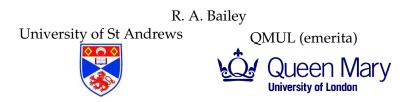
Block designs with very low replication, and other challenges in design of experiments



James Hutton Institute, 30 March 2015

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In the last 10 years, Cullis and colleagues in Australia (and independently Bueno and Gilmour) have suggested replacing many occurrences of the the control by double replicates of a small number of new varieties: for example, 24 new varieties with one plot each, 6 new varieties with two plots each, and the control on two further plots. In breeding trials of new varieties, typically there is very little seed of each of the new varieties.

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This is an improvement if there are no blocks.

How do we allow for variation between the plots?

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If field operations have been primarily in one direction for a long time, then it is reasonable to divide the fields into blocks whose length runs along that direction.

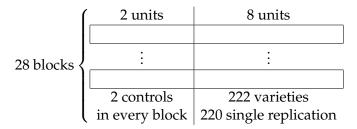
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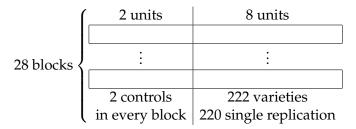
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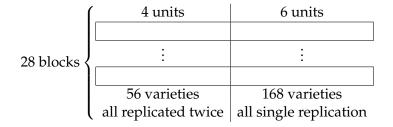
Even more extreme: 2 uninteresting controls in each block.

Two possible designs for 224 varieties in 28 blocks of 10

28 blocks {	2 units	8 units
	÷	÷
	2 controls	222 varieties
	in every block	220 single replication

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We are given b blocks of size k. We are given v varieties. Assume that

average replication
$$= \bar{r} = \frac{bk}{v} \le 2.$$

How should we allocate varieties to blocks?

A-optimal designs

We measure the response *Y* on the plot with variety *i* in block *D*, and assume that

 $Y = \tau_i + \beta_D +$ random noise,

where the random noise is $N(0, \sigma^2)$, independently for each plot.

Put

 $V_{ij} \sigma^2 = variance of the best linear unbiased estimator for <math>\tau_i - \tau_j$;

 $V_T = \sum_{i=1}^{v-1} \sum_{j=i+1}^{v} V_{ij} \quad \propto \quad \text{sum of variances of variety differences.}$

A block design is A-optimal if it minimizes V_T .

Definition Call a variety a a drone if it has replication 1; Definition Call a variety a a drone if it has replication 1; a queen-bee if it occurs in every block; Definition Call a variety a a drone if it has replication 1; a queen-bee if it occurs in every block; a worker otherwise. Definition Call a variety a a drone if it has replication 1; a queen-bee if it occurs in every block; a worker otherwise. Definition Call a variety a a drone if it has replication 1; a queen-bee if it occurs in every block; a worker otherwise.

Is it better to put all the drones into one block (or a few blocks), or are they better distributed equally among all the blocks?

Block ABlock Bn dronesm drones

If we move all the drones in block *B* into block *A* then we reduce *nm* variances from $2 + V_{AB}$ to 2, where V_{AB} is the variance of the estimator of the difference between the block effects of *A* and *B* in the design obtained by ignoring the drones.

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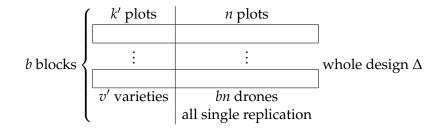
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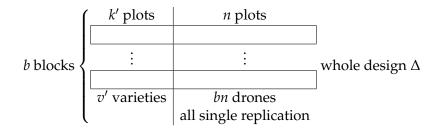
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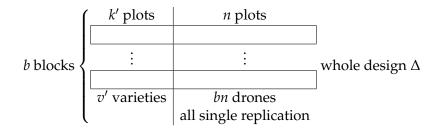
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Then we have to remove *m* non-drones from block *A*, and this increases the variances between these n + m drones and the remaining v - n - m varieties. This more than compensates for the original reduction in variance.

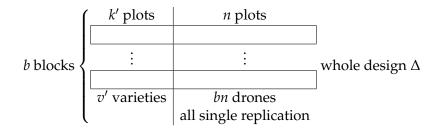




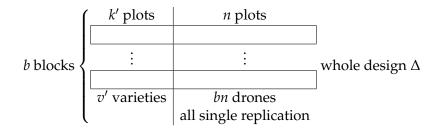
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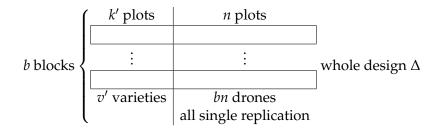


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$$n \ge n_0 = \left\lfloor \frac{2v - bk}{b} \right\rfloor \qquad k' \le k_0 = k - n_0$$

Theorem (cf. Herzberg and Jarrett, 2007) If there are n drones in each block of Δ , and the core design Γ has v' varieties in b blocks of size k' then the sum of the variances of variety differences in Δ

$$= V_T(\Delta) = bn(bn + v' - 1) + V_T(\Gamma) + nV_{BT}(\Gamma) + n^2V_B(\Gamma),$$

where

- $V_T(\Gamma) = the sum of the variances of variety differences in \Gamma$
- $V_B(\Gamma) = the sum of the variances of block differences in \Gamma$
- $V_{BT}(\Gamma)$ = the sum of the variances of sums of one treatment and one block in Γ .

Sum of variances in whole design if Γ is equi-replicate

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so $V_B(\Gamma)$ and $V_{BT}(\Gamma)$ are both increasing functions of $V_T(\Gamma)$.

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so $V_B(\Gamma)$ and $V_{BT}(\Gamma)$ are both increasing functions of $V_T(\Gamma)$. Consequence

For a given choice of k', use the core design Γ which minimizes $V_T(\Gamma)$.

Sum of variances in whole design if there are many drones

$$V_T(\Delta) = bn(bn + v' - 1) + V_T(\Gamma) + nV_{BT}(\Gamma) + n^2 V_B(\Gamma)$$

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Consequence

If v is large then n is large, so we need to focus on reducing $V_B(\Gamma)$, so it may be best to increase the number of drones and decrease k' (the size of blocks in the core design Γ), so that average replication within Γ is more than 2. If there are 4(2 + n) varieties in 4 blocks of size 4 + n, the design on the left is A-better than the design on the right if and only if n < 50.

1 2 3 4	<i>n</i> drones	1 2 3	n+1 drones
1 2 5 6	<i>n</i> drones	1 2 4	n+1 drones
3 6 7 8	<i>n</i> drones	1 3 4	n+1 drones
4 5 7 8	<i>n</i> drones	2 3 4	n+1 drones

Theorem

Suppose that we are given b blocks of size k, and v varieties. For i = 1, 2, let design Δ_i have core subdesign Γ_i with block size k_i . If Γ_1 is the dual of a balanced incomplete block design and $k_1 > k_2$ then Δ_2 is worse than Δ_1 on the A criterion, no matter how big v is. If there are 4n + 6 varieties in 4 blocks of size 3 + n, the design on the left is A-better than the design on the right, for all values of *n*.

1	2	3	<i>n</i> drones	1	2	n+1 drones
1	4	5	<i>n</i> drones	1	2	n+1 drones
2	4	6	<i>n</i> drones	1	2	n+1 drones
3	5	6	<i>n</i> drones	1	2	n+1 drones

Given *b*, *v* and *k*, how do we find an A-optimal design for *v* varieties in *b* blocks of size *k* when

$$\frac{bk}{2} \le v \le b(k-1) + 1?$$

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Average replication ≤ 2 Maximum *v* for estimability

Case 1. b = 2 or b = 3 (very small b).

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$$k_0 = k - \left\lfloor \frac{2v - bk}{b} \right\rfloor$$
 = biggest space per block for non-drones.

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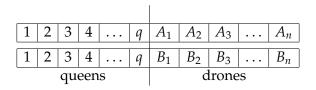
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Case 3. $k_0 \ge b - 1$.
Case 4. $2 < k_0 < b - 1$ (small k_0 but not Case 2).

$$k_0 = k - \left\lfloor \frac{2v - bk}{b} \right\rfloor$$
 = biggest space per block for non-drones.

Morgan and Jin (2007) showed that the A-optimal designs are those with 2n drones and q queen bees, where $n = n_0 = v - k$ and $q = k' = k_0 = k - n_0 = 2k - v$.



Case 1 continued. 3 blocks of size k

Using the nice theorem, RAB has shown that the A-optimal designs are as follows when v is divisible by 3 (and presumably small changes deal with the other cases). There are 3w workers and 3n drones,

where 3w = 3k - v and $n = n_0 = k - 2w$ and $k' = k_0 = 2w$.

1	2	4	5	•••	3w - 2	3w - 1	A_1	A_2	A_3	•••	A_n
1	3	4	6	•••	3 <i>w</i> −2	3w	<i>B</i> ₁	<i>B</i> ₂	<i>B</i> ₃	•••	B _n
2	3	5	6		3w - 1	3w	<i>C</i> ₁	<i>C</i> ₂	<i>C</i> ₃		C_n
w copies of design using all pairs from 3							Ċ	lrone	es		

Case 2. v = b(k-1) + 1

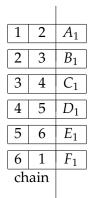
This is the maximum number of varieties that can be tested in b blocks of size k with all comparisons estimable.

Mandal, Shah and Sinha (1991), for k = 2, and Bailey and Cameron (2013), for general block size, showed that, no matter how many blocks there are, the A-optimal design has the following form.

1	A_1	A_2	A_3		A_{k-1}	
1	<i>B</i> ₁	<i>B</i> ₂	<i>B</i> ₃	•••	B_{k-1}	
1	<i>C</i> ₁	<i>C</i> ₂	<i>C</i> ₃		C_{k-1}	
1	D_1	D_2	<i>D</i> ₃	•••	D_{k-1}	
1	E_1	E_2	<i>E</i> ₃		E_{k-1}	
1 queen	v-1 drones					

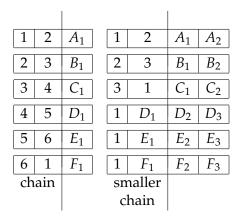
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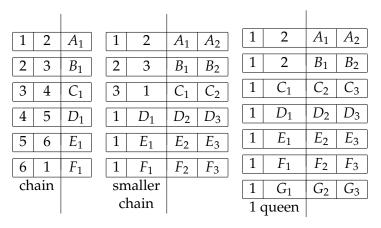
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small *k* and *b* increase *k* if $b \ge 5$



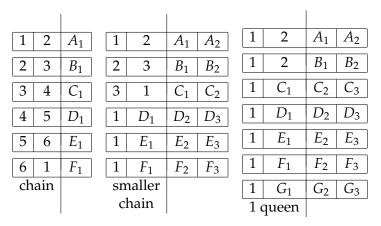
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small *k* and *b* increase *k* if $b \ge 5$ then increase *b*



Youden and Connor (1953) had recommended chain designs.

For simplicity, assume that b divides 2v, so that

$$n_0 = \frac{2v - bk}{b} =$$
 minimum number of drones per block.

Then

$$\frac{b(2k-b+1)}{2} \ge v \ge \frac{bk}{2} \ge \frac{b(b-1)}{2}.$$

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Then

$$\frac{b(2k-b+1)}{2} \ge v \ge \frac{bk}{2} \ge \frac{b(b-1)}{2}.$$

Let Γ_0 be the design for b(b-1)/2 varieties replicated twice in *b* blocks of size b-1in such a way that there is one variety in common to each pair of blocks. This is A-optimal for these numbers.

If $k_0 = s(b-1)$ then take Γ to be *s* copies of Γ_0 . This is always *A*-optimal.

If $k_0 > b - 1$ but k_0 is not a multiple of b - 1, then the following strategy seems likely to be good (but it is not A-optimal when $b = k_0 = 4$ and v is very large).

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Construction Method

1. put n_0 drones in each block;

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Construction Method

- 1. put n_0 drones in each block;
- 2. put in one copy of Γ_0 ;

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- 1. put n_0 drones in each block;
- 2. put in one copy of Γ_0 ;
- 3. *put in as many further copies of* Γ_0 *as possible;*

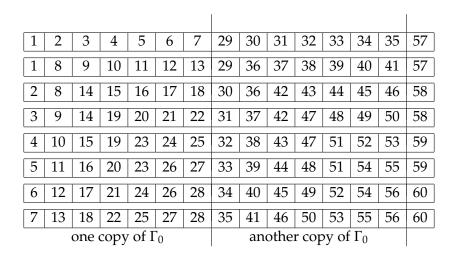
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Construction Method

- 1. put n_0 drones in each block;
- 2. put in one copy of Γ_0 ;
- 3. *put in as many further copies of* Γ_0 *as possible;*
- 4. in any remaining space, use a good design for workers with replication 2 (so long as there is at least one copy of Γ₀, it probably doesn't make much difference which one is used).

Case 3. Example: b = 8 and k = 15 (so $60 \le v \le 92$)

60 varieties: all workers ($n_0 = 0$)



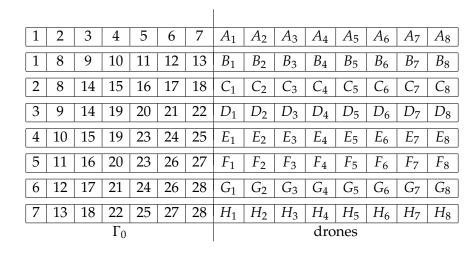
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76 varieties: 44 workers, 32 drones ($n_0 = 4$)

1	2	3	4	5	6	7	29	30	31	32	A_1	A_2	A_3	A_4
1	8	9	10	11	12	13	33	34	35	36	<i>B</i> ₁	<i>B</i> ₂	<i>B</i> ₃	<i>B</i> ₄
2	8	14	15	16	17	18	37	38	39	40	C_1	<i>C</i> ₂	<i>C</i> ₃	<i>C</i> ₄
3	9	14	19	20	21	22	41	42	43	44	D_1	<i>D</i> ₂	<i>D</i> ₃	D_4
4	10	15	19	23	24	25	29	33	37	41	E_1	<i>E</i> ₂	E_3	<i>E</i> ₄
5	11	16	20	23	26	27	30	34	38	42	F_1	<i>F</i> ₂	F_3	F_4
6	12	17	21	24	26	28	31	35	39	43	G_1	<i>G</i> ₂	G ₃	<i>G</i> ₄
7	13	18	22	25	27	28	32	36	40	44	H_1	H_2	H_3	H_4
			Γ					6 wc				dro	nes	
							re	plica	ation	2				

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92 varieties: 28 workers, 64 drones ($n_0 = 8$)



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by interchanging the roles of blocks and varieties.)

- $V_T(\Gamma_i)$ = the sum of the variances of variety differences in Γ_i
- $V_B(\Gamma_i)$ = the sum of the variances of block differences in Γ_i
- $V_{BT}(\Gamma_i)$ = the sum of the variances of sums of one treatment and one block in Γ_i .

If there are n_i drones in each block then, in the whole design Δ ,

$$V_T(\Delta) = bn_i(bn_i + v'_i - 1) + V_T(\Gamma_i) + n_i V_{BT}(\Gamma_i) + n_i^2 V_B(\Gamma_i).$$

For various values of $k_i \le k_0$, find the best core subdesign Γ_i for v'_i varieties in *b* blocks of

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Use this formula to find the core subdesign which gives the smallest $V_T(\Delta)$.

As the number of varieties increases, it becomes more important to choose Γ_i with a small value of $V_B(\Gamma_i)$.

Case 4 continued. $k_0 = 4 < b - 1$, $V_B(\Gamma_i) \div b(b - 1)/2$

Best design for *b* blocks known to RAB

	Γ_1	Γ2	Γ_3	Γ_4
k_i	2	3	3	4
	2 queens,	2 queens,	<i>b</i> workers	2 <i>b</i> workers
	both boring	2 workers (rep 2)	rep 3	rep 2
b=6	1	1-	0.85	0.87
b = 7	1	1-	0.86	0.92
b = 8	1	1-	0.89	0.93
b = 9	1	1-	0.92	
b = 10	1	1-		
b = 11	1	1-		
<i>b</i> = 12	1	1-	0.98	
b = 13	1	1-	1	1.07
b = 14	1	1-		
b = 15	1	1-	1.01	1.08

Case 4 continued. $k_0 = 4 < b - 1$, $V_B(\Gamma_i) \div b(b - 1)/2$

Best design for *b* blocks known to RAB

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		Γ_1	Γ_2	Γ_3	Γ_4
	k_i	2	3	3	4
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		both boring	2 workers (rep 2)	rep 3	rep 2
l	v = 6	1	1-	0.85	0.87
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As *v* increases, Γ_3 becomes better than Γ_4 .

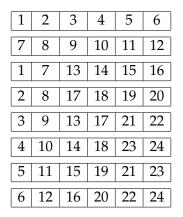
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As *v* increases, Γ_3 becomes better than Γ_4 . If $b \ge 14$, then, as *v* increases, Γ_1 and Γ_2 become better than Γ_3 . Case 4 continued. $2 < k_0 < b - 1$ when b = 8: $k_0 = 6$

 $k = k_0 = 6$, and 24 varieties, all workers, all replicated twice.



(One worker for each pair of blocks except for $\{A, B\}$, $\{C, D\}$, $\{E, F\}$ and $\{G, H\}$.)

Case 4 continued. k = 5 and k = 6 when b = 8: $k_0 = 5$

k = 5k = 620 varieties: 28 varieties: 20 workers, no drones 20 workers, 8 drones A_1 B_1 C_1 D_1 E_1 F_1 G_1 H_1

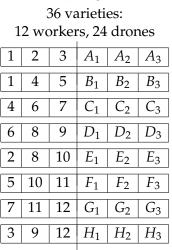
Case 4 continued. k = 5 and k = 6 when b = 8: $k_0 = 4$

k = 5k = 624 varieties: 32 varieties: 16 workers, 8 drones 8 workers, 24 drones 2 3 4 A_1 1 2 4 A_1 A_2 A_3 5 B_1 5 B_2 B_3 6 7 8 2 3 B_1 9 10 11 12 C_1 3 4 6 C_1 C_2 C_3 13 14 15 16 D_1 4 5 7 D_1 D_2 D_3 5 9 13 E_1 5 8 E_1 E_2 E_3 1 6 \overline{F}_3 2 10 14 F_1 7 1 F_1 F_2 6 6 3 11 15 G_1 8 2 G_1 G_2 7 7 G₃ 12 16 H_1 3 H_1 H_2 4 8 8 1 H_3 k'=4k'=3rep = 2rep 3

Case 4 continued. k = 5 and k = 6 when b = 8: $k_0 = 3$

k = 528 varieties: 12 workers, 16 drones 3 A_1 2 A_2 B_1 B_2 1 4 5 7 4 6 C_1 C_2 8 9 D_1 D_2 6 \overline{E}_2 2 8 10 E_1 5 10 11 F_1 F_2 7 11 12 G_1 G_2 3 9 12 H_1 H_2

k = 6



The overall message is that there can be phase changes as the spare capacity for replication (bk - v) decreases. Therefore it is necessary to compare core subdesigns Γ_i with different block size k_i . The overall message is that there can be phase changes as the spare capacity for replication (bk - v) decreases. Therefore it is necessary to compare core subdesigns Γ_i with different block size k_i .

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This is work is progress, not a finished project.

- Replication.
- ► Blocking.
- Randomization.

- Replication.
- Blocking.
- Randomization.

What is old? What is new? In 2012 the UK's Food and Environment Research Agency conducted an experiment to find out "the effects of neonicotinoid seed treatments on bumble bee colonies under field conditions" (from a DEFRA report available on the web, Crown copyright 2013).

fera.co.uk/ccss/documents/defraBumbleBeereportPS2371V4A.pdf

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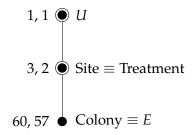
Site	Treatment of oilseed rape seeds
Site A, near Lincoln	no treatment
Site B, near York	Modesto TM
Site C, near Scunthorpe	Chinook TM

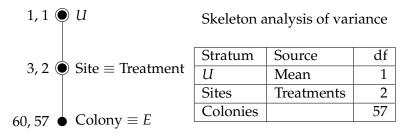
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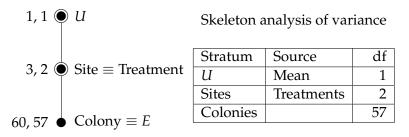
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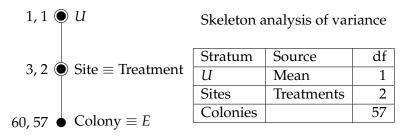
Twenty colonies of bumble bees were placed at each site. Various outcomes were measured on each colony.



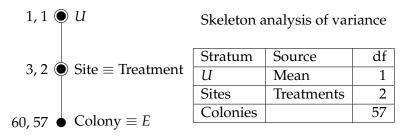




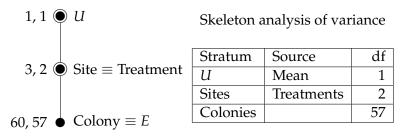
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The Hasse diagram can clearly show such false replication before the experiment is carried out.

False replication is one of the oldest and most common mistakes in design of experiments.

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Why is it still happening in experiments undertaken or commissioned by publicly funded bodies?

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For example, Ching-Shui Cheng discussed supersaturated designs (for *m* 2-level factors in *n* experimental units, with n < m). One of the classical surrogates for model identifability is equal replication ("balance") of the levels of each factor. He showed that this is not a good guide when the models have high dimension.

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and use the blocks in both the design and the analysis, in order to remove bias from the estimates of treatment differences AND from the estimate of experimental error in order to increase power.

We know how to do this for several systems of blocks, under orthogonality.

In experiments in human-computer interaction,

it is common to ask each participant to undertake a certain task under several different scenarios.

With four scenarios, the experiment might use 20 participants once each on four days, and assign scenarios using Latin squares.

A	B	C	D	A	В	С	D	Α	В	C	D	A	В	C	D	Α	В	C	D
В	C	D	Α	В	С	D	A	В	С	D	Α	В	С	D	A	В	С	D	Α
С	D	A	В	С	D	Α	В	С	D	A	В	С	D	A	В	С	D	A	В
D	A	В	С	D	A	В	С	D	A	В	С	D	A	В	C	D	A	В	C

Blocking: the old. III

Skeleton analysis of variance

Stratum	Source	df
U	Mean	1
Days		3
Participants		19
Days#Participants	Scenarios	3
	residual	54

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This is wrong on two counts:

- (i) We do not test for differences between days and between participants: we expect such differences and fit them in the model.
- (ii) If we do not remove these differences, we decrease the power for detecting differences between scenarios.

As new technologies are used in experimentation (for example, in genomics) there may be systems of blocks which the experimenters do not recognise because they are not called 'blocks'. How do we ensure that known methods of construction, randomization and data analysis are not lost? As new technologies are used in experimentation (for example, in genomics) there may be systems of blocks which the experimenters do not recognise because they are not called 'blocks'. How do we ensure that known methods of construction, randomization and data analysis are not lost?

In agricultural fields in some countries, it is more realistic to model spatial variation by spatial correlation than by discrete blocks. How should we construct and randomize such experiments, and analyse data from them? In some recent experiments, the experimental units can be thought of as the nodes of a graph, with edges between some nodes. For example, Gerry Humphris of the University of St Andrews is experimenting with non-medical interventions such as sending a text message to known binge drinkers on Friday afternoons. One drinker (a node) may alter his behaviour and thus affect the behaviour of other people that he knows (the nodes joined to him in the graph). How should we construct and randomize such experiments, and analyse data from them? There is a large body of theory about how to randomize experiments in *simple orthogonal block structures* in such a way that estimators of treatment effects and of their variances are both unbiased over the randomization.

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This has not stopped people from developing software for throwing away "undesirable" outcomes of randomization.

- D. T. Bowman: TFPlan: software for restricted randomization in field plot design. *Agronomy Journal* 92, (2000), 1276–1278.
- A talk presented at the Tenth Working Seminar on Statistical Methods in Variety Testing at Będlewo, Poland, in July 2014.

For spatial correlation, and for designs on the nodes of graphs,

- how should we randomize?
- what criterion of validity should we use?