

# INCOMPLETE SPLIT-PLOT DESIGNS BASED ON $\alpha$ -DESIGNS

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A COMPROMISE BETWEEN TRADITIONAL SPLIT-PLOT DESIGNS AND RANDOMISED COMPLETE BLOCK DESIGNS

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# Introduction

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- > **Example of trial to be performed**
- > 2-factorial design
  - > Treatment factor 1 with few levels (e.g.  $\pm$  Herbicides)
  - > Treatment factor 2 with many levels (e.g. a large number of varieties)
- > **Some possible designs**
- > Split-plot
- > Randomised complete block designs
- > Incomplete split-plot

# What is an incomplete split-plot

## Small example: $\pm$ Herbicide, 9 varieties

Randomised complete block design

<u>9</u>	<u>5</u>	2	<u>8</u>	9	5	4	<u>6</u>	<u>4</u>	1	<u>1</u>	8	<u>7</u>	3	<u>2</u>	7	<u>3</u>	6
<u>5</u>	7	2	9	4	<u>4</u>	8	3	5	6	<u>7</u>	<u>1</u>	<u>6</u>	<u>2</u>	<u>8</u>	1	<u>3</u>	<u>9</u>

Traditional split-plot

2	9	5	8	1	4	7	6	3	<u>9</u>	<u>8</u>	<u>5</u>	<u>6</u>	<u>4</u>	<u>1</u>	<u>2</u>	<u>3</u>	<u>7</u>
<u>7</u>	<u>2</u>	<u>9</u>	<u>3</u>	<u>8</u>	<u>4</u>	<u>1</u>	<u>6</u>	<u>5</u>	7	1	6	5	2	8	9	3	4

Incomplete split-plot

8	2	5	<u>8</u>	<u>2</u>	<u>5</u>	1	4	7	<u>7</u>	<u>4</u>	<u>1</u>	<u>3</u>	<u>6</u>	<u>9</u>	9	6	3
<u>9</u>	<u>7</u>	<u>8</u>	7	8	9	<u>4</u>	<u>6</u>	<u>5</u>	6	5	4	2	1	3	<u>2</u>	<u>1</u>	<u>3</u>

# Construction

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- › Can be based on different types of incomplete block designs
  - › Closed to use to use  $\alpha$ -designs (generalised lattice)
- ›  $\alpha$ -designs (generalized lattice)
  - › Resolvable
  - › Flexible
    - › Available for almost any number of varieties ( $v > 2$ )
    - › Wide range of replications ( $2 \leq k \leq 10$ )
    - › Broad range of block sizes (and  $v$  does not need to be a multiple of  $k$ )

# Construction – 4 different methods

$v=12, r=2$  (in A and B) or 4 (in C and D),  $s=4, k=3$ .

Design method	Rep	Layout in the “field”							
A	1	D H L	C G K	c g k	b f j	a e i	A E I	B F J	d h l
	2	c h i	C H I	d e j	B G L	A F K	b g l	D E J	a f k
B	1	B F J	b f j	A E I	a e i	d l h	D L H	C G K	c g k
	2	d e j	D E J	C H I	c h i	a f k	A F K	B G L	b g l
C	1	D H L	C G K	c e l	b h k	a g j	A E I	B F J	d f i
	2	c f l	C H I	a h j	B G L	A F K	b e k	D E J	d g i
D	1	B F J	b h k	a g j	A E I	D H L	d f i	c e l	C G K
	2	d g i	D E J	C H I	c f l	A F K	a h j	B G L	b e k

# Analyses

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$$Y_{rtv} = \mu + \delta_v + \tau_t + (\delta\tau)_{vt} + A_r + E_{rtv}$$

RCB

$$Y_{rtv} = \mu + \delta_v + \tau_t + (\delta\tau)_{vt} + A_r + C_{rt} + E_{rtv}$$

Split-plot

$$Y_{rbtv} = \mu + \delta_v + \tau_t + (\delta\tau)_{vt} + A_r + D_{rb} + E_{rbtv}$$

A and C

$$Y_{rgbtv} = \mu + \delta_v + \tau_t + (\delta\tau)_{vt} + A_r + C_{rg} + D_{rgb} + E_{rgbtv}$$

B and D

# Intra-block efficiencies

Trial definition				Efficiency of $\alpha$ -design used for construction <sup>1</sup>		Efficiency of variety comparisons		Efficiency of F-test for the hypothesis of no interaction	
Replicates <i>r</i>	Treatments <i>t</i>	Varieties <i>v</i>	Blocks <i>s</i>	Method A&B	Method C&D	Method A&B	Method C&D	Method A&B	Method C&D
2	2	80	10	0.806	0.864	0.806	0.806	0.886	0.848
2	2	40	5	0.823	0.876	0.823	0.823	0.897	0.863
2	2	48	6	0.817	0.872	0.817	0.817	0.894	0.858
2	2	48	8	0.753	0.826	0.753	0.752	0.851	0.801
3	2	35	5	0.845	0.872	0.845	0.844	0.882	0.843

<sup>1</sup>) Also efficiency factor for varieties if no interaction effect are included in the model

- > If no interaction effect then method C and D are better than method A and B
- > If test for interaction is important then method A and B are better than method C and D
- > If variation between group of blocks are larger than between blocks then
  - > Method B is better than method A and method D is better than method C

The projects, in which the designs were used, expected interaction between the two factors and were very interested in testing for interaction, so method B were chosen

# Incomplete split-plot

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- › Practical compromise
  - › Easier than RCB, more difficult than split-plot
  - › May require more guard-area than split-plot, but less than RCB
- › Efficiency
  - › Herbicides: more whole plots, comparison within pair (group) of incomplete block and thus moderate distance between incomplete “whole-plots”: More efficient than split-plot
  - › Varieties: few plots within each incomplete “whole plot” and thus small distance between sub-plots: More efficient than RCB and split-plot

# Performed experiments

Trial no	Number of				
	Repli- cates <i>r</i>	Treat- ments <i>t</i>	Varie- ties <i>v</i>	Blocks <i>s</i>	Plots per block <i>k</i>
A	2	2	48	6	8
B	2	2	48	8	6
C	2	2	48	6	8
D	3	2	35	5	7
E	3	2	8	2	4

# Field layout of trial E

Each plot is  
1.5 m × 11.0 m  
Each block is  
12.0 m × 11.0 m

Replicate 1						Replicate 2					
<u>45</u>	<u>03</u>	42	34	17	<u>14</u>	38	<u>42</u>	36	<u>29</u>	<u>31</u>	16
<u>20</u>	<u>25</u>	08	27	30	<u>31</u>	08	<u>25</u>	24	<u>14</u>	<u>02</u>	23
<u>35</u>	<u>04</u>	46	32	29	<u>15</u>	45	<u>46</u>	17	<u>18</u>	<u>12</u>	44
<u>18</u>	<u>16</u>	12	19	43	<u>09</u>	22	<u>09</u>	10	<u>41</u>	<u>11</u>	01
<u>24</u>	<u>37</u>	47	05	02	<u>48</u>	40	<u>21</u>	27	<u>05</u>	<u>20</u>	48
<u>21</u>	<u>10</u>	06	41	28	<u>07</u>	28	<u>19</u>	03	<u>37</u>	<u>32</u>	47
<u>33</u>	<u>23</u>	13	38	22	<u>01</u>	04	<u>15</u>	33	<u>43</u>	<u>34</u>	26
<u>40</u>	<u>11</u>	39	44	26	<u>36</u>	07	<u>30</u>	13	<u>39</u>	<u>06</u>	35
35	16	<u>47</u>	<u>19</u>	<u>17</u>	14	<u>40</u>	25	<u>03</u>	37	34	<u>35</u>
24	37	<u>08</u>	<u>44</u>	<u>28</u>	31	<u>45</u>	15	<u>27</u>	29	20	<u>01</u>
18	10	<u>39</u>	<u>32</u>	<u>43</u>	15	<u>08</u>	09	<u>13</u>	39	12	<u>26</u>
20	04	<u>13</u>	<u>05</u>	<u>22</u>	09	<u>07</u>	19	<u>10</u>	14	32	<u>48</u>
40	25	<u>06</u>	<u>38</u>	<u>29</u>	01	<u>28</u>	21	<u>17</u>	41	11	<u>16</u>
45	23	<u>46</u>	<u>34</u>	<u>30</u>	36	<u>38</u>	42	<u>36</u>	18	06	<u>44</u>
33	03	<u>42</u>	<u>41</u>	<u>26</u>	48	<u>04</u>	46	<u>33</u>	43	02	<u>47</u>
21	11	<u>12</u>	<u>27</u>	<u>02</u>	07	<u>22</u>	30	<u>24</u>	05	31	<u>23</u>

# Field layout of trial E

Each plot is  
2.5 m × 12.5 m  
Each block is  
10.0 m × 12.5 m

Rep 3	Variety	7.0	1.0	3.0	6.0	3.1	6.1	7.1	1.1
	plotno	6	7	18	19	30	31	42	43
Rep 2	Variety	8.1	2.1	4.1	5.1	5.0	2.0	8.0	4.0
	plotno	5	8	17	20	29	32	41	44
Rep 1	Variety	7.1	6.1	5.1	8.1	7.0	8.0	6.0	5.0
	plotno	4	9	16	21	28	33	40	45
Rep 1	Variety	2.0	4.0	3.0	1.0	2.1	4.1	3.1	1.1
	plotno	3	10	15	22	27	34	39	46
Rep 1	Variety	4.0	1.0	7.0	5.0	4.1	1.1	7.1	5.1
	plotno	2	11	14	23	26	35	38	47
Rep 1	Variety	3.0	8.0	6.0	2.0	8.1	6.1	3.1	2.1
	plotno	1	12	13	24	25	36	37	48

# Efficiency of the designs, Yield

Trial no.	Design <sup>1</sup> : Comparison	Grain yield, hkg ha <sup>-1</sup>				
		LSD			Relative reduction, %	
		Plan	Split-plot	RCB	Split plot	RCB
A	Variety:Treat	6.5	7.9	8.04	18	19
	Treat:Variety	6.7	10.1	8.04	34	17
	Treat	2.2	11.3	1.16	81	-90
	Variety	4.6	5.6	5.69	18	19
B	Variety:Treat	8.4	8.5	9.19	1	9
	Treat:Variety	8.4	16.9	9.19	50	9
	Treat	1.7	23.2	1.33	93	-28
	Variety	6.2	6.0	6.50	-3	5
C	Variety:Treat	4.9	7.9	8.92	38	45
	Treat:Variety	5.3	19.7	8.92	73	41
	Treat	2.4	27.0	1.29	91	-86
	Variety	3.5	5.6	6.31	37	45
D	Variety:Treat	2.0	2.8	2.86	29	30
	Treat:Variety	2.2	3.1	2.86	29	23
	Treat	1.0	1.7	0.48	41	-108
	Variety	1.4	2.0	2.02	30	31
E	Variety:Treat	5.8	5.7	7.4	-2	21
	Treat:Variety	5.9	7.6	7.4	23	21
	Treat	3.7	7.8	2.6	53	-40
	Variety	4.2	4.1	5.2	-4	19

# Efficiency of the designs, %Mildew

Trial no.	Design <sup>1</sup> : Comparison	Powdery mildew severity, transformed %				
		LSD			Relative reduction, %	
		Plan	Split-plot	RCB	Split plot	RCB
A	Variety:Treat	0.609	0.625	0.625	3	3
	Treat:Variety	0.609	0.695	0.625	12	3
	Treat	0.086	0.577	0.090	85	4
	Variety	0.440	0.442	0.442	0	0
C	Variety:Treat	0.950	1.032	1.032	7	7
	Treat:Variety	0.970	1.149	1.032	16	6
	Treat	0.263	0.954	0.149	72	-77
	Variety	0.672	0.730	0.730	8	8
D	Variety:Treat	0.437	0.437	0.438	0	0
	Treat:Variety	0.437	0.455	0.438	4	0
	Treat	0.074	0.175	0.074	58	0
	Variety	0.309	0.309	0.310	0	0

# Efficiency of the designs, other variables

Variable	Design: Comparison	LSD			Relative reduction, %	
		Plan	Split-plot R	RCB	Split-plot R	RCB
Wheat June 19 g 0.25m <sup>-2</sup>	Variety:Herbicide	161	167	195	3	17
	Herbicide:Variety	174	210	195	17	11
	Herbicide	119	204	69	42	-73
	Variety	115	118	138	3	17
Log Weed June 19 Log g 0.25m <sup>-2</sup>	Variety:Herbicide	2.8	2.8	2.8	1	0
	Herbicide:Variety	2.8	2.9	2.8	4	0
	Herbicide	1.0	2.1	1.0	53	0
	Variety	2.0	2.0	2.0	1	0
Vegetation cover (%) June 5	Variety:Herbicide	.052	.055	.077	6	33
	Herbicide:Variety	.055	.057	.077	3	30
	Herbicide	.032	.041	.027	21	-17
	Variety	.036	.039	.055	6	33
LAI June 21	Variety:Herbicide	0.59	0.60	0.59	2	1
	Herbicide:Variety	0.59	0.62	0.59	4	0
	Herbicide	0.26	0.44	0.21	41	-24
	Variety	0.42	0.42	0.42	2	1
MTA June 21	Variety:Herbicide	5.7	5.8	5.8	0	2
	Herbicide:Variety	5.7	6.0	5.8	4	2
	Herbicide	2.0	4.3	2.1	53	2
	Variety	4.1	4.1	4.1	0	2
Log DIFN June 21	Variety:Herbicide	0.52	0.52	0.53	-0	1
	Herbicide:Variety	0.53	0.57	0.53	7	0
	Herbicide	0.26	0.45	0.19	43	-38
	Variety	0.37	0.37	0.38	-1	1

# Discussion and conclusions

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- › **Efficiency**
- › Compared to randomised complete block design
  - › Incomplete split-plot were most often less efficient when comparing the main effect of treatments
    - › Larger number of independent plots in RCB
  - › Incomplete split-plot most often more efficient for other comparisons
    - › Larger block size in RCB
- › Compared to traditional split-plot
  - › Incomplete split-plot were most often more efficient for all types of comparisons
    - › Especially for comparing treatment means (many more degrees of freedom and smaller blocks)

# Discussion and conclusions

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- › Increase in efficiency
  - › In most cases larger for grain yield than for mildew
    - › Probably because mildew is less sensible to soil variation
  - › Small for trial E when comparing mean of varieties and varieties within treatment
    - › Relative small reduction in block sizes
  - › Small for trial B when comparing mean of varieties and varieties within treatment
    - › Reason unknown

# Discussion and conclusions

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- › Practical considerations
  - › Treatment applications
    - › Easier than randomised complete block design
    - › More difficult than split-plot design
  - › Guard areas
    - › Less than for randomised complete block design
    - › More than for split-plot design
- › Design and statistical analysis
  - › More complex than both randomised complete block design and split-plot design
  - › Appropriate software are available and with today's computer power this should not be a problem