.53	(11.9)	22
101 2	102 4	103 7	104 1	105 6	106 3	107 5	7	3	23.8	4.3	18.7	29

### Unbalanced "Edge effect"

### Treatment 1 occurs at edge 4 times, T2 and T3 at edge only 2 times



# Balanced Treatment Distribution and Edge Effect

Average distance from 21.3 to 24.4
Distances range from 18.7 to 27.2
"Edge effect" is balanced



### Randomize All Replicates

- This frame displays when a nonrandomized replicate is defined in Settings
- Select "Randomize All Replicates" to follow recommended statistical practice

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	Rep Length	103	77	51	25			statistical practice to randomize all replicates.
	Surface/Area	0.059	0.056*	0.059	0.090			

# ARM Tools Can Assist With Important Trial Activities

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### **Assessment Data Review**

- Check for measurement or entry mistakes
  - If possible, perform <u>immediately after making</u> <u>the assessment</u>, while it is still feasible to verify and correct any mistake
  - If do check "later", only option may be to mark any questionable measurement as <u>missing data</u>, losing 1 error DF for each missing data point

Goal: preserve error DF for better precision

### **Assessment Data Review Tools**

Assessment Data - Line 1 Column Number

### Tools button group

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	and the second se			P(Friedman's X2): 0.072	Hide empty fields
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- Pop-out" editor panel
- Presents analysis of the current data column
- Offers to attempt auto-fix for any violations of AOV assumptions
- Find statistical outliers

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- Column navigation
- Column description
- Min & Max limits
- Actual value range
- Transformation formula description
- Click "Refresh" to update after changing current data column

Column 5 Propert	ties	<b></b>
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Min/Max entry:	0	100
Low/High value:	0.00	15.00
Formula AL:	LOG([5]	+ 1)

Refre

- Descriptive statistics from full AOV of data
- Displays violations of AOV assumptions
- "Fix" prompts if can resolve violations



of variance/skewness/kurtosis

Help

Fix

No

ARM - SPECIAL CONFIRMATION



Yes

- Search for outliers in current data column using a standard outlier test
- "Find Next" locates each statistical outlier



"Damaged" <sup>1.90</sup> Find Next
drops value from outlier test and AOV

- If you will consistently review every assessment using Column Properties
- Then it is safe to set this summary option

Fields to Print	Global - General	Global - Page Heading
AOV Means Table Report Options Missing data estimates		ary General
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### **Assessment Data Review Tools**

9

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### Tools button group

Assessment Data - Line 1 Column Number

Column Properties panel

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P(Friedman's X	(2):	0.072		Hide empty fields
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				GPS.

### **Box-Whisker Graph**

### 'Box percentiles' and 'Outliers based on' options of 'Plot' or 'Subsamples' for boxwhisker





Display data



0.00

0.00

#### 35

### **Assessment Map**

"Heat map" shows response differences per

assessment data column by color intensity:

- Treatment consistency
- Possible site variations as dark or light zones

Plot problems



### **Assessment Map**

### 'Color by current treatment' option:

- Easily review variation in each treatment
- Simpler to find position in each replicate for trial with many treatments



### More Information in ARM Help

He	p		
	Contents •		
	Search for Help On 🕨		
2	Topic F1		
+	Study Definition F5		
1	References (pdf)	Quick Reference	
	How To Topics	Trial Map Overview	
	Tutorial +	Study Rules	
	Media	ARM Action Codes Overview	
or 🐔	BBCH Growth Stages (pdf)	Data Transformations Overview	×
d	Check for Updates	Recommended Assessment Review Methods Overview	×
DI	Detect and Repair	Treatment Homogeneity and Normality Overview	ent

Recommended Assessment Review Methods

Recommended methods to review assessment data columns are located in Properties panel of the Assessment Data editor:

- Column Properties sub-panel
- Tools section of Properties panel

Recommended practices are to:

- Review new assessments while still at trial site, to most efficiently verify whether any unusual values may be from a data entry mistake or a possible site (non-treatment) effect.
- Use Column Properties to review for entry mistakes or violations of AOV assumptions
- Use Tools for detailed review:

Column Number		5 *	Properties					
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	3	7	1	t	101	3	2.00	Assessment Map
		+		14				



- Planning
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### Statistical Analysis summary

- - Assessment Data Summary
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- **Dose-Response Analysis**

### **Statistical Analysis Best Practices**

- Use "Least squares" analysis options on AOV and Factorial AOV Reports
- Use "Post-hoc power analysis" to better plan follow-up experiments
- Do "protected" mean comparison tests
- Be careful using LSD to hand-compare treatment pairs on AOV Means when a data correction transformation is applied

### **AOV Means Table Report Options**

### "Analysis method" options:

Least square estimation (like SAS GLM)Print adjusted means

Pre-mix Ingredient	Fields to Print	Global - General	Global - Page Heading
AOV Means Table	Report Options	General Summa	ary General Sum
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adi isted treatment mea	30		
Use adjusted mean	an I as primary mean		0.0000
Use adjusted mean	an I as primary mean		
Use adjusted mean	an as primary mean		Analysis method
Use adjusted mean	an as primary mean		Analysis method Traditional AOV

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## **Factorial** AOV Report Options

### "Analysis method" options:

Least square estimation (like SAS GLM)
Print adjusted means (if missing data)

Factorial AOV Table Report Options

Report Options General	Summary Report Preview	
Mean comparison test Test: Significance or alpha lev Only when significan Symbol indicating no sign treatment means:	Tukey's HSD         vel:       5%         t AOV treatment P(F)         nificant difference between       -	AOV tables to print Complete Pooled error Both Analysis method
Adjusted treatment mean Use adjusted mean a	as primary mean	<ul> <li>Traditional AOV</li> <li>Least square estimation</li> </ul>
Descriptive statistics for LSD (or HSD if Tuke) Standard Deviation Coefficient of Variation	each factor mean section y's) on	

# Post-hoc Power Analysis (for Follow-Up Experiments)

 Optional descriptive statistic on AOV Means Table report



- Lists, for each assessment column, the minimum number of replicates required to statistically separate treatment means based on Treatment P(F) and current significance level
- Use for planning future trials

# Post-hoc Power Analysis (for Follow-Up Experiments)

- In example, LSD can distinguish a 25% mean difference, yet largest existing difference is 18%
- Current AOV Trt P(F) is 0.2979, so use 0.30+ significance level to separate treatment means
- Need 8+ replicates to reject null hypothesis at 0.05 significance

Crop Variety	CEZANNE
Trt	
No.	24
2	85.33 a
3	81.67 a
4	98.00 a
5	95.33 a
LSD P=.05 (% mean diff)	21.808 (25%)
Standard Deviation	10.915
CV	12.12
Grand Mean	90.083
Minimum Replicates (power = 80)	8
Largest Mean Difference (% mean diff)	16.333 (18%)
Treatment F	1.541
Treatment Prob(F)	0.2979

### "Protected" Mean Comparison Test

Mean comparison test is only performed on assessment data columns when the  $\alpha$  Signif. Level >= AOV Treatment Prob(F)

$\alpha$ SL=0.0	)5
Trt No.	9*
1	2.50 b
2	22.50 ab
3	33.00 a
4	29.00 ab
5	17.50 ab
LSD P=05 CV	19.666 61.07
Replicate F Replicate Prob(F) Treatment F Treatment Prob(F)	0.626 0.6121 3.466 0.0421

<u>αSL=0.0</u>	)1
Trt No.	9*
1	2.50 -
2	22.50 -
3	33.00 -
4	29.00 -
5	17.50 -
LSD P=01 CV	27.570 61.07
Replicate F Replicate Prob(F) Treatment F Treatment Prob(F)	0.626 0.6121 3.466 0.0421

## **AOV Means Table Report**

If data is transformed by square root, log, or arcsine SQR %: shows range of LSD values

used to compare the largest and smallest non-zero means - is a range because they are all non-linear transformations

Pest Code Part Rated Rating Date Rating Type Rating Unit Days After First/La ARM Action Codes	st Applic.	SEPTTR LEAF3 P Jun-18-2014 PESSEV % 64 15 AL	SEPTTR LEAF2 P Jul-2-2014 PESSEV % 78 29 AA
Trt Treatment No. Name	Rate Rate Unit	5*	7*
1 Untreated Che	eck	6.89 a	15.14 a
2 Tub	0.5 l/ha	1.28 b	1.23 bc
3 Tub	1 l/ha	1.04 b	0.40 c
4 Tilt 250	0.5 l/ha	1.60 b	1.51 bc
5 Mico 60 Fungol	1.5 ∥ha 1.25 ∥ha	1.19 b	3.09 b
LSD P=.05 CV		1.878 - 3.785 39.69t	1.218 - 4.299 23.81t

### **AOV Means Table Report**

Example using this log transform:

Compare 6.89 vs. 1.60 using LSD≈3.785

(largest mean vs. next largest)

Rating Typ∈ Rating Unit ARM Action	PESSEV % AL
Trt Treatm No. Name	5*
1 Untreat	6.89 a
2 Tub	1.28 b
3 Tub	1.04 b
4 Tilt 250	1.60 b
5 Mico 60 Fungol	1.19 b
LSD P= 05	1 878 - 3 785

Compare 1.04 vs. 1.19 using LSD≈1.878 (smallest mean vs. next smallest)

## Additional Information, See Statistical Handbook 3<sup>rd</sup> Edition

A Statistical Handbook for Agricultural Field Trials Specialists Third Edition

by Gérard de Kerchove d'Exaerde



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## Thank You

# Contact us with questions at GDM.ARM.Support@gdmdata.com



