

# **Welcome to the Introduction to Augmented Design Webinar**

**Today's Presenter: Dr. Jennifer Kling**

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# Introduction to Augmented Designs

Applications in Plant Breeding

Jennifer Kling

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# Outline – Augmented Designs

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- Essential features
  - When are they used in plant breeding?
- Design options
  - Today - one-way control of heterogeneity
- Augmented Block Design - Example
  - Randomization and Field Plan
  - Analysis with SAS
  - Interpretation of Results
- Overview of variations on the basic design
- Software and Further References

# Augmented Designs - Essential Features

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- Introduced by Federer (1956)
- **Controls** (check varieties) are replicated in a standard experimental design
- **New treatments** (genotypes) are not replicated, or have fewer replicates than the checks – they **augment** the standard design

# Before Augmented Designs

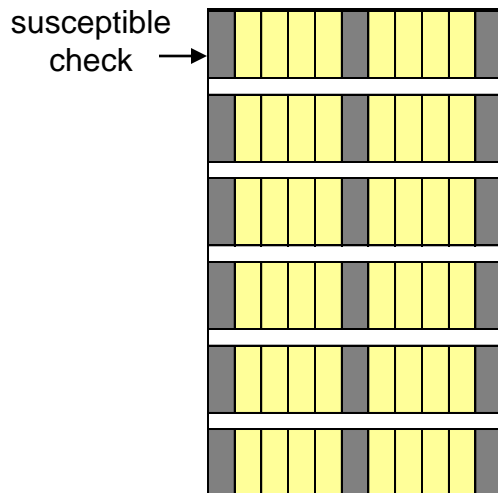
Breeding for Resistance to  
*Striga* in Sorghum at ICRISAT

**Systematic use of checks**



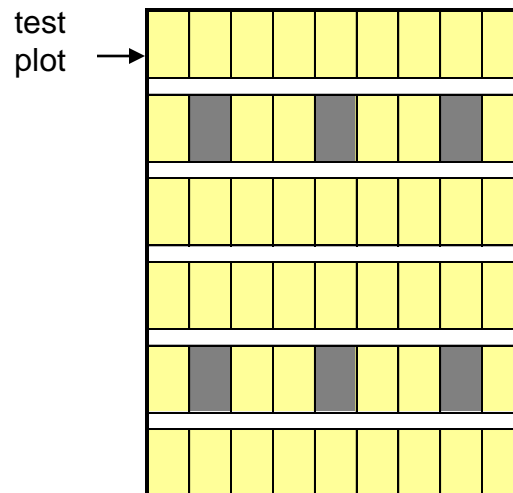
Stage I

Observation Nursery  
(unreplicated)



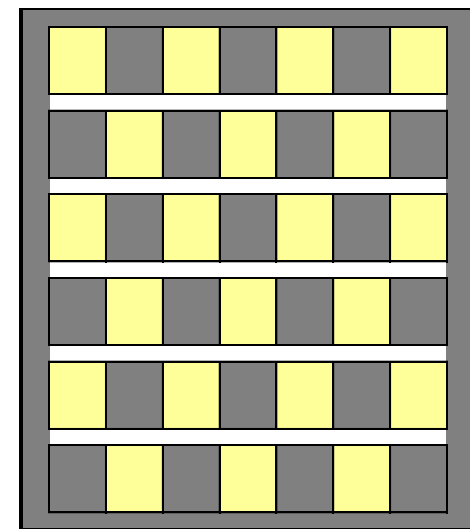
Stage II

Preliminary Screening  
(replicated)



Stage III

Advanced Screening  
(replicated)



Checkerboard

# Augmented Designs - Advantages

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- Fewer check plots required than for designs with systematic repetition of a single check
- Provide an estimate of **standard error** that can be used for comparisons
  - among the new genotypes
  - between new genotypes and check varieties
- Observations on new genotypes can be adjusted for field heterogeneity (blocking)

# Some Disadvantages

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- Considerable resources are spent on production and processing of control plots
- Relatively **few degrees of freedom for experimental error**, which reduces the power to detect differences among treatments
- Unreplicated experiments are inherently imprecise, no matter how sophisticated the design

# When are they used?

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## ■ Early generations

- seed is limited
- too many new entries to replicate
- difficult to maintain homogeneity within blocks with so many entries

## ■ On-farm research

- growers may prefer to grow a single replication
- may not be able to accommodate all entries

# Design Options

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- Choose a design that is appropriate for controlling the heterogeneity in the experimental area
  - One-way blocking
    - Randomized Complete Block Design
    - Incomplete Block Designs (e.g. Lattice Design)
  - Two-way blocking
    - Latin Square (Complete Blocks)
    - Youden Square (Incomplete Blocks)
    - Row-Column Designs

# Design Options

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- Underlying design refers to assignment of checks to the experimental units
- All augmented designs are **incomplete** with respect to the new entries
- Can be replicated in different environments
  - Need to consider relative efficiency compared to other possible designs

# Augmented Block Design Example

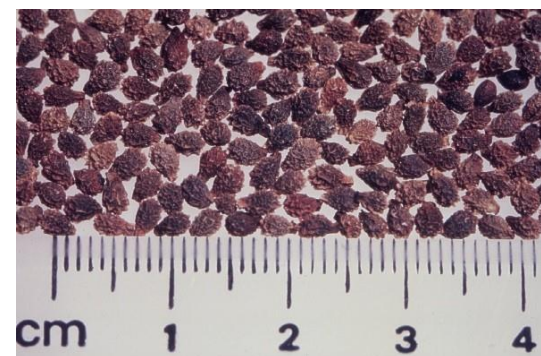
---

- Control heterogeneity in one direction
- Simplest case:
  - Checks occur once in every block
  - New entries occur once in the experiment
- This is an **ARCBD**

# Meadowfoam (*Limnanthes alba*)

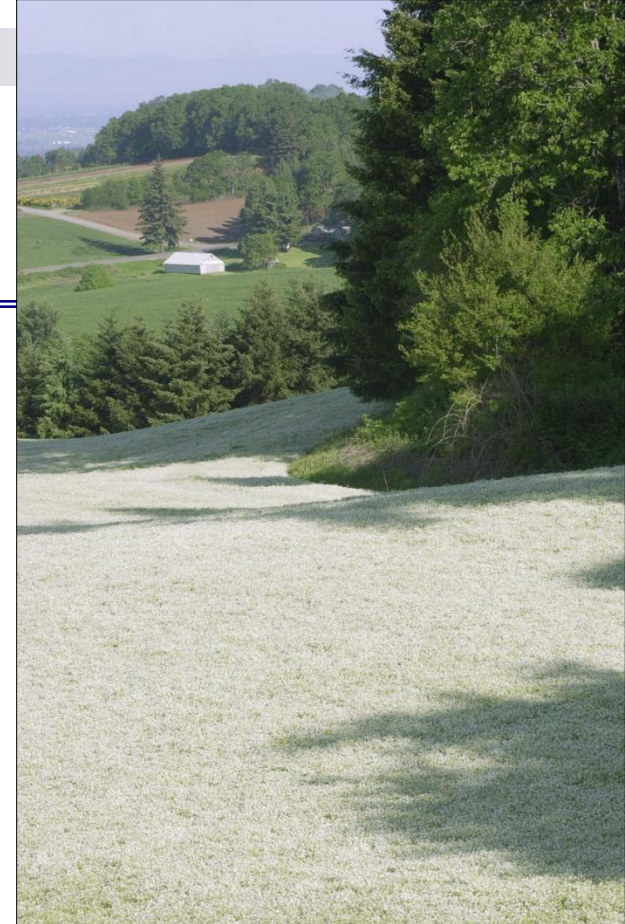
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- Native plant first produced as a crop in 1980
- Seed oil with novel long-chain fatty acids
  - light-colored and odor free
  - exceptional oxidative stability
- Used in personal care products
- Potential uses
  - fuel additive
  - vehicle lubricants
  - pharmaceutical products



# Meadowfoam in Oregon

- Good rotation crop in the Willamette Valley
  - Winter annual
  - Plant and seed meal are high in glucosinolates
  - Same equipment as grass seed



# Pollinators

## Honeybees



**Blue Orchard  
Bees**



**Blue Bottle  
Flies**

# The Germplasm

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- Diverse breeding populations were inherited from a retired breeder (Gary Jolliff)
- Populations were regenerated in the greenhouse and selfed
- $S_2$  lines were transplanted to the field and allowed to outcross
- Insufficient seed for replicated progeny trials
- **Goal** - form a broadbased pool for recurrent selection
  - Screen  $S_2$ -testcrosses
  - Recombine selected  $S_2$  parents


# The Experiment (2007-2008)

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- New treatments:
  - 50  $S_2$ -testcross families
- Check varieties:
  - Ross (C1)
    - Cycle 4 of an elite population (OMF58)
    - Widely grown commercial variety
  - OMF183 (C2)
    - Cycle 5 from OMF58
  - Starlight (C3)
    - released variety derived from this germplasm collection

# The Design

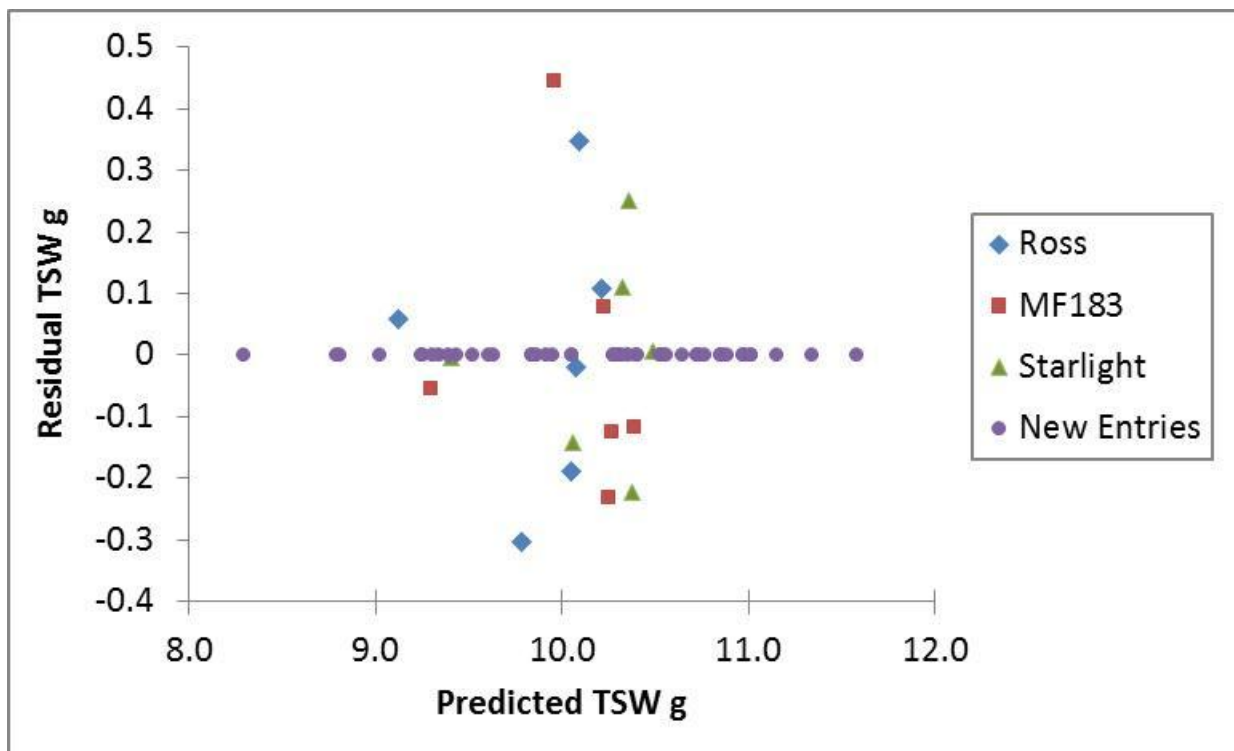
---

- Block in one direction
- 3 checks ( $c=3$ )
- 6 blocks ( $r=6$ )
- Error degrees of freedom =  $(r-1)(c-1) = 10$  
- 50 new entries ( $n=50$ )
- Number of plots =  $n + r*c = 50 + 6*3 = 68$
- Plots per block =  $68/6 = 11.3 \rightarrow 12$  ( $c=3, n=9$ )
- Last block has only 8 plots ( $c=3, n=5$ )
  - Unequal numbers of new entries per block is OK

# Statistical Model

$$Y_{ij} = \mu + \beta_i + c_j + \tau_{k(i)} + \varepsilon_{ij}$$

mean + blocks + checks + new entries + error



# Field Plan

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- Include sufficient number of checks and replicates to provide a good estimate of experimental error and adequate power for detecting differences among varieties
- Arrange blocks along field gradient to maximize variation among blocks and minimize variation within blocks
- Assign each of the checks at random to each of the blocks

# Designation of plots in blocks

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Block 6

168	167	166	165	164	163	162	161
-----	-----	-----	-----	-----	-----	-----	-----

Block 5

149	150	151	152	153	154	155	156	157	158	159	160
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

Block 4

148	147	146	145	144	143	142	141	140	139	138	137
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

Block 3

125	126	127	128	129	130	131	132	133	134	135	136
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

Block 2

124	123	122	121	120	119	118	117	116	115	114	113
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

Block 1

101	102	103	104	105	106	107	108	109	110	111	112
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

# Checks assigned to plots in each block

Block 6	168	167	166	165	164	163	162	161
				C2			C1	C3

Block 5	149	150	151	152	153	154	155	156	157	158	159	160
			C1						C3			C2

Block 4	148	147	146	145	144	143	142	141	140	139	138	137
	C3				C2			C1				

Block 3	125	126	127	128	129	130	131	132	133	134	135	136
						C1					C2	C3

Block 2	124	123	122	121	120	119	118	117	116	115	114	113
	C1		C3							C2		

Block 1	101	102	103	104	105	106	107	108	109	110	111	112
			C2				C1					C3



# Meadowfoam progeny trials

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# Data Collection

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- Flowering dates, plant height, disease resistance, lodging, seed yield, 1000-seed weight, (oil content)
- For this example: 1000-seed weight (**TSW**)
  - Relatively high heritability
  - Positively correlated with oil content and oil yield

# Fixed or Random?

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- Generally assume that blocks are random
  - They represent a larger group of potential blocks
  - We want to make inferences beyond the particular sample of blocks in the experiment
- Genotypes
  - Checks are fixed effects – we are interested in making comparisons with these specific varieties
  - New entries could be fixed or random
    - May be fixed in this case – want to select the winners
    - Most commonly random – particularly in genetic studies

# SAS data input – genotypes fixed

```
DATA anyname;
```

```
INPUT Plot Entry Name$ Block TSW;
```

```
datalines;
```

```
101      8          31          1          9.25
102     45         126          1          9.43
103     90        MF183          1         10.30
104      9          34          1          9.02
105     27          89          1         10.97
106     28          90          1         10.52
107     89          Ross          1          9.86
108     42         121          1         10.56
109     18          68          1         10.39
110     14          57          1         10.97
111     41         118          1         10.32
112     91   Starlight          1         10.44
.         .           .           .           .
.         .           .           .           .
161     91   Starlight          6         10.15
162     89          Ross          6         10.44
163     25          85          6         10.85
164     46         132          6          9.52
165     90        MF183          6         10.14
166     26          87          6         10.73
167     33         101          6          9.24
168     17          66          6          8.79
```

```
;
```

*Other options for data input*

- Import Wizard
- Infile statements
- SAS libraries

# Analysis #1 – new entries fixed

---

```
PROC MIXED;  
TITLE 'Augmented Design using PROC MIXED - entries fixed';  
CLASS block entry;  
MODEL TSW=entry;  
RANDOM block;  
/*compare all means using Tukey's test*/  
LSMEANS entry/pdiff adjust=tukey;  
/*test for entries that exceed Starlight*/  
LSMEANS entry/pdiff=CONTROLU('91') adjust=dunnett;  
ods output lsmeans=TSWadj diffs=TSWpdiff;  
RUN;
```

Use export wizard to export

- TSWadj*
- TSWpdiff*

# Results for Analysis #1 (fixed entries)

---

## The Mixed Procedure Covariance Parameter Estimates

Cov Parm	Estimate
Block	0.1381
Residual	0.06981

## Fit Statistics

-2 Res Log Likelihood	17.7
AIC (smaller is better)	21.7
AICC (smaller is better)	22.7
BIC (smaller is better)	21.3

## Type 3 Tests of Fixed Effects

Effect	Num DF	Den DF	F Value	Pr > F
Entry	52	10	7.51	0.0008

# Output from Dunnett Test

*TSWpdiff.xls*

Effect	Entry	_Entry	Estimate	StdErr	DF	tValue	Probt	Adjustment	Adjp
Entry	1	91	000.3879	000.3131	10	1.24	0.1219	Dunnett-Hsu	0.9351
Entry	2	91	-000.5862	000.3131	10	-1.87	0.9547	Dunnett-Hsu	1.0000
Entry	3	91	000.4366	000.3131	10	1.39	0.0967	Dunnett-Hsu	0.8879
Entry	4	91	-000.1362	000.3131	10	-0.44	0.6636	Dunnett-Hsu	1.0000
Entry	5	91	000.5366	000.3131	10	1.71	0.0587	Dunnett-Hsu	0.7459
Entry	6	91	-000.4782	000.3131	10	-1.53	0.9212	Dunnett-Hsu	1.0000
Entry	7	91	-000.7034	000.3131	10	-2.25	0.9758	Dunnett-Hsu	1.0000
Entry	8	91	-001.0565	000.3131	10	-3.37	0.9965	Dunnett-Hsu	1.0000
Entry	9	91	-001.2865	000.3131	10	-4.11	0.9989	Dunnett-Hsu	1.0000
Entry	10	91	-000.0282	000.3131	10	-0.09	0.5350	Dunnett-Hsu	1.0000
Entry	11	91	001.4666	000.3131	10	4.68	0.0004	Dunnett-Hsu	0.0149
Entry	12	91	000.4518	000.3131	10	1.44	0.0898	Dunnett-Hsu	0.8700
Entry	13	91	000.3266	000.3131	10	1.04	0.1607	Dunnett-Hsu	0.9726
Entry	14	91	000.6635	000.3131	10	2.12	0.0300	Dunnett-Hsu	0.5249
Entry	15	91	-000.4921	000.3131	10	-1.57	0.9265	Dunnett-Hsu	1.0000

# Analysis #2 – new entries random

```
DATA arcabd;  
set anyname;  
if(entry>50) then new=0;  
else new=1;  
if(new) then entryc=999;  
else entryc=entry;  
RUN;
```

Plot	Entry	Name	Block	TSW	new	entryc
101	8	31	1	9.25	1	999
102	45	126	1	9.43	1	999
103	90	MF183	1	10.3	0	90
104	9	34	1	9.02	1	999
105	27	89	1	10.97	1	999
106	28	90	1	10.52	1	999
107	89	Ross	1	9.86	0	89
108	42	121	1	10.56	1	999
109	18	68	1	10.39	1	999
110	14	57	1	10.97	1	999
111	41	118	1	10.32	1	999
112	91	Starlight	1	10.44	0	91
113	31	96	2	11.58	1	999
114	3	15	2	9.95	1	999
115	90	MF183	2	9.24	0	90

# Analysis #2 – new entries random

---

```
PROC MIXED;  
TITLE 'Augmented Design using PROC MIXED - entries random';  
CLASS block entry entryc;  
MODEL TSW=entryc;  
RANDOM block entry*new/solution;  
LSMEANS entryc;  
ods output solutionr=eblups;  
RUN;
```

Use export wizard to export

*eblups*

# Results for Analysis #2 (random entries)

---

## The Mixed Procedure Covariance Parameter Estimates

Cov Parm	Estimate
Block	0.01418
new*Entry	0.3659
Residual	0.1748

## Fit Statistics

-2 Res Log Likelihood	136.5
AIC (smaller is better)	142.5
AICC (smaller is better)	142.9
BIC (smaller is better)	141.9

## Type 3 Tests of Fixed Effects

Effect	Num DF	Den DF	F Value	Pr > F
entryc	3	10	0.89	0.4779

# Estimated Best Linear Unbiased Predictors

*eblups.xls*

Effect	Block	Entry	Estimate	StdErrPred	DF	tValue	Probt
Block	1		000.0189	000.1018	10	0.19	0.8564
Block	2		-000.1264	000.1018	10	-1.24	0.2427
Block	3		000.0188	000.1018	10	0.18	0.8569
Block	4		000.0733	000.1018	10	0.72	0.4882
Block	5		000.0156	000.1018	10	0.15	0.8811
Block	6		-000.0002	000.1044	10	0.00	0.9983
new*Entry		1	000.2991	000.3561	10	0.84	0.4206
new*Entry		2	-000.2461	000.3561	10	-0.69	0.5052
new*Entry		3	-000.0869	000.3561	10	-0.24	0.8122
new*Entry		4	000.0585	000.3561	10	0.16	0.8729
new*Entry		5	-000.0192	000.3561	10	-0.05	0.9581
new*Entry		6	-000.4199	000.3561	10	-1.18	0.2656
new*Entry		7	-000.8584	000.3561	10	-2.41	0.0366
new*Entry		8	-000.6590	000.3561	10	-1.85	0.0939
new*Entry		9	-000.8146	000.3561	10	-2.29	0.0452
new*Entry		10	-000.1153	000.3561	10	-0.32	0.7527
new*Entry		11	000.6102	000.3561	10	1.71	0.1173
new*Entry		12	000.2095	000.3561	10	0.59	0.5693
new*Entry		13	-000.1613	000.3561	10	-0.45	0.6602
new*Entry		14	000.5051	000.3561	10	1.42	0.1865
new*Entry		15	-000.2965	000.3561	10	-0.83	0.4245
new*Entry		16	000.0487	000.3561	10	0.14	0.8940

# Variations – two-way control of heterogeneity

Design	Features/Applications	Reference
Modified Youden Square	Two-way control of heterogeneity	Federer and Raghavarao, 1975
Augmented Row-column	Entries adjusted for adjacent checks; requires many checks	Federer, Nair and Raghavarao, 1975
Modified Augmented (MAD Type 1)	Systematic placement of controls; Requires square plots	Lin and Poushinsky, 1983
Modified Augmented (MAD Type 2)	Systematic placement of controls; Long, rectangular plots	Lin and Poushinsky, 1985



# More Variations

Design	Features/Applications	Reference
<b><i>Factorial treatments</i></b>		
Augmented Split Block	Intercropping, stress adaptation	Federer, 2005
<b><i>Incomplete Blocks</i></b>		
Augmented Lattice Square	Accommodates larger number of checks	Federer, 2002
$\alpha$ - $\alpha$ Lattice	Accommodates larger number of checks	Williams and John, 2003
Augmented p-rep (partially replicated)	Replicates of entries used to make block adjustments and estimate error	Williams, Piepho, and Whitaker, 2011
<b><i>Multiple locations</i></b>		
RCBD, ICBD, row-column, etc.	SAS analysis provided	Federer, Reynolds, and Crossa, 2001

# Multiple Locations – Augmented or Lattice Design?

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## ■ Lattice Designs

- Sites are complete replications
- Information from all plots are used to adjust means for field effects
  - May have greater precision and power to detect differences among entries

## ■ Augmented Designs

- Flexible arrangement of new entries in field plans
- Estimate of experimental error obtained from each location
  - Assess site quality
  - Evaluate GXE interactions
  - More flexibility in combining information across sites

# Software for Augmented Designs

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- **AGROBASE GenII** by Agronomix, Inc.
  - Randomize and analyze modified augmented type-2 designs
- CIMMYT – sas macro called **UNREPLICATE**  
<http://apps.cimmyt.org/english/wps/biometrics/>
  - Developed in 2000 – uses some older SAS syntax
- Statistical Package for Augmented Designs (**SPAD**) – Indian Agricultural Research Institute, New Delhi  
[www.iasri.res.in/iasrisoftware/SPAD.ppt](http://www.iasri.res.in/iasrisoftware/SPAD.ppt)  
<http://www.iasri.res.in/design/Augmented%20Designs/home.htm>

# References

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- Full list of references provided as supplement
- Federer, W.T. 1961. Augmented designs with one-way elimination of heterogeneity. *Biometrics* 17(3): 447-473
- Wolfinger, R.D., W.T. Federer, and O. Cordero-Brana. 1997. SAS PROC GLM and PROC MIXED for recovering blocking and variety information in augmented designs. *Agronomy Journal* 89: 856-859

# Acknowledgements

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  - Alicia Wilson



# Questions?

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