

EXPERIMENTS IN SQUARE LATTICE WITH A COMMON TREATMENT IN ALL BLOCKS

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INTRODUCTION

Since a long time ago, the **Instituto Agronômico de Campinas**, a research institute located em Campinas, São Paulo, Brasil, uses for its experiments with corn (maize) varieties and hybrids, square lattices with k^2 treatments and blocks of $k + 1$ plots, the extra plot in each block receiving a standard variety or hybrid, the same for all blocks, not included among the k^2 original treatments. It is clear, therefore, that these square lattice experiments include, on the whole, $k^2 + 1$ treatments in blocks of $k + 1$ plots. For example, in a 3^2 lattice, with 2 orthogonal replications, and treatments 1, 2, ..., 9, plus treatment A (standard variety), the blocks would be as follows:

| |
|-------------------|
| Block 1 : 1 2 3 A |
| Block 2 : 4 5 6 A |
| Block 3 : 7 8 9 A |
| Block 4 : 1 4 7 A |
| Block 5 : 2 5 8 A |
| Block 6 : 3 6 9 A |

This paper deals with the intrablock analysis of these designs.

INTRABLOCK ANALYSIS

In these designs we have $v = k^2 + 1$ treatments in a square lattice with m orthogonal replications, $b = mk$ blocks, of $k + 1$ plots.

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There are k^2 regular treatments (1, 2, ..., k^2), plus a common treatment A. The parameter λ_{ij} is equal to 1 for regular treatments that appear in the same block (first associates), equal to zero for regular treatments that do not appear in the same block (second associates). But when one of the treatments is the common one, then $\lambda_{ij} = m$. So, the normal equations (KEMPTHORNE, 1952; PIMENTEL GOMES, 1969) will have coefficients.

$$c_{ij} = m\left(1 - \frac{1}{k+1}\right) = \frac{mk}{k+1},$$

$$c_{ij} = -\frac{1}{k+1} \quad (\text{first associates}),$$

$$c_{ij} = 0 \quad (\text{second associates}),$$

$$c_{iA} = -\frac{m}{k+1} \quad (\text{a regular treatment and the common treatment}),$$

$$c_{AA} = mk\left(1 - \frac{1}{k+1}\right) = \frac{mk^2}{k+1}.$$

Notice: For the example above, the equation corresponding to treatment 1 is:

$$\frac{mk}{k+1}t_1 - \frac{1}{k+1}t_2 - \frac{1}{k+1}t_3 - \frac{1}{k+1}t_4 - \frac{1}{k+1}t_7 - \frac{m}{k+1}t_A = Q_1$$

and for treatment A (common) it is:

$$-\frac{m}{k+1}t_1 - \frac{m}{k+1}t_2 - \dots - \frac{m}{k+1}t_9 + \frac{mk^2}{k+1}t_A = Q_A.$$

Since matrix C = (c_{ij}) is singular, we introduce a restriction for the treatment effects, which can be

$$k^2$$

$$\sum_{i=1}^{k^2} t_i + kt_A = 0.$$

The solution of the system of normal equation gives then:

$$t_A = \frac{1}{mk} Q_A ,$$

$$t_i = \frac{1}{m} Q_i + \frac{1}{k(mk - k + m)} Q_A +$$

$$+ \frac{1}{m(mk - k + 1)} \left[S_1(Q_i) + S_2(Q_i) + \dots + S_m(Q_i) \right] ,$$

where $S_j(Q_i)$ is the sum of Q's in the j^{th} replication, in the block where the i^{th} treatment appears.

The sum of squares for treatments (adjusted) is:

$$\begin{aligned} SST(\text{adjusted}) &= \frac{1}{m} \sum Q_i^2 + \frac{m-1}{m(mk - k + 1)} Q_A^2 + \\ &+ \frac{1}{m(mk - k + m)} \sum_{j,j'} S_{jj'}^2(Q) , \end{aligned}$$

where $S_{jj'}(Q)$ is the sum of Q's in the j^{th} block of the j^{th} replication. So the analysis of variance is obtained as explained in table I.

TABLE I - Analysis of variance.

| Source of variation | D.F. | S:S. |
|--|-------------------|----------------|
| Replications | $m - 1$ | As usual |
| Blocks within replications (unadjusted) | $m(k - 1)$ | As usual |
| Treatments (adjusted) | k^2 | By formula |
| Residual | By subtraction | By subtraction |
| Total | $m k (k + 1) - 1$ | As usual |

The adjusted treatment means are:

$$\hat{m}_j = \frac{G}{mk(k+1)} + t_j, \quad \hat{m}_A = \frac{G}{mk(k+1)} + t_A,$$

where G is the grand total of all plots.

CONTRASTS BETWEEN TREATMENT MEANS

There are 3 cases to be studied.

1st associates: Two regular treatments occurring in the same block, for instance, treatments 1 and 2 in the example above:

$$V(\hat{m}_j - \hat{m}_l) = \frac{2\sigma^2}{m} \left[1 + \frac{m-1}{mk-k+m} \right].$$

2nd associates: Two regular treatments which do not occur in the same block, for instance treatments 1 and 6 in the example above:

$$V(\hat{m}_j - \hat{m}_l) = \frac{2\sigma^2}{m} \left[1 + \frac{m}{mk-k+m} \right].$$

3rd associates: A regular treatment and the common treatment:

$$V(\hat{m}_j - \hat{m}_A) = \sigma^2 \left[\frac{1}{m} + \frac{1}{mk} + \frac{k-1}{k(mk-k+m)} \right].$$

EXAMPLE OF ANALYSIS

We shall take as example a 5×5 square lattice with 4 orthogonal replications and a common treatment A present in all blocks. The experiment was carried out with corn (maize), and harvest expressed in kg./ha. (table II).

TABLE II - Yields, in kg/ha, of corn (maize) in the trial in the quadruple square lattice, with a common treatment in all blocks, used as example.

| Block Number | | | | | | | | | Block Totals |
|---------------------------------|-----------|-----------|-----------|-----------|-----------|----------|-------|--------|-----------------|
| 1st replicate | | | | | | | | | |
| 1 | 6126 (5) | 6497 (11) | 6309 (8) | 6271 (19) | 5743 (22) | 6602 (A) | 37548 | | |
| 2 | 6809 (2) | 6642 (10) | 5111 (13) | 4646 (24) | 5240 (16) | 6173 (A) | 34621 | | |
| 3 | 3670 (4) | 6899 (7) | 5770 (21) | 4167 (18) | 4195 (15) | 6430 (A) | 31131 | | |
| 4 | 6610 (12) | 7166 (20) | 5925 (23) | 5332 (1) | 5509 (9) | 4608 (A) | 35150 | | |
| 5 | 6175 (14) | 7413 (3) | 5768 (6) | 6059 (17) | 5704 (25) | 7202 (A) | 38321 | | |
| | | | | | | | | 176771 | |
| 2nd replicate | | | | | | | | | |
| 6 | 6218 (1) | 6692 (18) | 6621 (10) | 6321 (14) | 6318 (22) | 5787 (A) | 37957 | | |
| 7 | 5580 (2) | 5586 (15) | 4682 (23) | 6155 (19) | 8237 (6) | 5487 (A) | 35727 | | |
| 8 | 6199 (25) | 3844 (4) | 5549 (12) | 5550 (16) | 6480 (8) | 4844 (A) | 32466 | | |
| 9 | 5960 (11) | 5019 (24) | 6300 (7) | 6993 (20) | 5996 (3) | 6280 (A) | 36548 | | |
| 10 | 6986 (17) | 5191 (9) | 7204 (5) | 6999 (21) | 6394 (13) | 6001 (A) | 38775 | | |
| | | | | | | | | 181473 | |
| 3rd replicate | | | | | | | | | |
| 11 | 6350 (25) | 6519 (1) | 5195 (13) | 6187 (19) | 5412 (7) | 4415 (A) | 34078 | | |
| 12 | 4542 (8) | 4330 (8) | 5580 (2) | 3847 (21) | 4339 (14) | 5273 (A) | 27911 | | |
| 13 | 4491 (4) | 6285 (17) | 4927 (11) | 3998 (23) | 3846 (10) | 6044 (A) | 29591 | | |
| 14 | 5374 (22) | 5690 (15) | 4230 (3) | 4177 (9) | 5416 (16) | 5165 (A) | 30052 | | |
| 15 | 6064 (6) | 5992 (18) | 5780 (5) | 5102 (24) | 4692 (12) | 4887 (A) | 31817 | | |
| | | | | | | | | 153449 | |
| 4th replicate | | | | | | | | | |
| 16 | 6052 (7) | 6439 (6) | 6600 (10) | 5855 (8) | 7160 (9) | 6687 (A) | 38793 | | |
| 17 | 4125 (1) | 5822 (2) | 2956 (4) | 7022 (5) | 7804 (3) | 5273 (A) | 33002 | | |
| 18 | 4235 (13) | 4867 (11) | 4734 (15) | 6342 (14) | 7691 (12) | 5744 (A) | 33613 | | |
| 19 | 5199 (16) | 3985 (20) | 5029 (19) | 4998 (18) | 6223 (17) | 4823 (A) | 30257 | | |
| 20 | 5129 (29) | 5880 (25) | 3609 (24) | 5718 (23) | 5538 (22) | 6044 (A) | 31118 | | |
| | | | | | | | | 166783 | |

For treatments 1, 2, ..., 25 and A we compute now the totals T_i ($i = 1, 2, \dots, 25, A$) and the adjusted treatment totals.

$$Q_i = T_i - \sum_j \frac{n_{ij}}{k+1} B_j,$$

TABLE III - Data analyzed, with values of T_i , Q'_i and \hat{m}_i for each treatment.

| Treat. n. ^o | 1 st rep. | 2 nd rep. | 3 rd rep. | 4 th rep. | T_i | Q'_i | \hat{m}_i |
|---------------------------|-------------------------|-------------------------|-------------------------|-------------------------|---------|--------|-------------|
| | | | | | | | |
| 1 | 5332 | 6218 | 6519 | 4125 | 22,194 | -7023 | 5318 |
| 2 | 6809 | 5580 | 5580 | 5822 | 23,791 | 11485 | 6146 |
| 3 | 7413 | 5996 | 4230 | 7804 | 25,443 | 14735 | 6369 |
| 4 | 3670 | 3844 | 4491 | 2956 | 14,961 | -36424 | 3948 |
| 5 | 6126 | 7204 | 5780 | 7022 | 26,132 | 15650 | 6406 |
| 6 | 5768 | 8237 | 6064 | 6439 | 26,508 | 14390 | 6466 |
| 7 | 6899 | 6300 | 5412 | 6052 | 24,663 | 7428 | 5899 |
| 8 | 6309 | 6480 | 4542 | 5855 | 23,186 | 2398 | 5846 |
| 9 | 5509 | 5191 | 4177 | 7160 | 22,037 | -10548 | 5241 |
| 10 | 6642 | 6621 | 3846 | 6600 | 23,709 | 1292 | 5606 |
| 11 | 6497 | 5960 | 4927 | 4867 | 22,251 | -3794 | 5440 |
| 12 | 6610 | 5549 | 4692 | 7691 | 24,542 | 14206 | 6211 |
| 13 | 5111 | 6394 | 5195 | 4235 | 20,935 | -15477 | 4927 |
| 14 | 6175 | 6321 | 4339 | 6342 | 23,177 | 1260 | 5829 |
| 15 | 4195 | 5586 | 5690 | 4734 | 20,205 | -9293 | 5175 |
| 16 | 5240 | 5550 | 5416 | 5199 | 21,405 | 1034 | 5660 |
| 17 | 6059 | 6986 | 6285 | 6223 | 25,553 | 16374 | 6451 |
| 18 | 4167 | 6692 | 5892 | 4998 | 21,749 | -668 | 5636 |
| 19 | 6271 | 6155 | 6187 | 5029 | 23,642 | 4242 | 5941 |
| 20 | 7166 | 6993 | 4330 | 3985 | 22,474 | 4978 | 5948 |
| 21 | 5770 | 6999 | 3847 | 5129 | 21,745 | 1535 | 5642 |
| 22 | 5743 | 6318 | 5374 | 5538 | 22,973 | 1163 | 5673 |
| 23 | 5925 | 4682 | 3998 | 5718 | 20,323 | 9648 | 5130 |
| 24 | 4646 | 5019 | 5102 | 3609 | 18,376 | -23848 | 4589 |
| 25 | 5704 | 6199 | 6350 | 5080 | 23,333 | 4015 | 5827 |
| A | 6602 | 5787 | 4415 | 6687 | | | |
| | 6173 | 5487 | 5273 | 5273 | | | |
| | 6430 | 4844 | 6044 | 5744 | | | |
| | 4608 | 6280 | 5165 | 4823 | | | |
| | 7202 | 6001 | 4287 | 6044 | 113,169 | 538 | 5658 |

where $N = (n_{ij})$ is the incidence matrix, and B_j is the total of block j . It is known that the incidence matrix is obtained from elements n_{ij} , with $n_{ij} = 1$ if treatment i occurs in block j , and $n_{ij} = 0$, if it does not occur.

However, it is easier to calculate $Q'_i = (k + 1) Q_i$, as done in table III. We have:

$$T_1 = 22194$$

$$\begin{aligned} Q'_1 &= 6 T_1 - (B_4 + B_6 + B_{11} + B_{17}) \\ &= 6 \times 22194 - (35150 + 37957 + 34078 + 33002) \\ &= -7023 \end{aligned}$$

$$\begin{aligned} Q'_A &= 6 Q_A = 6 \times 113169 - 678476 \\ &= 538. \end{aligned}$$

Formulas given in section 2 can be easily changed to use Q' - values instead of Q -values. We obtain:

$$t_A = \frac{1}{mk(k+1)} Q_A'$$

$$= (1/120) 538$$

$$= 4.5 \approx 4$$

$$\begin{aligned} t_i &= \frac{1}{m(k+1)} Q_j' + \frac{1}{k(k+1)(mk-k+m)} Q_A' + \\ &+ \frac{1}{(k+1)m(mk-k+m)} \left[S_1 (Q_1') + S_2 (Q_2') + \dots + S_m (Q_m') \right] \end{aligned}$$

$$= (1/24) (-7023) + (1/570) (538) + (1/456) (-20403)$$

$$= -336.$$

On the other hand, we have

$$\hat{m}_1 = 5654 - 336 = 5318,$$

$$\hat{m}_A = 5654 + 4 = 5658.$$

The sum of squares for treatments (adjusted) is:

$$\begin{aligned} SST(\text{adjusted}) &= \frac{1}{m(k+1)^2} \sum Q_i^2 + \frac{m-1}{m(k+1)^2(mk-k+m)} Q_A^2 + \\ &+ \frac{1}{m(k+1)^2(mk-k+m)} \sum_{j,j'} S^2_{jj'} (Q_i) \end{aligned}$$

$$\begin{aligned} SST(\text{adjusted}) &= (1/144) 3,888,633,692 + (1/912) 289,444 + \\ &+ (1/2736) 9,235,411,576 = 30,380,234. \end{aligned}$$

The analysis of variance obtained is given in table IV.

TABLE IV - Analysis of variance of data in table II.

| Source of variation | D.F. | S.S. | M.S. | F |
|--------------------------|------|------------|-----------|-------|
| Blocks | 19 | 35,779,509 | | |
| Treatments (adjusted) | 25 | 30,380,234 | 1,215,209 | 1,67* |
| Error | 75 | 54,631,221 | 728,416 | |

For first associates we have:

$$\begin{aligned}\hat{V}(\hat{m}_i - \hat{m}_j) &= \frac{2(728416)}{4} (1 + 3/19) \\ &= (0,5789) 728416 \\ &= 421680.\end{aligned}$$

For second associates the estimate of variance is:

$$\begin{aligned}\hat{V}(\hat{m}_i - \hat{m}_u) &= \frac{2(728416)}{4} (1 + 4/19) \\ &= (0,6053) 728416 \\ &= 440910.\end{aligned}$$

Finally, for a contrast between the common treatment and any other treatment we have:

$$\begin{aligned}\hat{V}(\hat{m}_i - \hat{m}_A) &= 728416 \left(\frac{1}{4} + \frac{1}{20} + \frac{1}{76} \right) \\ &= 247278.\end{aligned}$$

For the usual 5×5 aquare lattice we should obtain

$$\hat{V}(\hat{m}_i - \hat{m}_u) = 0,6000 s^2$$

for first associates, and

$$\hat{V}(\hat{m}_i - \hat{m}_u) = 0,6333 s^2$$

for second associates. We conclude, therefore, that the new design gives lower estimates for these variances.

SUMMARY

This paper deals with a generalization of square lattice designs, with k^2 treatments in blocks of $k + 1$ plots, the extra plot in each block receiving a standard treatment, the same for all blocks. The new design leads to lower variances for contrasts between adjusted treatment means.

RESUMO

O Instituto Agronômico de Campinas vem, há muitos anos, utilizando nos seus ensaios de milho, reticulados quadrados com k^2 tratamentos em blocos de $k + 1$ parcelas, sendo a parcela extra de cada bloco cultivada com um cultivar padrão (variedade ou híbrido), não incluído entre os k^2 tratamentos originais. Conclui-se, pois, que esses delineamentos incluem $k^2 + 1$ tratamentos, em blocos de $k + 1$ parcelas.

O presente trabalho deduz fórmulas para a análise da variância desses delineamentos, e para a estimativa das médias ajustadas de tratamentos. Fórmulas para a variância de diversos contrastes são deduzidas. Finalmente, apresenta-se um exemplo, detalhadamente analisado, de um ensaio em reticulado quadrado com $k^2 = 25$, e 4 repetições ortogonais, instalado com 26 cultivares, em blocos de 6 parcelas.

LITERATURE CITED

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