Conflicts between optimality criteria for block designs with low replication

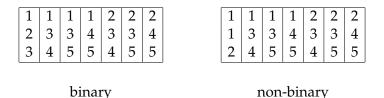
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Lisboa, April 2012

I have v treatments that I want to compare. I have b blocks. Each block has space for k treatments (not necessarily distinct).

How should I choose a block design?



A design is **binary** if no treatment occurs more than once in any block.

1	1	2	3	4	5	6
2	4	5	6	10	11	12
3	7	8	9	13	14	15

1	1	1	1	1	1	1
2	4	6	8	10	12	14
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replications differ by ≤ 1

queen-bee design

The replication of a treatment is its number of occurrences.

A design is a **queen-bee** design if there is a treatment that occurs in every block.

1	2	3	4	5	6	7	
	3						
4	5	6	7	1	2	3	

balanced (2-design)

non-balanced

A binary design is **balanced** if every pair of distinct treaments occurs together in the same number of blocks.

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The concurrence graph *G* of the design has the treatments as vertices.

There are no loops.

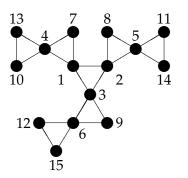
If $i \neq j$ then there are λ_{ij} edges between *i* and *j*.

So the valency d_i of vertex i is

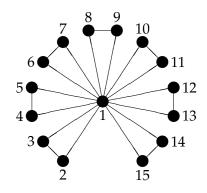
$$d_i = \sum_{j
eq i} \lambda_{ij}.$$

Concurrence graphs of two designs: v = 15, b = 7, k = 3

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Call the remaining eigenvalues *nontrivial*. They are all non-negative.

We measure the response *Y* on each unit in each block.

If that unit has treatment i and block m, then we assume that

$$Y = \tau_i + \beta_m +$$
random noise.

We want to estimate contrasts $\sum_i x_i \tau_i$ with $\sum_i x_i = 0$.

In particular, we want to estimate all the simple differences $\tau_i - \tau_j$.

Put V_{ij} = variance of the best linear unbiased estimator for $\tau_i - \tau_j$.

We want all the V_{ij} to be small.

Assume that all the noise is independent, with variance σ^2 . If $\sum_i x_i = 0$, then the variance of the best linear unbiased estimator of $\sum_i x_i \tau_i$ is equal to

$$(x^{\top}L^{-}x)k\sigma^{2}.$$

In particular, the variance of the best linear unbiased estimator of the simple difference $\tau_i - \tau_j$ is

$$V_{ij} = \left(L_{ii}^- + L_{jj}^- - 2L_{ij}^-\right)k\sigma^2.$$

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Put \bar{V} = average value of the V_{ij} . Then

$$\bar{V} = \frac{2k\sigma^2 \operatorname{Tr}(L^-)}{v-1} = 2k\sigma^2 \times \frac{1}{\operatorname{harmonic mean of } \theta_1, \dots, \theta_{v-1}},$$

where $\theta_1, \ldots, \theta_{v-1}$ are the nontrivial eigenvalues of *L*.

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► E-optimal if minimizes the largest value of $x^{\top}L^{-}x/x^{\top}x$; —equivalently, it maximizes the minimum non-trivial eigenvalue θ_1 of the Laplacian matrix *L*:

If there is a balanced incomplete-block design (BIBD) for v treatments in b blocks of size k, then it is A, D and E-optimal.

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Hence a general idea that

- designs optimal on any of these criteria should be close to balanced
- designs optimal on one of these criteria are not very bad on either of the others.

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So a design is D-optimal if and only if its concurrence graph has the maximal number of spanning trees.

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This is easy to calculate by hand when the graph is sparse.

Electrical networks

We can consider the concurrence graph as an electrical network with a 1-ohm resistance in each edge. Connect a 1-volt battery between vertices *i* and *j*. Current flows in the network, according to these rules.

1. Ohm's Law:

In every edge, voltage drop = current \times resistance = current.

2. Kirchhoff's Voltage Law:

The total voltage drop from one vertex to any other vertex is the same no matter which path we take from one to the other.

3. Kirchhoff's Current Law:

At every vertex which is not connected to the battery, the total current coming in is equal to the total current going out.

Find the total current *I* from *i* to *j*, then use Ohm's Law to define the effective resistance R_{ij} between *i* and *j* as 1/I.

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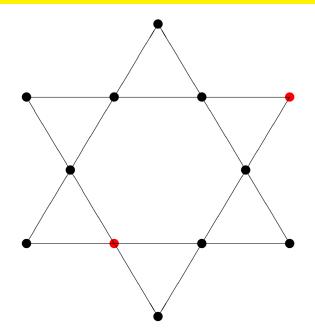
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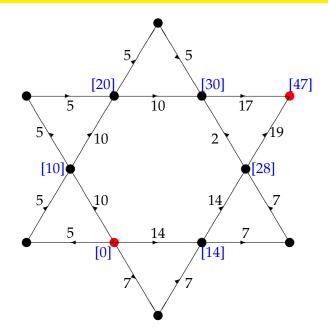
$$V_{ij} = R_{ij} \times k\sigma^2.$$

Effective resistances are easy to calculate without matrix inversion if the graph is sparse.

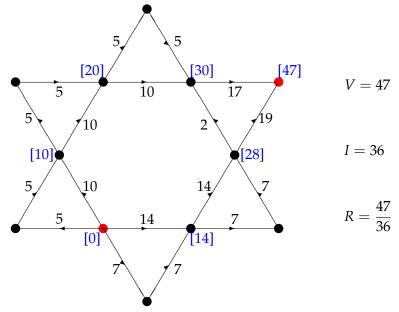
Example calculation



Example calculation



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16/47

Lemma

Let G have an *edge-cutset* of size c (set of c edges whose removal disconnects the graph) whose removal separates the graph into components of sizes m and n. Then

$$\theta_1 \leq c\left(\frac{1}{m} + \frac{1}{n}\right).$$

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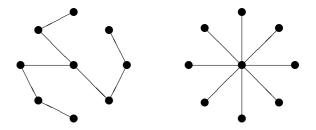
$$\theta_1 \leq c\left(\frac{1}{m} + \frac{1}{n}\right).$$

If *c* is small but *m* and *n* are both large, then θ_1 is small.

Block size 2: least replication

If k = 2 then the design is the same as its concurrence graph, and connectivity requires $b \ge v - 1$.

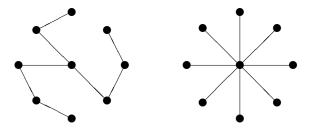
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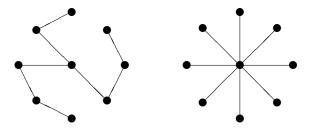


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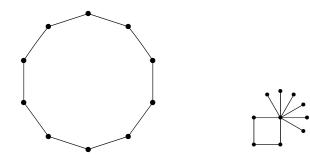
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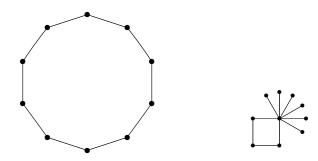
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The only A- or E-optimal designs are the stars.

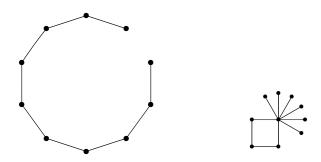
If k = 2 and b = v then the design consists of a cycle with trees attached to some vertices.



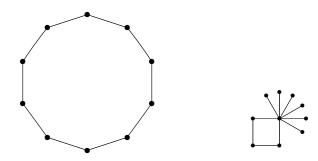
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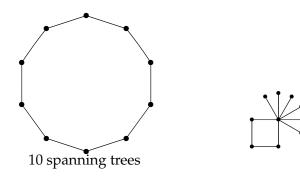
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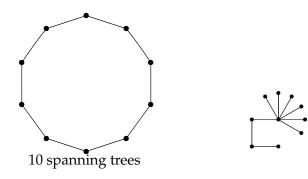
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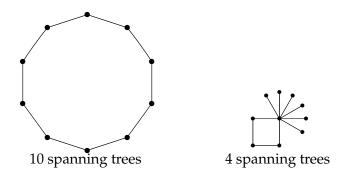
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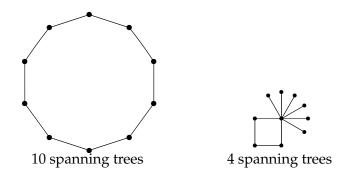


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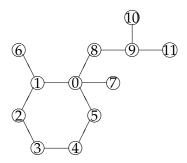
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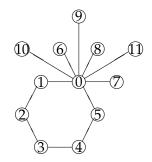
For a spanning tree, remove one edge without disconnecting the graph.



The cycle is uniquely D-optimal when b = v.

Block size 2: one more block: A and E





For a given size of cycle, the total variance is minimized and the smallest non-trivial Laplacian eigenvalue is maximized when everything outside the cycle is attached as a leaf to the same vertex of the cycle.

D-optimal designs	cycle	always
A-optimal designs	cycle square with leaves attached triangle with leaves attached	$if v \le 8$ if $9 \le v \le 12$ if $12 \le v$
E-optimal designs	cycle triangle or digon with leaves	$ \begin{array}{l} \text{if } v \leq 6 \\ \text{if } 6 \leq v \end{array} $

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The A-optimal designs and E-optimal designs are far from equi-replicate.

The change is sudden, not gradual.

An old collaborator, 1980s

"We all know that the A-optimal designs are essentially the same as the D-optimal designs. Surely you've got enough mathematics to prove this?"

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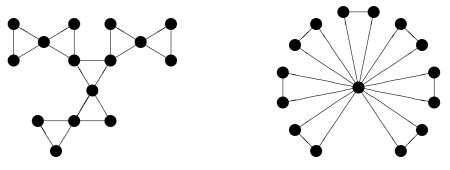
That old collaborator, December 2008

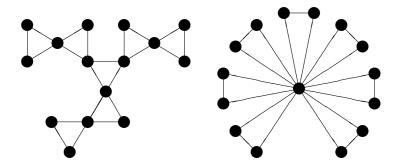
"It seems to be just block size 2 that is a problem."

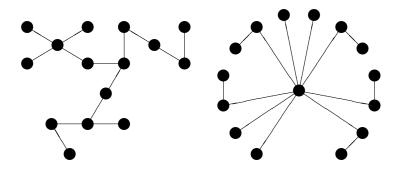
The remaining arguments extend easily to general block size.

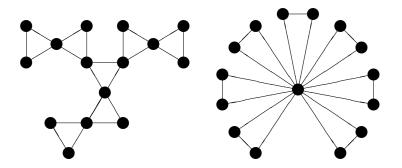
When k = 3, for a connected design, we need $2b \ge v - 1$.

If 2b + 1 = v then all designs are gum-trees, in the sense that there is a unique sequence of blocks from any one treatment to another.



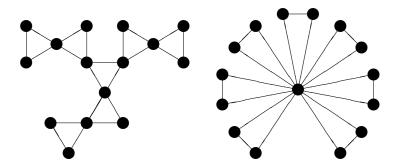






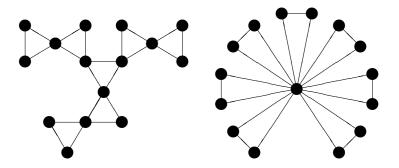
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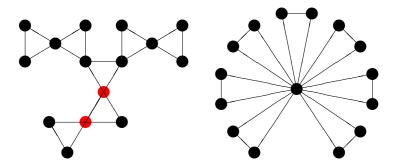


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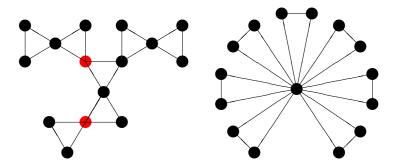
Every gum-tree with b blocks of size 3 has 3^b spanning trees. The D-criterion does not differentiate them.



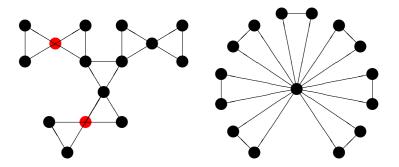
Let R_{ij} be the effective resistance between treatments *i* and *j*.



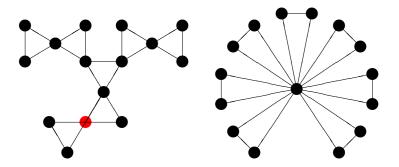
Let R_{ij} be the effective resistance between treatments *i* and *j*. If $i, j \in$ same block then $R_{ij} = \frac{2}{3}$.



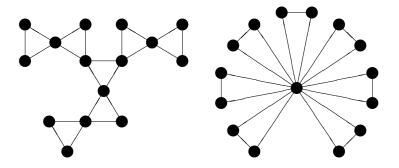
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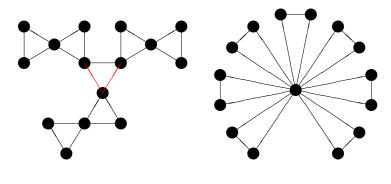


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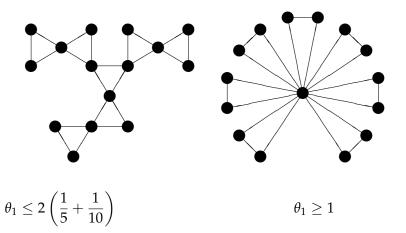
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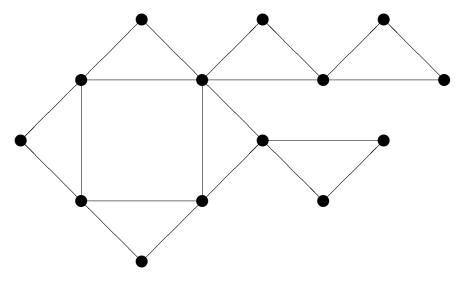
$$\theta_1 \le 2\left(\frac{1}{5} + \frac{1}{10}\right)$$

 $\theta_1 \geq 1$

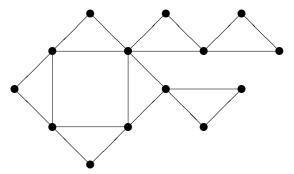


The only E-optimal designs are the queen-bee designs.

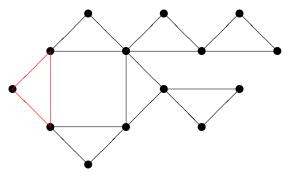
If 2b = v then *G* is a gum-cycle with gum-trees attached.



Suppose that there are *s* blocks in the gum-cycle.



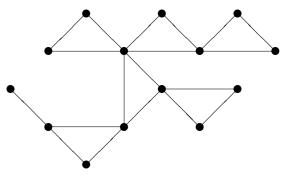
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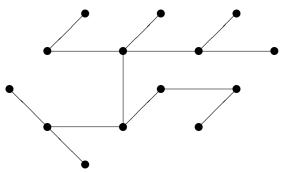
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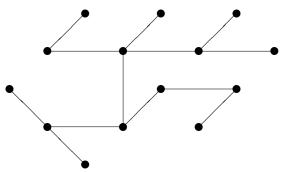
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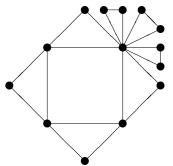


For a spanning tree: choose a block in the gum-cycle sremove its central edge and one other 2 remove an edge from each other block 3^{b-1} There are $2s \times 3^{b-1}$ spanning trees. This is maximized when s = b. This argument extends to all block sizes.

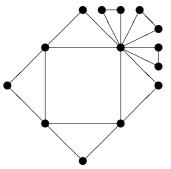
If v = b(k - 1) then the only D-optimal designs are the gum-cycles.

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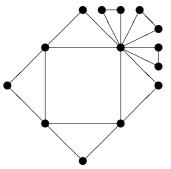


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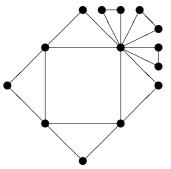
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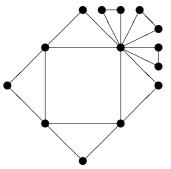
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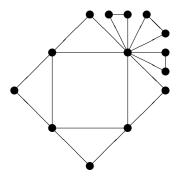


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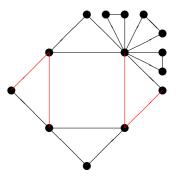
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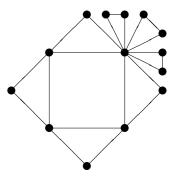


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Average replication	optimal design (probably)
2 and a little above	many small designs (including many leaves) glued at the control
around 3	one large random almost-regular graph with average replication 3.5, also quite a lot of edges from points in this to the control, and a bunch of leaves rooted at the control
4 and above	a random almost-regular graph (maybe with a few leaves)

Large blocks; many unreplicated treatments

The milling phase of a wheat variety trial has 224 varieties in 14 blocks of size 20.

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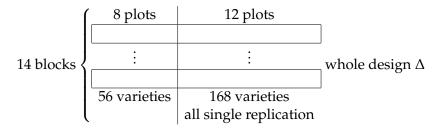
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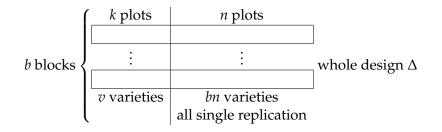


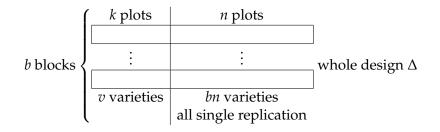
Subdesign Γ has 56 varieties in 14 blocks of size 8.

Yates (1936) concluded that square-lattice incomplete-block designs are superior to the inclusion of controls, but all of his examples had average replication equal to three or more.

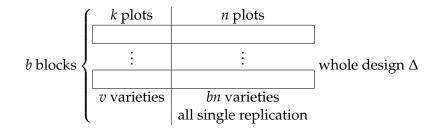
Yates (1936) concluded that square-lattice incomplete-block designs are superior to the inclusion of controls, but all of his examples had average replication equal to three or more.

Here we assume that average replication is (much) less than two.

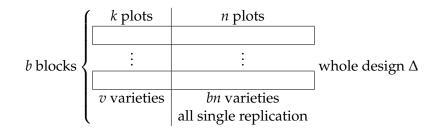




Whole design Δ has v + bn varieties in b blocks of size k + n;

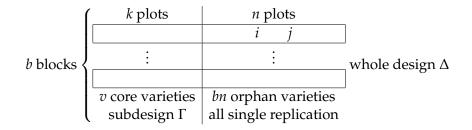


Whole design Δ has v + bn varieties in b blocks of size k + n; the subdesign Γ has v core varieties in b blocks of size k;

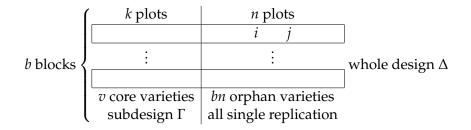


Whole design Δ has v + bn varieties in b blocks of size k + n; the subdesign Γ has v core varieties in b blocks of size k; call the remaining varieties orphans.

Pairwise variance: two orphans in the same block

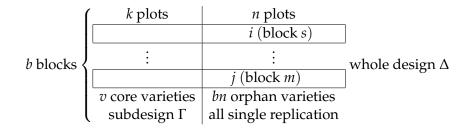


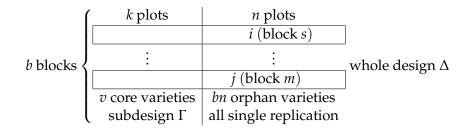
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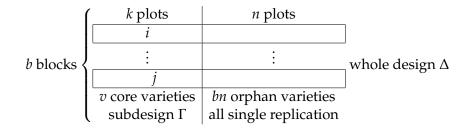
$$\operatorname{Var}_{\Delta}(\hat{\tau}_i - \hat{\tau}_j) = 2\sigma^2.$$

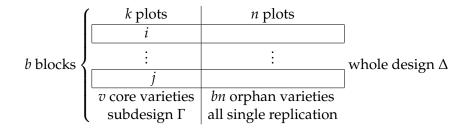
Pairwise variance: two orphans in different blocks





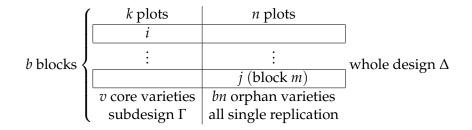
$$\operatorname{Var}_{\Delta}(\hat{\tau}_i - \hat{\tau}_j) = 2\sigma^2 + \operatorname{Var}_{\Gamma}(\hat{\beta}_s - \hat{\beta}_m).$$



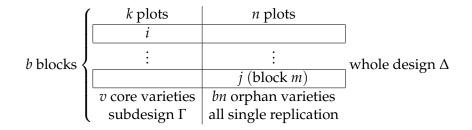


$$\operatorname{Var}_{\Delta}(\hat{\tau}_i - \hat{\tau}_j) = \operatorname{Var}_{\Gamma}(\hat{\tau}_i - \hat{\tau}_j).$$

Pairwise variance: one core variety and one orphan



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$$\operatorname{Var}_{\Delta}(\hat{\tau}_i - \hat{\tau}_j) = \sigma^2 + \operatorname{Var}_{\Gamma}(\hat{\tau}_i + \hat{\beta}_m).$$

Sum of the pairwise variances

Theorem (cf Herzberg and Jarrett, 2007) The sum of the variances of treatment differences in Δ

$$= constant + V_1 + nV_3 + n^2V_2,$$

where

- V_1 = the sum of the variances of treatment differences in Γ
- $V_2 = the sum of the variances of block differences in \Gamma$
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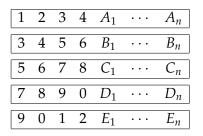
(If Γ is equi-replicate then V_2 and V_3 are both increasing functions of V_1 .)

Consequence

For a given choice of k, make Γ as efficient as possible.

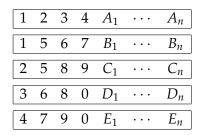
Consequence

If *n* or *b* is large, it may be best to make Γ a complete block design for *k*' controls, even if there is no interest in comparisons between new treatments and controls, or between controls.

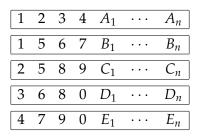


Youden and Connor (1953): "experiments in physics do not need much replication because results are not very variable" —introduced chain block designs

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subdesign is dual of BIBD (Herzberg and Andrews, 1978)

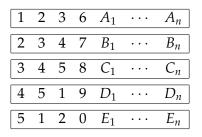


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$$\begin{bmatrix} K_1 & K_2 & 1 & 2 & A_1 & \cdots & A_n \\ \hline K_1 & K_2 & 3 & 4 & B_1 & \cdots & B_n \\ \hline K_1 & K_2 & 5 & 6 & C_1 & \cdots & C_n \\ \hline K_1 & K_2 & 7 & 8 & D_1 & \cdots & D_n \\ \hline \hline K_1 & K_2 & 9 & 0 & E_1 & \cdots & E_n \\ \end{bmatrix}$$

better for large *n* if b > 13 even if there is no interest in controls

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