



To characterize A- optimal row-column designs for complete Diallel Cross method (2)

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Abstract

The purpose of this paper is to characterize A-optimal designs for complete diallel cross (CDC) method (2) for conducting an experiment with 'p' inbred lines when heterogeneity is to be eliminated in two directions. Some methods of construction of this design are given. AMS Subject Classification: 62 k 05

Keywords: Complete diallel cross; Latin square; Row-column design; Nested balanced incomplete block design; General combining ability; Optimality

Introduction

By a complete diallel cross (CDC) mating design is one in which a set of p inbred lines is chosen and crosses among these lines are made. The procedure gives rise to a maximum of p^2 combinations. Griffing (1956) gave four classifications of CDC mating designs as : CDC method (1) mating design contains p^2 crosses, CDC method (2) mating design contains $p(p+1)/2$ crosses, CDC method (3) mating design contains $p(p-1)$ crosses and CDC method (4) mating design contains $p(p-1)/2$ crosses. The CDC mating designs are often of interest to breeders to investigate the genetic properties and potentials of inbred lines of individuals (for application in plant breeding see, Hallaur and Miranda, 1988). When the experimenter is interested in comparisons of hybrid with their parents, it is advisable to include the parents in the experimental material in the experiment and use CDC method (1) or (2) so that comparisons of hybrid with the parents can be made [1] Suppose there are p inbred lines and it is desired to perform a CDC experiment involving $v = p(p+1)/2$ crosses of the two types $(i \times j)$ and $(j \times i)$ for $i, j = 0, 1, \dots, p-1$, where cross $(i \times j) = (j \times i)$. This is the CDC method (2) mating design of [1] with p inbred lines, having p self and $p(p-1)/2$ distinct F1 crosses, who gave the detailed analysis of such mating designs laid out in a randomized complete block design. Most commonly, CDC mating design have been evaluated using completely or randomized complete block designs as environment designs. In most cases, however, the number of

crosses is too large, leading to an overall inefficient experiment. It is for this reason incomplete block designs were introduced by many authors to resolve the problem of estimating the genetic parameters precisely. However, this approach is not quite appropriate if one is interested in statistical properties, like optimality, of the design. Even an optimal incomplete block design may turn out to be of poor efficiency when used for a CDC experiment. This is because, the analysis of a CDC experiment in incomplete blocks depends on the incidence of inbred lines, rather than on that of treatments or crosses in blocks. Several authors such as [2-9] investigated optimal block designs for CDC method (4) in the 1-way elimination of heterogeneity. [10,11] constructed A- optimal block design for CDC method (2) by using mutually orthogonal Latin squares and nested balanced incomplete block design, respectively, in the 1-way elimination of heterogeneity. The most of the available literature restricts to universal optimality and combinatorial aspects for CDC experiment method (4) and method (2) in the 1-way elimination of heterogeneity set up except [12-13] who studied CDC method (4) in row-column (2-way elimination of heterogeneity) designs. However, row-column designs for complete diallel cross method (2) have not received any attention so far in statistical literature. The primary objective of this paper is to characterize the optimal row-column designs for CDC method (2). Here we are giving some methods of construction of these designs through Latin squares

and nested balanced incomplete block designs. Row-column designs control heterogeneity in the experimental material in two directions. Therefore it increases the precision of the estimates of genetic parameters. The paper is organized as follows. In section 2 we give some preliminary results. In section 3 we give the condition of characterization of these designs. In section 4 we describe the method of analysis of these designs. In section 5 we give the condition of A- optimality of these designs. In section 6 we describe the efficiency factor of these designs in comparison to randomized block designs. In section 7 we give several methods of construction of these designs.

Preliminaries

Suppose that we wish to conduct the diallel cross experiments involving p inbred lines giving rise to a total of $n_c = p(p+1)/2$ distinct crosses of the types $i \times j$ and $(i \times j)$ for $i, j = 0, 1, 2, \dots, p-1$, where cross $(i \times j) = (j \times i)$, in a row-column design with $a \geq p$ rows and $b \geq p$ columns. This is the complete diallel cross (CDC) method (2) of [1], having p self and $p(p-1)/2$ distinct F1 crosses, where p is a prime or power of a prime or prime factors, who gave the detailed analysis of such mating designs laid out in a randomized complete block design. Assume that only one cross of the type $i \times j = j \times i$ and $i \times i$, where $i, j = 0, 1, 2, \dots, p-1$, are applied in each of the $N = a \times b$ plots. We assume the usual fixed-effects additive linear model.

$$y_{(i,j)kl} = \mu + g_i + g_j + \beta_k + \gamma_l + e_{(ij)kl} \quad (4.1)$$

Let $y_{(i,j)kl}$ be the observation on $(i \times j)$ cross applied in the k th row and l th column, μ is grand mean effect, g_i (g_j) is the i th (j th) line gca effect, β_k is the k th row effect, γ_l is the l th column effect and the $e_{(ij)kl}$ are uncorrelated random errors with zero mean and constant variance. The various effects are assumed to satisfy the side conditions.

$$\sum_{i=1}^p g_i = \sum_{k=1}^a \beta_k = \sum_{l=1}^b \gamma_l = 0$$

It is of interest to estimate the general combining ability contrast between i th and j th lines g_j , $i, j = 0, 1, \dots, p-1$. In order for these contrast to be estimable, a necessary condition is that $v \leq ab$, where $v = p(p+1)/2$. Let, $(i, j = 0, 1, \dots, p-1)$ denote the corresponding least squares estimators of g_j , $i, j = 0, 1, \dots, p-1$

Definition: On the above criterion, a design in which $p(p+1)/2$ crosses are allocated in an $a \times b$ array, is a design for CDC method (2) if the least square estimators of gca effects of i th and j th lines $g_i - g_j$ contrast, $i, j = 0, 1, \dots, p-1$, satisfy

$$Var(\hat{g}_i - \hat{g}_j) = \delta^2 \sigma^2, i, j = 0, 1, \dots, p-1 \quad (4.2)$$

where δ^2 is some constant and depend on the particular design employed.

Characterization of CDC Method (2) Design

Consider a row-column design d having m_{ik} incidences of i th line in the k th row and n_{il} incidences of i th line in the l th column ($0 \leq i \leq p-1, 1 \leq k \leq a, 1 \leq l \leq b$). Let $M = \{m_{ik}\}$ and $N = \{n_{il}\}$ denote row and column incidence matrices, respectively. Further let $r_i = \sum_{k=1}^a m_{ik} = \sum_{l=1}^b n_{il}$ be the number of replication of i th treatment, $r = (r_1, r_2, \dots, r_p)$ is a row vector of the replications, $r_{ii'}$ is the number of crosses in d in which i and i' lines appear together and

$$w_{ii'} = \sum_{k=1}^a m_{ik} m_{i'k} \text{ and } v_{ii'} = \sum_{l=1}^b n_{il} n_{i'l}$$

Define

$$\mu_{ii'} = [1/bw_{ii'} + 1/av_{ii'} - r_i r_{i'} / p r_i]$$

We then have the following theorem.

Theorem: A design in which $p(p+1)/2$ crosses are allocated in an $a \times b$ array is a design for CDC method (2) if the least square estimators of the difference of gca's of two lines, say, i and j i.e $g_i - g_j$, $i, j = 0, 1, \dots, p-1$, contrasts satisfy

$$\delta^2 = \frac{2}{(r_i - \mu_{ii} + r_{i'} - \mu_{i'i})}$$

Proof. Consider a diallel cross experiment method (2) involving p lines in a row-column design d with k rows and l columns. The model (2.1) can be written in the matrix form as

$$Y = \mu 1_n + \Delta'_1 g + \Delta'_2 \beta + \Delta'_3 \gamma + e \quad (5.1)$$

Where Y is an $N \times 1$ vector of observed responses, μ is the general mean, g, β and γ are the column vectors of p general combining ability (gca) parameters, a row effects and l column effects, respectively, are the corresponding design matrices, respectively $\Delta'_1 (N \times p), \Delta'_2 (N \times k), \Delta'_3 (N \times l)$ and e denotes the vector of independent random errors having mean 0 and covariance matrix $\sigma^2 I_N$. The various effects are assumed to satisfy the side conditions

$$\sum_{i=1}^p \tau_i = \sum_{k=1}^a \beta_k = \sum_{l=1}^b \gamma_l = 0$$

Let $G_d = \Delta_1 \Delta'_1 = (g_{dii'})$, $g_{dii} = s_{di} = r_i$ i.e the number of times i th line occurs in the crosses in the whole design d and for $i \neq i'$, $g_{dii'}$ is the number of crosses in d in which i and i' lines appear together, $M = \Delta_1 \Delta'_2$ be the $p \times k$ incidence matrix of lines vs rows, $N = \Delta_1 \Delta'_3$ be the $p \times l$ incidence matrix of treatments vs columns and $\Delta_2 \Delta'_3 = 1_l 1'_l$. Let r_{dc} denote the number of times the c th cross appears in the design d , $c = 1, 2, \dots, v$ and similarly r_i denote the number of times the i th line occurs in the design d , $i = 0, 1, \dots, p-1$. Under (4.1), it can be shown that the reduced normal equations for estimating the treatment effects, after eliminating the effect of rows and columns, are

$$C_d \hat{g} = Q$$

$$C_d = G_d - \frac{1}{b} M M' - \frac{1}{a} N N' + \frac{s_d s_d'}{s_d' 1} \tag{5.2}$$

Where

Where C_d is a $p \times p$ information matrix of the lines and $G_d = \Delta_i \Delta_i' = (g_{di})$, $M = (m_{ik})$ is the number of times the line i occurs in row k of d , $N = (n_{il})$ is the number of times the cross i occurs in the column l . Thus, the entries of C_d are

$$C_{ii} = s_{di} - \frac{1}{b} w_{ii} - \frac{1}{a} v_{ii} + \frac{(s_{di})^2}{s_{di}' p} = s_{di} - \mu_{ii} \tag{5.3}$$

and

$$C_{i' i'} = g_{di'} - \frac{1}{b} w_{i' i'} - \frac{1}{a} v_{i' i'} + \frac{(s_{di'})^2}{s_{di}' p} = g_{di'} - \mu_{i' i'} \tag{5.4}$$

Using equations (4.3) and (4.4), we get variance of $(\hat{g}_i - \hat{g}_j)$ as given below.

$$\text{var}((\hat{g}_i - \hat{g}_j)) = \frac{2}{(s_{di} - \mu_{ii} + g_{di'} - \mu_{i' i'})} \sigma^2$$

Hence theorem is proved.

Analysis of CDC Method (2) Design

We next consider the analysis aspects of CDC method (2) designs. Let

$$T_i = \sum_{jkl} y_{ijkl} = i^{th} \text{ line total } (0 \leq i \leq p-1)$$

$$A_k = \sum_{ijl} y_{ij.l} = k^{th} \text{ row total } (1 \leq k \leq a)$$

$$B_l = \sum_{ijk} y_{ijk.} = l^{th} \text{ column total } (1 \leq l \leq b)$$

Further let

$$A_k^* = \sum_{i=0}^a m_{ik} A_k, B_l^* = \sum_{i=1}^b n_{il} B_l, \text{ and } G = \sum_{i=0}^{p-1} T_i = \sum_{k=1}^a A_k = \sum_{l=1}^b B_l = \text{Grand Total}$$

Define the adjusted total for i th line total as

$$Q_i = T_i - \frac{1}{b} A_k^* - \frac{1}{a} B_l^* + \frac{2s_d G}{ab}$$

Hence it can be shown that the least square estimator of $(\hat{g}_i - \hat{g}_j)$ is as given below.

$$(\hat{g}_i - \hat{g}_j) = \frac{Q_i - Q_j}{(s_{di} - \mu_{ii} + g_{di'} - \mu_{i' i'})} \tag{6.1}$$

Note that these designs are not orthogonal but are variance balanced. Hence the analysis of these designs is simple. The analysis of variance (ANOVA) for CDC method (2) is given below in Table1.

Note: The specific combining effects and its sums of squares can also be obtained by using above row-column design for CDC method (2) design [10]

Condition of A-Optimality CDC Method (2) Design

It is not easy to construct A-optimal CDC method (2) design. It is easier to construct CDC method (2) designs that possess

some additional symmetry properties e.g. (i) Designs must be equireplicate in lines. (ii) The number of crosses in which two lines appear together must be equal. (iii) In addition the number of rows and columns should be equal. Let us take a Latin square of standard form with p symbols [14]. If we identify the symbols of a Latin square as lines of a CDC method (2) and perform crosses among the lines appearing in same cell in the first row and also perform crosses between the lines of the first row from corresponding lines in other rows. Thus we get a block design d for CDC method (2) involving $p(p+1)/2$ crosses of the type $(i \times i)$ and $(i \times j)$, where $i, j = 0, 1, \dots, p-1$. The crosses of the type $(i \times j) = (j \times i)$ occurs two times in a design d while crosses of the type $(i \times i)$ occurs once in a design d . The above block design can be arranged in p rows and p columns. Then we get a row-column design d^* with parameters $v = p(p+1)/2$, $b = p$ and $a = p$. In d^*

$m_{d^*ik} = \frac{m_{d^*i.}}{k} = 2$, the design is orthogonal with respect to lines, row vs blocks as classification or a row-regular setting with respect to lines, for $i, j = 0, 1, \dots, p-1, k = 1, \dots, p$.

$nd^*i.l = x$ where $x = \text{int}(mk/p) = 2$ i.e the lines appear two times in columns as blocks, the design is orthogonal with respect to lines, blocks vs rows as a classification, where $m = 2$.

A design $d^* \in \mathcal{ED}$ is said to be A-optimal if and only if $\text{tr}(V_{d^*}) \leq \text{tr}(V_d)$ means that trace of variance-covariance matrix $d^* \in \mathcal{ED}$ is less than equal to trace of variance-covariance matrix d .

Now we give the following theorem without proof [13]

Theorem 5.1: Let $d^* \in \mathcal{ED}(p, b, k)$ be a row-column design for diallel crosses satisfying

$$(i) \text{ trace}(C_d) = k^{-1} b \{ 2k(k-1-2x) + p x(x+1) \}$$

(ii) $(C_d) = (p-1)^{-1} k^{-1} b \{ 2k(k-1-2x) + p x(x+1) \} (I_p - p^{-1} \mathbf{1} \mathbf{1}' p)$ is completely symmetric.

where $x = [2k/p]$, I_p is an identity matrix of order p and $1_p 1_p'$ is a $p \times p$ matrix of all ones. Furthermore, using $d^* \in \mathcal{ED}(p, b, k)$ all elementary contrasts among gca effects are estimated with variance

$$(iii) [2b^{-1}(p-1)k / \{ 2k(k-1-2x) + p x(x+1) \}] \sigma^2.$$

Then according to Kiefer (1975), $d^* \in \mathcal{ED}(p, b, k)$ is universally optimal and, and in particular minimizes the average variance of the best linear unbiased estimator of all elementary contrasts among the gca effects.

By using equations (5.3) and (5.4) we obtain the information matrix C_{d^*}

$$C_{d^*} = 2(p-1) [I_p - \frac{2}{(p-1)} \mathbf{1}_p \mathbf{1}_p'] \tag{7.1}$$

Where I_p is an identity matrix of order p and 1_p is a unit column vector of order p .

The information matrix C_{d^*} is a completely symmetric matrix and its trace = $2(p-1)2$, is greater than the trace $(C_d) = 6p$ given in Theorem (7.1) (i) and also its variance = $1/p-1$ is less than the

variance of Cd = 2(p-1)/6p given in (iii). Hence our proposed design d* is A- optimal.

Efficiency Factor

Efficiency of any design is always compared with no other design except randomized block design. Suppose that instead of the proposed design d*, one adopts a randomized complete block design with 2 blocks, each block having all $p(p+1)/2$ crosses. [1] on page 473 shows that the variance of any elementary contrast among the gca effects is $2\sigma_1^2/(p+2)$, where σ_1^2 is the per observation variance in the case of a randomized block design experiment. It is clear from (5.3) that using design d* each elementary contrast among gca effects is estimated with variance $\sigma_1^2/(p-1)$. Hence the efficiency factor of design d* compared to a randomized block design under the assumption of equal intra block variances is given by

$$E = \frac{2/(p+2)}{1/(p-1)} = \frac{2(p-1)}{p+2} > 1 \tag{8.1}$$

Hence our proposed design is more efficient than randomized block design when $p > 4$.

Methods of Construction D*

Here we are giving some methods of construction of these designs by using Latin squares which possess the above property and also by using some NBIB designs giving by [15] For definition of Latin square and NBIBD [14,15], respectively.

a. Method: Let $p = 2m+1, m \geq 2$. Then cyclically developing the following $2m+1$ columns

$$(0, 2m), (1, 2m-1), (2, 2m-2), \dots, (2m, 0)$$

Yields a row-column for CDC method (2) with parameters $v = p(p+1)/2, b = p$ and $a = p$.

Remark 1. The m columns form a NBIBD with parameters $p = 2m+1, b_1 = m,$

$$k_1 = m(2m+1), k_2 = 2, \lambda_2 = 1.$$

Example 1. For $m = 2$, we get the following row-column design for CDC method (2) with parameters $v = 15, b = 5$ and $a = 5$.

Row-Column Design for CDC method (2)

	B_1	B_2	B_3	B_4	B_5
R_1	0×4	1×3	2×2	3×1	4×0
R_2	1×0	2×4	3×3	4×2	0×1
R_3	2×1	3×0	4×4	0×3	1×2
R_4	3×2	4×1	0×0	1×4	2×3
R_5	4×3	0×2	1×1	2×0	3×4

b. Method: Start with a self-orthogonal Latin square and its transpose; See [16]. Superimpose one over other then we get a row-column design for CDC method (2) with parameters $v = p(p+1)/2, b = p$ and $a = p$.

Example 2: Consider the following self-orthogonal Latin square and its transpose of order 5. After superimposing, we get row-column design for CDC method (2) with parameters $v = 15, b = 5$ and $a = 5$.

0	1	2	3	4	0	3	1	4	2
3	4	0	1	2	1	4	2	0	3
1	2	3	4	0	2	0	3	1	4
4	0	1	2	3	3	1	4	2	0
2	3	4	0	1	4	2	0	3	1

Row-Column Design for CDC method (2)

	B_1	B_2	B_3	B_4	B_5
R_1	0×0	1×3	2×1	3×4	4×2
R_2	3×1	4×4	0×2	1×0	2×3
R_3	1×2	2×0	3×3	4×1	0×4
R_4	4×3	0×1	1×4	2×2	3×0
R_5	2×4	3×2	4×0	0×3	1×1

c. Method: [10] constructed optimal block design for CDC method (2) by superimposing two mutually orthogonal Latin squares of order p . From their designs row-column designs for CDC method (2) with parameters $v = 15, b = 5$ and $a = 5$ can be obtained. The plan is given below.

Row-Column Design for CDC method (2)

	B_1	B_2	B_3	B_4	B_5
R_1	0×0	1×1	2×2	3×3	4×4
R_2	1×2	2×3	3×4	4×0	0×1
R_3	2×4	3×0	4×1	0×2	1×3
R_4	3×1	4×2	0×3	1×4	2×0
R_5	4×3	0×4	1×0	2×1	3×2

Remark 2. Our proposed design consumes less experimental units in comparison to [10]

Method: [17,18] obtained mutually orthogonal Latin squares of order 12 (3×4) and 10(2×5), superimposing these Latin squares we can obtain row-column designs for CDC method (2) with parameters $v = 78, b = 12$ and $a = 12$ and $v = 55, b = 10$ and $a = 10$.

d. Method. [15] gave NBIBD with parameters $(p, b_1, b_2, r, k_1, k_2)$ where $k_1 = p-1, k_2 = 2$ and $p = b_1$. If p is odd, a starter [19,20] in an abelian group of order p is a partition of the non-zero elements of the group into pairs $x_i, y_i (i = 1, 2, \dots, (p-1)/2)$ such that the $(p-1)$ differences $(x_i - y_i)$ and $(y_i - x_i)$ are all different. The $p-1$ differences are thus the $p-1$ non-zero elements of the group. With only a slight change, a starter can thus be used to produce the initial block in the representation of an NBIBD with $b_1 = p, k_1 = p-1$ and $k_2 = 2$. We searched these designs from the catalogue of MPR (2001) and listed in Table 1. These designs can be used for the construction row-column designs for CDC method (2). We are giving example below for the construction of row-column design for CDC method (2).

Table 1: Analysis of Variance.

Source	Degrees of Freedom	Sums of squares
Rows	a-1	$\frac{1}{b} \sum_{j=1}^a A_j^2 - \frac{G^2}{ab}$
Columns	b-1	$\frac{1}{a} \sum_{i=1}^b B_i^2 - \frac{G^2}{ab}$
Crosses	p(p+1)/2 -1	$\sum_{ij}^{p(p+1)/2} y_{ij..}^2 - \frac{G^2}{b}$
General Combining ability	p-1	$\sum_{i=0}^{p-1} \hat{g}_i Q_i$
Error	By subtraction	By subtraction
Total	ab-1	$\sum_{ijkl} y_{ijkl}^2 - \frac{G^2}{b}$

Table 2: A-optimal CDC method (2) designs generated by using NBIB designs Morganet al. (2001).

S.No.	p	b	k	Source
1	7	7	7	(1 6 2 5 4 3) mod 7
				(1 3 2 6 4 5) mod 7
2	9	9	9	(1 8 2 7 3 6 4 5) mod 9
				(1 6 2 8 3 4 5 7) mod 9
3	11	11	11	(1 10 2 9 3 8 4 7 5 6) mod 11
				(1 2 3 6 4 8 5 10 9 7) mod 11
4	7	7	7	(0 1 4 2) (0 2 1 4) (0 4 2 1) mod 7
5	13	13	13	(1 12 5 8) (2 11 3 10) (4 9 6 7) mod 13
				(1 4 2 7) (3 12 6 8) (9 10 5 11) mod 13
6	13	13	13	(3 10 4 9 1 12) (5 8 11 2 6 7) mod 13
				(1 4 3 12 9 10) (2 7 6 8 5 11) mod 13
7	13	13	13	(1 12 2 11 3 10 4 9 5 8 6 7) mod 13
				(1 4 2 7 3 12 5 11 6 8 9 10) mod 13
8	15	15	15	(1 6 3 10) ((1 13 2 8) (1 2 6 8) (1 2 7 14))
				(1 4 2 8) (1 12 2 4) (1 5 2 7) mod 15
				(1 3 6 10) ((1 18 2 13) (1 6 2 8) (1 2 7 14) (1 4 2 8) (1 4 2 12) (1 2 5 7) mod 15)

Example 3: Consider the S.No.1 NBIBD from Table 2 with parameters p = 7, b = 7 and k = 7. We see that there are two distinct starters. We put these starters vertically and cyclically developing mod (7) and attaching the cross of the type i x i in a column where

the ith does not occur Then we get a row-column design for CDC method (2) with parameters v = 28, b = 7, and a = 7 as given below.

- (1×6) (2×0) (3×1) (4×2) (5×3)(6×4) (0×5)
- (2×5) (3×6) (4×0) (5×1) (6×2)(0×3) (1×4)
- (3×4) (4×5) (5×6)(6×0) (0×1)(1×2) (2×3)
- (1×3)(2×4) (3×5)(4×6) (5×0) (6×1) (0×2)
- (2×6)(3×0) (4×1) (5×2) (6×3) (0×4) (1×5)
- (4×5)(5×6) (6×0) (0×1) (1×2) (2×3) (3×4)
- (0×0)(1×1) (2×2) (3×3) (4×4) (5×5) (6×6)

Remark 3. From the designs constructed by Method 1 and Method 4, one can obtain row-column design for CDC method (2) with parameters v = p (p+1)/2, a = (p+1)/2, b = p.

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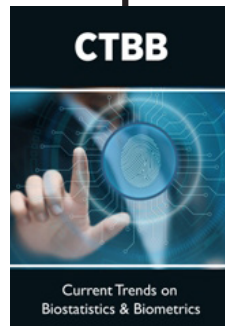


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