

## OPTIMAL CROSS-OVER DESIGNS FOR FULL INTERACTION MODELS

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We consider repeated measurement designs when a residual or carry-over effect may be present in at most one later period. Since assuming an additive model may be unrealistic for some applications and lead to biased estimation of treatment effects, we consider a model with interactions between carry-over and direct treatment effects. When the aim of the experiment is to study the effects of a treatment used alone, we obtain universally optimal approximate designs. We also propose some efficient designs with a reduced number of subjects.

**1. Introduction.** In repeated measurement designs or crossover designs, interference is often observed between a direct treatment effect and the treatment applied in the previous period. We denote by  $\xi_{uv}$  the effect of treatment  $u$  when it is preceded by treatment  $v$ . There are several ways to model such effects. The simplest one is to assume that there is no interference. In that case,  $\xi_{uv} = \tau_u$ , the direct treatment effect.

For a parsimonious interference model, we may assume that the direct and the carry-over effects are additive. In that case,  $\xi_{uv} = \tau_u + \lambda_v$ , where  $\tau_u$  is the direct effect of treatment  $u$  and  $\lambda_v$  is the carry-over effect due to treatment  $v$ . In practice, this model is often unrealistic.

Kempton et al. (2001) propose an interference model in which a treatment which has a large direct effect will also have a large carry-over effect. More precisely, they assume that the carry-over effect is proportional to the direct effect. Bailey and Kunert (2006) obtain optimal designs under this model.

Afsarinejad and Hedayat (2002) proposed another way to enrich the additive models: they assume that the carry-over effect of a treatment depends on whether that treatment is preceded by itself or not. In that case  $\xi_{uv} = \tau_u + \lambda_v + \chi_{uv}$ , where  $\chi_{uv} = 0$  if  $u \neq v$  and  $\chi_{uu}$  represents the specific effect of treatment  $u$  preceded by itself. For that model, optimal designs are obtained by Kunert and Stufken (2002, 2008) when the parameters of

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*MSC 2010 subject classifications:* Primary 62K05, 62K10; secondary

*Keywords and phrases:* cross-over designs, repeated measurement designs, interference models, optimal design, total effects.

interest are the direct treatment effects, and by Druilhet and Tinsson (2009) when the parameters of interest are the total effects  $\tau_u + \lambda_u + \chi_{uu}$ .

The finest possible model, proposed by Sen and Mukerjee (1987), assumes full interactions between carry-over and direct treatment effects, which means that no constraints on  $\xi_{uv}$  are assumed. For a full interaction model, there is no natural way to define a direct treatment effect. For example, Park et al. (2011) obtained efficient designs when the parameters of interest are the standard least-squares means of treatments, i.e.  $t^{-1} \sum_v \xi_{uv}$  for  $1 \leq u \leq t$ , where  $t$  is the number of treatments to be compared. Under a full interaction model, the contrasts of the least-squares means depend on all the other treatment effects through their interactions.

When the aim of the experiment is to select a single treatment which will be used alone, i.e. preceded by itself, the relevant effects to be considered are total effects  $\phi_u = \xi_{uu}$  for  $1 \leq u \leq t$ , which correspond to the effect of a treatment preceded by itself: see Bailey and Druilhet (2004) for a review of situations where total effects have to be considered.

Kushner (1997) and Kunert and Martin (2000) proposed a method for obtaining optimal cross-over designs for direct treatment effects in the framework of approximate designs by using Schur-complement properties. The method has three main steps: (i) expressing the information matrix of the whole design as a sum of the information matrices for the sequences of treatments given to individual subjects (Section 3.1); (ii) considering so-called symmetric designs, in which the proportion of subjects given any sequence is invariant under the symmetric group of all permutations of the treatments (Section 3.2); applying maximin procedures to equivalence classes of sequences (Section 4).

A first generalization of these techniques for more general effects was proposed by Druilhet and Tinsson (2009). In this paper, we propose a higher level of generalization by using group theory to obtain optimal designs for total effects under the full interaction interference model. We also propose efficient designs of reduced sizes.

**2. The designs and the model.** We consider a design  $d$  with  $n$  subjects and  $k$  periods. Let  $t$  be the number of treatments. For  $1 \leq i \leq n$  and  $1 \leq j \leq k$ , denote by  $d(i, j)$  the treatment assigned to subject  $i$  in period  $j$ . We assume the following full treatment  $\times$  carry-over interaction model for the response  $y_{ij}$ :

$$(1) \quad y_{ij} = \beta_i + \xi_{d(i,j), d(i,j-1)} + \varepsilon_{ij},$$

where  $\beta_i$  is the effect of subject  $i$  and  $\xi_{uv}$  is the effect of treatment  $u$  when preceded by treatment  $v$ . For the first period, we assume a specific carry-

over effect that can be represented by a fictitious treatment labelled 0:  $\xi_{u0}$  represents the effect of treatment  $u$  with no treatment before. The residual errors  $\varepsilon_{ij}$  are assumed to be independent identically distributed with expectation 0 and variance  $\sigma^2$ . In most applications, a period effect is included in the model. It will be seen in Section 3.3 that optimal designs found for Model (1) are also optimal when period effects are added.

In vector notation, Model (1) can be written:

$$Y = B\beta + X_d\xi + \varepsilon,$$

where  $Y$  is the  $nk$ -vector of responses with entries  $y_{ij}$  in lexicographic order, and  $\beta$  is the  $n$ -vector of subject effects. The entries of the  $t(t+1)$ -vector  $\xi$  are denoted by  $\xi_{uv}$ , also sorted in lexicographic order. The matrices associated with these effects are respectively given by  $B$  and  $X_d$ . Note that  $B = I_n \otimes \mathbb{1}_k$ , where  $I_n$  denotes the identity matrix of order  $n$ , the symbol  $\otimes$  denotes the Kronecker product and  $\mathbb{1}_k$  is the  $k$ -dimensional vector of ones. Also,  $X_d$  is an  $nk \times t(t+1)$  matrix whose entries are all 0 apart from a single 1 in each row. In particular,  $X_d \mathbb{1}_{t(t+1)} = \mathbb{1}_{nk}$ . We have  $\mathbb{E}(\varepsilon) = 0$  and  $\text{Var}(\varepsilon) = \sigma^2 I_{nk}$ .

We denote by  $\phi$  the  $t$ -vector of total effects, which corresponds to the situation where a treatment is preceded by itself. We have  $\phi_u = \xi_{uu}$ , for  $u = 1, \dots, t$ . Denote by  $K$  the  $t(t+1) \times t$  matrix with entries  $K_{uv}^w = 1$  if  $u = v = w$  and 0 otherwise for  $u, w = 1, \dots, t$  and  $v = 0, \dots, t$ , where  $w$  is the single index for the columns and  $uv$  is the double index for the rows, similar to the index for the vector  $\xi_{uv}$ . We have

$$(2) \quad \phi = K'\xi.$$

### 3. Information matrices for total effects.

3.1. *Information matrix for  $\xi$  and  $\phi$ .* Put  $\omega_B = B(B'B)^{-1}B'$ , which is the projection matrix onto the column space of  $B$ , and  $\omega_B^\perp = I_{nk} - \omega_B = I_n \otimes Q_k$  with  $Q_k = \omega_{\mathbb{1}_k}^\perp = I_k - k^{-1}J_k$ , where  $J_k = \mathbb{1}_k \mathbb{1}_k'$ . The information matrix  $C_d[\xi]$  for the vector  $\xi$  is given by (see e.g. Kunert, 1983):

$$C_d[\xi] = X_d' \omega_B^\perp X_d.$$

Note that  $\omega_B^\perp X_d \mathbb{1}_{t(t+1)} = \omega_B^\perp \mathbb{1}_{nk} = \mathbf{0}$ , and so

$$(3) \quad C_d[\xi] \mathbb{1}_{t(t+1)} = \mathbf{0}.$$

Denote by  $X_{di}$  the  $k \times t(t+1)$  design matrix for subject  $i$  and by  $C_{di}[\xi] = X_{di}' Q_k X_{di}$  the information matrix corresponding to subject  $i$  alone. We

have  $X'_d = (X'_{d1}, \dots, X'_{dn})$  and

$$C_d[\xi] = \sum_{i=1}^n C_{di}[\xi] = \sum_{i=1}^n X'_{di} Q_k X_{di}.$$

Note that  $X_{di}$  and therefore  $C_{di}[\xi]$  depend only on the sequence of treatments applied to subject  $i$ . Denote by  $\mathcal{S}$  the set of all sequences of  $k$  treatments. For a design  $d$  and a sequence  $s \in \mathcal{S}$ , denote by  $\pi_d(s)$  the proportion of subjects that receive  $s$ , and denote by  $X_s$  and  $C_s[\xi]$  the associated matrices. We have

$$(4) \quad C_d[\xi] = n \sum_{s \in \mathcal{S}} \pi_d(s) C_s[\xi] = n \sum_{s \in \mathcal{S}} \pi_d(s) X'_s Q_k X_s.$$

The information matrix for the parameter of interest  $\phi = K'\xi$  may be obtained from  $C_d[\xi]$  by the extremal representation (see Gaffke, 1987 or Pukelsheim, 1993):

$$(5) \quad C_d[\phi] = C_d[K'\xi] = \min_{L \in \mathcal{L}_K} L' C_d[\xi] L,$$

where  $\mathcal{L}_K = \{L \in \mathbb{R}^{t(t+1) \times t} \mid L'K = I_t\}$  and the minimum is taken relative to the Loewner ordering. The minimum in (5) exists and is unique for a given design  $d$ . Put  $\mathcal{E}_d = \{L \in \mathcal{L}_K \mid L' C_d[\xi] L = C_d[\phi]\}$ .

In the sequel, the entries of  $L$ , or, more generally, of any matrix of size  $t(t+1) \times t$ , will be denoted by  $L_{uv}^w$ , for  $u, w = 1, \dots, t$ , and  $v = 0, \dots, t$ , where  $w$  is the column index and  $uv$  is the double index for the rows, similar to the vector  $\xi$  or the matrix  $K$ . The  $t \times t$  matrix  $L'K$  has entries  $(L'K)_{uv} = L_{vv}^u$ , for  $u, v = 1, \dots, t$ .

LEMMA 1. *For any design  $d$ , the row and column sums of  $C_d[\phi]$  are zero.*

PROOF. Since  $C_d[\phi]$  is symmetric, we have to prove that  $\mathbb{I}'_t C_d[\phi] \mathbb{I}_t = 0$ . Consider the  $t(t+1) \times t$  matrix  $L$  such that  $L_{vv}^u$  is equal to 1 if  $u = v$  and 0 otherwise. The matrix  $L$  satisfies  $L \mathbb{I}_t = \mathbb{I}_{t(t+1)}$  and the constraint  $L'K = I_t$ . It follows from (5) and (3) that  $0 \leq \mathbb{I}'_t C_d[\phi] \mathbb{I}_t \leq \mathbb{I}'_t L' C_d[\xi] L \mathbb{I}_t = \mathbb{I}'_{t(t+1)} C_d[\xi] \mathbb{I}_{t(t+1)} = 0$ .  $\square$

For a design  $d$ , denote by  $L^*$  a matrix in  $\mathcal{E}_d$ . Since, for any given  $L$ ,  $L' C_d[\xi] L$  is linear in  $C_d[\xi]$ , we have by (4):

$$(6) \quad C_d[\phi] = L^{*'} C_d[\xi] L^* = n \sum_{s \in \mathcal{S}} \pi_d(s) L^{*'} C_s[\xi] L^*.$$

This linearization is the basis of Kushner's methods.

3.2. *Approximate designs and symmetric designs.* An exact design is characterized, up to a subject permutation, by the proportions of sequences that appear in it. These proportions are multiples of  $n^{-1}$ . If we allow the proportions to vary continuously in  $[0, 1]$  with the only restriction that the sum must be equal to 1, we obtain an approximate design. By definition, the information matrices of  $\xi$  and  $\phi$  for an approximate designs are given by (4) and (5) as for an exact design. The second idea of Kushner's method is to find a universally optimal design in the set of approximate designs using the linearized expression (6). If the optimal approximate design is not an exact design, one can calculate a sharp lower bound for efficiency factors of competing exact designs.

We now recall the concepts of permuted sequence, symmetric design and symmetrized design as introduced by Kushner (1997). Let  $\sigma$  be a permutation of the treatment labels  $\{1, \dots, t\}$  and  $s$  a sequence of treatments. The *permuted sequence*  $s_\sigma$  is obtained from  $s$  by permuting the treatment labels according to  $\sigma$ . Similarly, the design  $d_\sigma$  is the design obtained from the design  $d$  by permuting the treatment labels according to  $\sigma$ . A design  $d$  is said to be a *symmetric design* if, for any sequence  $s$  and any permutation  $\sigma$ ,  $\pi_d(s_\sigma) = \pi_d(s)$ . For such a design,  $d$  and  $d_\sigma$  are identical up to a subject permutation, which may be written  $d = d_\sigma$ . From a design  $d$ , we define the *symmetrized design*  $\bar{d}$  by

$$(7) \quad \pi_{\bar{d}}(s) = \frac{1}{t!} \sum_{\sigma \in S_t} \pi_d(s_\sigma), \quad \forall s \in \mathcal{S},$$

where  $S_t$  is the set of all permutations of  $\{1, \dots, t\}$ . It is easy to see that the symmetrized design  $\bar{d}$  is a symmetric design.

To a permutation  $\sigma$  of treatment labels, we may associate a permutation  $\sigma^*$  of the carry-over effect labels  $\{0, 1, \dots, t\}$  where  $\sigma^*(0) = 0$  and  $\sigma^*(u) = \sigma(u)$  for  $u = 1, \dots, t$ . We also associate a permutation  $\tilde{\sigma}$  of  $\{1, \dots, t\} \times \{0, \dots, t\}$  defined by  $\tilde{\sigma}(u, v) = (\sigma(u), \sigma^*(v))$ . We denote by  $P_\sigma$ ,  $P_{\sigma^*}$  and  $P_{\tilde{\sigma}} = P_\sigma \otimes P_{\sigma^*}$  the corresponding permutation matrices: for example,  $P_\sigma(u, v) = 1$  if  $\sigma(u) = v$  and  $P_\sigma(u, v) = 0$  otherwise.

For  $L \in \mathcal{L}_K$ , put  $L_\sigma = P_{\tilde{\sigma}}' L P_\sigma$ . It can be checked that  $P_{\tilde{\sigma}}' K P_\sigma = K$  (see also the definition of the matrix  $L_{(1)}$  after Lemma 4).

LEMMA 2. *For any design  $d$  and any permutation  $\sigma$  in  $S_t$ , we have*

$$(8) \quad C_{d_\sigma}[\xi] = P_{\bar{\sigma}} C_d[\xi] P'_{\bar{\sigma}};$$

$$(9) \quad C_{d_\sigma}[\phi] = P_\sigma C_d[\phi] P'_\sigma;$$

$$(10) \quad C_{\bar{d}}[\xi] = \frac{1}{t!} \sum_{\sigma \in S_t} P_{\bar{\sigma}} C_d[\xi] P'_{\bar{\sigma}};$$

$$(11) \quad C_{\bar{d}}[\phi] \geq \frac{1}{t!} \sum_{\sigma \in S_t} P_\sigma C_d[\phi] P'_\sigma \quad \text{w.r.t. the Loewner ordering};$$

and  $L \in \mathcal{E}_d$  if and only if  $L_\sigma \in \mathcal{E}_{d_\sigma}$ .

PROOF. By definition of  $P_{\bar{\sigma}}$ ,  $X_{d_\sigma} = X_d P'_{\bar{\sigma}}$ , and so  $C_{d_\sigma}[\xi] = X'_{d_\sigma} \omega_B^\perp X_{d_\sigma} = P_{\bar{\sigma}} X'_d \omega_B^\perp X_d P'_{\bar{\sigma}} = P_{\bar{\sigma}} C_d[\xi] P'_{\bar{\sigma}}$ , which corresponds to (8). If  $L \in \mathcal{L}_K$  then  $L' C_{d_\sigma}[\xi] L = L' P_{\bar{\sigma}} C_d[\xi] P'_{\bar{\sigma}} L = P_\sigma L'_\sigma C_d[\xi] L_\sigma P'_\sigma$ . Now  $L'_\sigma K = P'_\sigma L' P_{\bar{\sigma}} P'_{\bar{\sigma}} K P_\sigma = P'_\sigma L' K P_\sigma$ . If  $L \in \mathcal{L}_K$  then  $L' K = I_t$ , so  $L'_\sigma K = I_t$  and  $L_\sigma \in \mathcal{L}_K$ . The same argument with  $\sigma^{-1}$  shows that if  $L_\sigma \in \mathcal{L}_K$  then  $L \in \mathcal{L}_K$ . The Loewner ordering is unchanged by permutations, so

$$C_{d_\sigma}[\phi] = \min_{L \in \mathcal{L}_K} (L' C_{d_\sigma}[\xi] L) = P_\sigma \left( \min_{L_\sigma \in \mathcal{L}_K} L'_\sigma C_d[\xi] L_\sigma \right) P'_\sigma = P_\sigma C_d[\phi] P'_\sigma,$$

and (9) is established. Moreover,  $L \in \mathcal{E}_d$  if and only if  $L_\sigma \in \mathcal{E}_{d_\sigma}$ . Formula (10) follows directly from (8) and (7). Formula (11) follows from (10) and the concavity of the minimum representation (5).  $\square$

We recall that a  $t \times t$  matrix  $C$  is completely symmetric if  $C = a I_t + b J_t$  for some scalars  $a$  and  $b$  or, equivalently, if  $P_\sigma C P'_\sigma = C$  for every permutation  $\sigma$  in  $S_t$ .

LEMMA 3. *If  $d$  is a symmetric design then  $C_d[\phi]$  is completely symmetric.*

PROOF. Since  $d$  is symmetric,  $d_\sigma = d$ . By (9),  $C_d[\phi] = C_{d_\sigma}[\phi] = P_\sigma C_d[\phi] P'_\sigma$  for any permutation  $\sigma$  in  $S_t$ . Therefore  $C_d[\phi]$  is completely symmetric.  $\square$

The key point to obtain an optimal design is to identify the structure of the  $t(t+1) \times t$  matrix  $L^*$  defined in (6), whose entries are denoted by  $L_{uv}^{*w}$ .

LEMMA 4. *If  $d$  is a symmetric design then the matrix  $L^*$  in (6) can be chosen so that it satisfies*

$$(12) \quad L_\sigma^* = L^*, \quad \forall \sigma \in S_t,$$

or, equivalently,

$$(13) \quad L_{\sigma(u)\sigma^*(v)}^{*\sigma(w)} = L_{uv}^{*w}, \quad \forall \sigma \in S_t.$$

PROOF. If  $\sigma \in S_t$  then  $d_\sigma = d$ , so  $\mathcal{E}_{d_\sigma} = \mathcal{E}_d$  and Lemma 2 shows that  $L_\sigma \in \mathcal{E}_d$ . Put  $L^* = (\sum_{\sigma \in S_t} L_\sigma) / t!$ , which satisfies (12). Since  $\mathcal{E}_d$  is closed under taking averages (see Druilhet and Tinson, 2009, proof of Lemma A1),  $L^*$  also belongs to  $\mathcal{E}_d$ .  $\square$

A consequence of (13) is that the entries  $L_{uv}^{*w}$  are constant for  $(u, v, w)$  belonging to the same orbit of the permutation group  $\{(\tilde{\sigma}, \sigma)\}_{\sigma \in S_t}$  acting on  $\{1, \dots, t\} \times \{0, \dots, t\} \times \{1, \dots, t\}$ . There are seven distinct orbits:

- $\mathcal{O}_1 = \{(u, u, u) \mid u = 1, \dots, t\}$ ,
- $\mathcal{O}_2 = \{(u, v, u) \mid u, v = 1, \dots, t, u \neq v\}$ ,
- $\mathcal{O}_3 = \{(u, v, v) \mid u, v = 1, \dots, t, u \neq v\}$ ,
- $\mathcal{O}_4 = \{(u, v, w) \mid u, v, w = 1, \dots, t, u \neq v \neq w \neq u\}$ ,
- $\mathcal{O}_5 = \{(u, 0, u) \mid u = 1, \dots, t\}$ ,
- $\mathcal{O}_6 = \{(u, 0, w) \mid u, w = 1, \dots, t, u \neq w\}$ ,
- $\mathcal{O}_7 = \{(u, u, w) \mid u, w = 1, \dots, t, u \neq w\}$ .

For  $q = 1, \dots, 7$ , denote by  $L_{(q)}$  the  $t(t+1) \times t$  matrix with entries  $L_{(q)uv}^w = 1$  if  $(u, v, w)$  belongs to the orbit  $\mathcal{O}_q$  and 0 otherwise. Note that  $L_{(1)} = K$ .

By construction of  $L_{(q)}$ , we have

$$(14) \quad P_\sigma' L_{(q)} P_\sigma = L_{(q)}, \quad \forall \sigma \in S_t \text{ and } q = 1, \dots, 7.$$

PROPOSITION 5. *For a symmetric design  $d$ , the matrix  $L^*$  in Lemma 4 may be written as*

$$(15) \quad L^* = L_\gamma = L_{(1)} + \sum_{q=2}^6 \gamma_q L_{(q)},$$

where  $\gamma = (\gamma_2, \dots, \gamma_7)$  is a vector of scalars.

PROOF. Since  $L^*$  satisfies (12), it is a linear combination of the matrices  $L_{(q)}$ :  $L^* = \sum_{q=1}^7 \gamma_q L_{(q)}$ . It can be checked that  $L_{(1)}'K = K'K = I_t$ ,  $L_{(7)}'K = J_t - I_t$  and  $L_{(q)}'K = 0$  for  $q = 2, \dots, 6$ . Consequently, the constraint  $L^{*'}K = I_t$  may be written  $\gamma_1 = 1$  and  $\gamma_7 = 0$ .  $\square$

3.3. *The model with period effects.* We consider here the same model as in Section 2 with the addition of a period effect. The response for subject  $i$  in period  $j$  is given by:

$$(16) \quad y_{ij} = \alpha_j + \beta_i + \xi_{d(i,j),d(i,j-1)} + \varepsilon_{ij},$$

where  $\alpha_j$  is the effect of period  $j$ . In vector notation, we have

$$Y = A\alpha + B\beta + X_d\xi + \varepsilon,$$

with  $A = \mathbb{I}_n \otimes I_k$ , where  $\alpha$  is the  $k$ -vector of period effects. Denote  $\theta' = (\xi', \alpha')$ . If  $d$  is an exact design, the information matrix for  $\theta$  is given by:

$$\tilde{C}_d[\theta] = \begin{pmatrix} C_d[\xi] & C_{d12} \\ C_{d21} & C_{d22} \end{pmatrix} = \begin{pmatrix} X_d' \omega_B^\perp X_d & X_d' \omega_B^\perp A \\ A' \omega_B^\perp X_d & A' \omega_B^\perp A \end{pmatrix},$$

where  $C_d[\xi]$  is the information matrix for  $\xi$  obtained in the model without period effects and  $C_{d22} = nQ_k$ .

The  $t$ -vector  $\phi$  of total effects defined by (2) may also be seen as a subsystem of the parameter  $\theta$ , because  $\phi = \tilde{K}'\theta$  with  $\tilde{K}' = (K', 0_{t \times k})$ . The information matrix  $\tilde{C}_d[\phi]$  for  $\phi$  under Model (16) may be obtained from  $\tilde{C}_d[\theta]$  by the extremal representation:

$$\tilde{C}_d[\phi] = \min_{\tilde{L} \in \mathcal{L}_{\tilde{K}}} \tilde{L}' C_d[\theta] \tilde{L},$$

where  $\mathcal{L}_{\tilde{K}} = \{\tilde{L} \in \mathbb{R}^{(t(t+1)+k) \times t} \mid \tilde{L}'\tilde{K} = I_t\}$ . Partitioning  $\tilde{L}'$  as  $(L' \mid N')$  with  $L$  and  $N$  of sizes  $t(t+1) \times t$  and  $k \times t$ , we have

$$(17) \quad \tilde{C}_d[\phi] = \min_{(L' \mid N')' \in \mathcal{L}_{\tilde{K}}} (L' C_d[\xi] L + L' C_{d12} N + N' C_{d21} L + N' C_{d22} N).$$

Note that  $(L' \mid N')' \in \mathcal{L}_{\tilde{K}}$  is equivalent to  $L \in \mathcal{L}_K$  for  $L$  and  $N$  with suitable dimensions. Choosing  $N = 0$  in (17), we have  $\tilde{C}_d[\phi] \leq C_d[\phi]$  with respect to the Loewner ordering, where  $C_d[\phi]$  is the information matrix for  $\phi$  under the model without period effects, as defined in (5). Therefore  $0 \leq \mathbb{I}_t' \tilde{C}_d[\phi] \mathbb{I}_t \leq \mathbb{I}_t' C_d[\phi] \mathbb{I}_t = 0$ . Hence the row and column sums of  $\tilde{C}_d[\phi]$  are all zero, and so  $Q_t \tilde{C}_d[\phi] Q_t = \tilde{C}_d[\phi]$ .

For  $\sigma \in S_t$ , define the permutation  $\tilde{\sigma}$  for the entries of  $\theta$  such that the entries of  $\xi$  are permuted according to  $\tilde{\sigma}$  and  $\alpha$  remain unchanged. The associated permutation matrix  $P_{\tilde{\sigma}}$  is the block diagonal matrix with diagonal blocks  $P_{\tilde{\sigma}}$  and  $I_k$ . For  $\tilde{L}$  in  $\mathcal{L}_{\tilde{K}}$ , put  $\tilde{L}_\sigma = P_{\tilde{\sigma}}' \tilde{L} P_{\tilde{\sigma}}$ . If  $\tilde{L}' = (L' \mid N')$  then  $\tilde{L}'_\sigma = (L'_\sigma \mid N'_\sigma)$ , where  $N_\sigma = N P_{\tilde{\sigma}}$ .

The two sentences have been changed by Pierre.



LEMMA 6. For any design  $d$  and any permutation  $\sigma$  of treatment labels, we have

$$(18) \quad C_{d_\sigma 12} = P'_\sigma C_{d12};$$

$$(19) \quad \tilde{C}_{d_\sigma}[\phi] = P'_\sigma \tilde{C}_d[\phi] P'_\sigma.$$

PROOF. Equation (18) follows from the fact that  $X_{d_\sigma} = X_d P'_\sigma$ . The proof of (19) is similar to the proof of (9), replacing  $\xi$ ,  $L$ ,  $\mathcal{L}_K$  and  $K$  by  $\theta$ ,  $\tilde{L}$ ,  $\mathcal{L}_{\tilde{K}}$  and  $\tilde{K}$  respectively.  $\square$

An exact design is said to be *strongly balanced on the periods* if it satisfies the following conditions:

- (i) for the first period, each treatment appears equally often,
- (ii) for any given period, except the first one, each treatment appears preceded by itself equally often,
- (iii) for any given period, except the first one, the number of times a treatment, say  $u$ , is preceded by another treatment  $v$  does not depend on  $u$  or  $v$ .

Note that a symmetric exact design is strongly balanced on the periods.

LEMMA 7. If a design  $d$  is strongly balanced on the periods and  $\sigma \in S_t$  then  $P'_\sigma X'_d A = X'_d A$ .

PROOF. The  $(uv, j)$ -entry of  $X'_d A$  is equal to the number of times that treatment  $u$  occurs in period  $j$  preceded by treatment  $v$ . Strong balance implies that there is a single value for  $v = 0$ , another single value for  $v = u$ , and another single value for  $v \notin \{0, u\}$ . Permutation of the treatments does not change this.  $\square$

Given a design  $d$ , let  $G_d$  be the subgroup of  $S_t$  consisting of those permutations  $\sigma$  satisfying  $d_\sigma = d$  (up to a subject permutation). Note that a symmetric design may be characterized by  $G_d = S_t$ . The subgroup  $G_d$  is said to be *transitive* on  $\{1, \dots, t\}$ , if, given  $u, v$  in  $\{1, \dots, t\}$ , there is some  $\sigma$  in  $G_d$  with  $\sigma(u) = v$ . The subgroup  $G_d$  is *doubly transitive* if, given  $u_1, u_2, v_1, v_2$  with  $u_1 \neq u_2$  and  $v_1 \neq v_2$  there is some  $\sigma$  in  $G_d$  with  $\sigma(u_1) = v_1$  and  $\sigma(u_2) = v_2$ .

PROPOSITION 8. If  $d$  is an exact design with strong balance on the periods and with transitive group  $G_d$ , then the information matrix for  $\phi$  is the same under Models (1) and (16), that is

$$\tilde{C}_d[\phi] = C_d[\phi].$$

*In particular, this is true if  $d$  is a symmetric design.*

PROOF. The method of proof of Lemma 4 shows that the matrix  $\tilde{L}$  used for minimizing may be chosen to satisfy  $P'_\sigma \tilde{L} P_\sigma = \tilde{L}$  for all  $\sigma$  in  $G_d$ . This means that  $L = L_\sigma$  and  $N = N_\sigma = NP_\sigma$  for all  $\sigma$  in  $G_d$ . If  $NP_\sigma = N$  for all  $\sigma$  in  $G_d$  and  $G_d$  is transitive then every row of  $N$  is a multiple of  $\mathbb{1}'_t$ .

We have  $C_{d12} = X'_d \omega_B^\perp A = X'_d A Q_k$ . Lemma 7 shows that if  $L = L_\sigma$  then  $L' C_{d12} = L'_\sigma X'_d A Q_k = L'_\sigma P'_\sigma X'_d A Q_k = P'_\sigma L' C_{d12}$ . If  $G_d$  is transitive then every column of  $L' C_{d12}$  is a multiple of  $\mathbb{1}_t$ .

Therefore, the expression in (17) is equal to  $L' C_d[\xi] L + c(L, N) J_t$  for some scalar  $c(L, N)$ . Hence

$$\begin{aligned} \tilde{C}_d[\phi] &= Q_t \tilde{C}_d[\phi] Q_t = Q_t \left( \min_{(L|N)' \in \mathcal{L}_{\tilde{K}}} L' C_d[\xi] L + c(L, N) J_t \right) Q_t \\ &= \min_{(L|N)' \in \mathcal{L}_{\tilde{K}}} (Q_t L' C_d[\xi] L Q_t) \\ &= Q_t \left( \min_{L \in \mathcal{L}_K} L' C_d[\xi] L \right) Q_t \\ &= Q_t C_d[\phi] Q_t = C_d[\phi]. \end{aligned}$$

□

For any design  $d$  whose  $G_d$  is doubly transitive,  $C_d[\phi]$  is completely symmetric (replace  $S_t$  by  $G_d$  in the proof of Lemma 3). Double transitivity implies strong balance on the periods, so then  $\tilde{C}_d[\phi]$  is also completely symmetric, by Proposition 8. In Section 5.6 we give some examples that show that strong balance on the periods is not sufficient for  $\tilde{C}_d[\phi]$  to be completely symmetric.

The results obtained in this section also hold for approximate designs. Since the restriction of  $A$  to a single sequence is equal to  $I_k$ , for an exact designs  $d$  we have

$$\tilde{C}_d[\theta] = n \sum_{s \in \mathcal{S}} \pi_d(s) \begin{pmatrix} X'_s Q_k X_s & X'_s Q_k \\ Q_k X_s & Q_k \end{pmatrix}.$$

This expression can also be used for approximate designs. Moreover, in the definition of a design being strongly balanced on the periods, “equally often” may be replaced by “in the same proportions” and “number of times” by “proportion of times”. Then, the proofs of Lemma 7 and Proposition 8 can be easily adapted to approximate designs by replacing  $A' X_d$  by  $n \sum_s \pi_d(s) X_s$ , replacing  $X'_d \omega_B^\perp A$  by  $\sum_s \pi_d(s) X_s Q_k$ , and so on.

**4. Universally optimal approximate designs.** From Kiefer (1975), a design  $d^*$  for which the information matrix  $C_{d^*}[\phi]$  is completely symmetric and that maximizes the trace of  $C_d[\phi]$  over all the designs  $d$  for  $t$  treatments using  $n$  subjects for  $k$  periods is universally optimal.

4.1. *Condition for optimal designs.* The following proposition shows that a universally optimal approximate design may be sought among symmetric designs.

PROPOSITION 9. *A symmetric design for which the trace of the information matrix is maximal among the class of symmetric designs is universally optimal among all possible approximate designs.*

PROOF. For any design  $d$ , taking the trace in (11), we have  $\text{tr}(C_{\bar{d}}[\phi]) \geq \text{tr}(C_d[\phi])$ . Since, by Lemma 3,  $C_{\bar{d}}[\phi]$  is completely symmetric,  $\bar{d}$  is always better than  $d$  with respect to universal optimality. If  $d^*$  maximizes the trace among the set of symmetric designs, then for any design  $d$ ,  $\text{tr}(C_{d^*}[\phi]) \geq \text{tr}(C_{\bar{d}}[\phi]) \geq \text{tr}(C_d[\phi])$ . Since  $C_{d^*}[\phi]$  is completely symmetric and maximizes the trace,  $d^*$  is universally optimal.  $\square$

For any sequence  $s$ , and  $1 \leq p, q \leq 7$ , put  $c_{spq} = \text{tr}\left(L'_{(p)} C_s[\xi] L_{(q)}\right)$ . Then combining (6), (5) and (15), we have for a symmetric design,

$$\text{tr}(C_d[\phi]) = \min_{\gamma_2, \dots, \gamma_6} \sum_{s \in \mathcal{S}} n \pi_d(s) \sum_{p=1}^6 \sum_{q=1}^6 \gamma_p \gamma_q c_{spq} \quad \text{with } \gamma_1 = 1.$$

LEMMA 10. *For a sequence  $s$  and a permutation  $\sigma$  on the treatment labels, we have:*

$$c_{s_\sigma pq} = c_{spq}.$$

PROOF.

$$\begin{aligned} c_{s_\sigma pq} &= \text{tr}(P'_\sigma L'_{(p)} C_{s_\sigma}[\xi] L_{(q)} P_\sigma), & \text{since } \text{tr}(AB) &= \text{tr}(BA), \\ &= \text{tr}(P'_\sigma L'_{(p)} P'_\sigma C_s[\xi] P_\sigma L_{(q)} P_\sigma), & \text{by (8),} \\ &= \text{tr}(L'_{(p)} C_s[\xi] L_{(q)}) = c_{spq}, & \text{by (14).} \end{aligned}$$

$\square$

Two sequences are said to be *equivalent* if one can be obtained from the other one by some permutation of treatment labels. We denote by  $\mathcal{C}$  the set of all possible *equivalence classes*. From Lemma 10,  $c_{spq}$  depends only on the

equivalence class  $\ell$  to which  $s$  belongs, and will be therefore denoted  $c_{\ell pq}$ . To each equivalence class  $\ell$ , we may also associate the non-negative convex quadratic polynomial with five variables  $\gamma = (\gamma_2, \dots, \gamma_6)$ :

$$h_{\ell}(\gamma) = \sum_{p=1}^6 \sum_{q=1}^6 \gamma_p \gamma_q c_{\ell pq} \quad \text{where } \gamma_1 = 1.$$

For a symmetric design, we may write  $\pi_{\ell}$  for the proportion of sequences which are in the equivalence class  $\ell$ . Then

$$\text{tr}(C_d[\phi]) = \min_{\gamma} \sum_{\ell \in \mathcal{C}} n \pi_{\ell} h_{\ell}(\gamma).$$

Therefore, we have the following proposition:

PROPOSITION 11. *An approximate symmetric design  $d^*$  with proportions  $\{\pi_{\ell}^*\}_{\ell \in \mathcal{C}}$  that achieves*

$$(20) \quad \max_{\{\pi_{\ell}\}_{\ell \in \mathcal{C}}} \min_{\gamma} \sum_{\ell \in \mathcal{C}} \pi_{\ell} h_{\ell}(\gamma)$$

*is universally optimal for  $\phi$  among all possible designs.*

4.2. *Determination of optimal proportions.* Each equivalence class of sequences is defined by a partition of the set  $\{1, 3, \dots, k\}$  into at most  $t$  parts. If  $t \geq k$ , the number of such partitions is the Bell number  $B_k$ , which grows with  $k$  more than exponentially (Cameron, 1994, Chapter 3). Thus it is not realistic to solve the maximin problem in (20) by hand.

It seems intuitive that sequences in an optimal symmetric design should satisfy two contradictory conditions: for accurate estimation of total effects, each treatment should be preceded by itself a large number of times; while, for efficiency in allowing for subjects, the replications within each sequence should be as equal as possible. As a compromise, this suggests sequences in which all occurrences of each treatment are in a run of consecutive periods. Indeed, in our numerical results in Section 5, all sequences in the optimal designs have this form. Each equivalence class of such sequences is defined by a so-called *composition* of  $k$ . However, the number of compositions of  $k$  is  $2^{k-1}$  (Cameron, 1994, Chapter 4), so, even if we restrict ourselves to such sequences, a hand search is still not realistic.

We propose now the following method derived from Kushner (1997). Consider

$$h^*(\gamma) = \max_{\ell \in \mathcal{C}} h_{\ell}(\gamma).$$

We use the following procedure.

- Step 1** Find  $\gamma^*$  that minimizes the function  $h^*(\gamma)$  and denote  $h^* = h^*(\gamma^*)$  the minimum.
- Step 2** Select the classes  $\ell$  of sequences such that  $h_\ell(\gamma^*) = h^*$  and denote  $\mathcal{C}^*$  this set.
- Step 3** Solve in  $\{\pi_\ell \mid \ell \in \mathcal{C}^*\}$  the linear system,  $\sum_{\ell \in \mathcal{C}^*} \pi_\ell \frac{dh_\ell}{d\gamma}(\gamma^*) = 0$ , for  $0 < \pi_\ell < 1$  and  $\sum_{\ell \in \mathcal{C}^*} \pi_\ell = 1$ ; denote  $\pi^* = \{\pi_\ell^* \mid \ell \in \mathcal{C}^*\}$  the solution (not necessarily unique).
- Step 4** Give the symmetric designs such that  $\pi_\ell = \pi_\ell^*$  for  $\ell \in \mathcal{C}^*$  and  $\pi_\ell = 0$  otherwise; these designs are universally optimal.

Step 1 is the most challenging. However, since  $h^*(\gamma)$  is a convex function, any standard optimization algorithm gives accurate values for  $\gamma^*$  and  $h^*$  in a short time even if the number of possible classes is large. When supported by the software, we used an exact optimization algorithm to obtain the values of  $\gamma^*$ .

For Step 2, the optimal sequences are part of the information found in Step 1. Since  $\mathcal{C}^*$  is usually rather small, Step 3 simply involves inverting a small square matrix whose entries have been found in Step 1. Step 4 then reports the results.

**5. Examples of optimal and efficient designs.** For some values of  $k$  and  $t$ , we give optimal approximate designs for  $\phi$ . For each given  $k$ , the first table gives the optimal proportions and the second table gives the efficiency factor for a symmetric design generated by a single sequence.

Consider a real-valued criterion  $\psi(C_d[\phi])$  which is concave, non-decreasing in  $C_d[\phi]$  with respect to the Loewner ordering, and invariant under simultaneous permutations of rows and columns. From Kiefer (1975), there is an approximate design  $d^*$  which maximizes  $\psi(C_d[\phi])$  over the set of approximate designs with the same values of  $k$  and  $t$ . The efficiency factor of a design  $d$  for criterion  $\psi$  can therefore be defined by

$$eff_\psi(d) = \frac{\psi(C_d[\phi])}{\psi(C_{d^*}[\phi])}.$$

For  $\psi(C) = \text{tr}(C)$ , we simply write

$$(21) \quad eff(d) = \frac{\text{tr}(C_d[\phi])}{\text{tr}(C_{d^*}[\phi])}.$$

When  $C_d[\phi]$  is completely symmetric,  $eff(d)$  is also the efficiency factor for the well known  $D$ -,  $A$ - and  $E$ -criteria (see Shah and Sinha, 1989 or Druilhet, 2004).

In our tables, we write  $0^+$  or  $1^-$  when a value is within 0.005 of 0, 1 respectively. For some values of  $k$  and  $t$  the optimal proportions have been calculated with formal calculus when tractable; all others have been obtained by numerical optimization.

The values  $h^*$  displayed correspond to those defined in Section 4.2 for an optimal design. The information matrix for a symmetric optimal approximate design with  $n$  subjects is therefore

$$C_d[\phi] = \frac{n h^*}{t-1} Q_t.$$

5.1. *3 periods.* Optimal proportions for some values of  $t$ :

$t$	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
<i>Prop.</i> [ 1 1 2 ]	$\frac{1}{2}$	$\frac{5}{13}$	$\frac{1}{3}$	$\frac{7}{23}$	$\frac{2}{7}$	$\frac{3}{11}$	$\frac{5}{19}$	$\frac{11}{43}$	$\frac{1}{4}$	$\frac{13}{53}$	$\frac{7}{29}$	$\frac{5}{21}$	$\frac{4}{17}$	$\frac{17}{73}$	$\frac{3}{13}$
<i>Prop.</i> [ 1 2 2 ]	$\frac{1}{2}$	$\frac{8}{13}$	$\frac{2}{3}$	$\frac{16}{23}$	$\frac{5}{7}$	$\frac{8}{11}$	$\frac{14}{19}$	$\frac{32}{43}$	$\frac{3}{4}$	$\frac{40}{53}$	$\frac{22}{29}$	$\frac{16}{21}$	$\frac{13}{17}$	$\frac{56}{73}$	$\frac{10}{13}$
$h^*$	$\frac{1}{3}$	$\frac{16}{39}$	$\frac{4}{9}$	$\frac{32}{69}$	$\frac{10}{21}$	$\frac{16}{33}$	$\frac{28}{57}$	$\frac{64}{129}$	$\frac{1}{2}$	$\frac{80}{159}$	$\frac{44}{87}$	$\frac{32}{63}$	$\frac{26}{51}$	$\frac{112}{219}$	$\frac{20}{39}$

Efficiency of symmetric designs generated by a single sequence:

$t$	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
<i>eff.</i> [ 1 1 2 ]	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>eff.</i> [ 1 2 2 ]	0	0.61	0.75	0.81	0.84	0.86	0.87	0.88	0.89	0.89	0.90	0.90	0.91	0.91	0.91

Example of universally optimal design for  $t = 4$ :

$$\left( \begin{array}{cc} 1 & 1 & 1 & 1 & 1 & 1 & 2 & 2 & 2 & 2 & 2 & 2 & 3 & 3 & 3 & 3 & 3 & 3 & 4 & 4 & 4 & 4 & 4 & 1 & 1 & 1 & 2 & 2 & 2 & 3 & 3 & 3 & 4 & 4 & 4 \\ 2 & 2 & 3 & 3 & 4 & 4 & 1 & 1 & 3 & 3 & 4 & 4 & 1 & 1 & 2 & 2 & 4 & 4 & 1 & 1 & 2 & 2 & 3 & 3 & 1 & 1 & 1 & 2 & 2 & 2 & 3 & 3 & 3 & 4 & 4 & 4 \\ 2 & 2 & 3 & 3 & 4 & 4 & 1 & 1 & 3 & 3 & 4 & 4 & 1 & 1 & 2 & 2 & 4 & 4 & 1 & 1 & 2 & 2 & 3 & 3 & 2 & 3 & 4 & 1 & 3 & 4 & 1 & 2 & 4 & 1 & 2 & 3 \end{array} \right)$$

5.2. *4 periods.* The optimal approximate designs are generated by the single sequence [ 1 1 2 2 ] for  $2 \leq t \leq 30$ . It is conjectured that this is true for any value of  $t$ .

5.3. *5 periods.* Optimal proportions for some values of  $t$ :

$t$	2	3	4	5	6	7	8	9	10	15	20	30
<i>Prop.</i> [ 1 1 2 2 2 ]	$\frac{1}{2}$	$\frac{7}{9}$	$\frac{17}{19}$	$\frac{47}{49}$	0.98	0.98	0.98	0.98	0.98	0.97	0.97	0.97
<i>Prop.</i> [ 1 1 1 2 2 ]	$\frac{1}{2}$	$\frac{2}{9}$	$\frac{2}{19}$	$\frac{2}{49}$	0	0	0	0	0	0	0	0
<i>Prop.</i> [ 1 1 2 3 3 ]	0	0	0	0	0.02	0.02	0.02	0.02	0.02	0.03	0.03	0.03
$h^*$	$\frac{7}{5}$	$\frac{68}{45}$	$\frac{148}{95}$	$\frac{388}{245}$	1.60	1.61	1.62	1.63	1.63	1.64	1.65	1.66



$t$	3	4	5	6	7
<i>Eff.</i> [1 1 1 2 2 2 2]	0.98	0.96	0.95	0.94	0.94
<i>Eff.</i> [1 1 1 2 2 3 3]	0.98	0.99	0.98	0.98	0.98
<i>Eff.</i> [1 1 2 2 3 3 3]	0.98	1 <sup>-</sup>	1 <sup>-</sup>	1 <sup>-</sup>	1

5.6. *Efficient designs with  $t(t-1)$  subjects.* For  $k = 6$  or  $k = 7$ , we saw that efficient symmetric designs may be obtained from single sequences having three treatments by permuting all the treatment labels. Such designs require  $t(t-1)(t-2)$  subjects, which may be too large. We can construct efficient designs that are strongly balanced on the periods, are generated by a single sequence, and require only  $t(t-1)$  subjects, as follows.

**Step 1** We start from a balanced incomplete-block design with block-size 3 and  $t$  treatments such that for any two different periods  $j_1$  and  $j_2$  and any two different treatments  $u$  and  $v$ , there exists exactly one subject that receives treatment  $u$  in period  $j_1$  and treatment  $v$  in period  $j_2$ . (This is called an orthogonal array of type I and strength two: see Rao, 1961.)

- If  $t$  is odd, use all the triplets  $[u, u+v, u+2v]$  modulo  $t$ , for  $u = 0, \dots, t-1$  and  $v = 1, \dots, t-1$ .
- If  $t$  is even, use the preceding construction for  $t-1$  and replace each triplet of the form  $[u, u+1, u+2]$  by the three sequences  $[t, u+1, u+2]$ ,  $[u, t, u+2]$  and  $[u, u+1, t]$ .

**Step 2** Then, we construct a design with  $k$  periods by replicating the three treatments in each triplet in such a way that we obtain a sequence in the same equivalence class as the one that generates the efficient design.

For example, take  $k = 7$  and  $t = 5$  with generating sequence [1 1 2 2 3 3 3]. The starting design with three periods is:

$$\begin{pmatrix} 1 & 1 & 1 & 1 & 2 & 2 & 2 & 2 & 3 & 3 & 3 & 3 & 4 & 4 & 4 & 4 & 5 & 5 & 5 & 5 \\ 2 & 3 & 4 & 5 & 3 & 4 & 5 & 1 & 4 & 5 & 1 & 2 & 5 & 1 & 2 & 3 & 1 & 2 & 3 & 4 \\ 3 & 5 & 2 & 4 & 4 & 1 & 3 & 5 & 5 & 2 & 4 & 1 & 1 & 3 & 5 & 2 & 2 & 4 & 1 & 3 \end{pmatrix}$$

The resulting design with seven periods generated by [1 1 2 2 3 3 3] is

$$\begin{pmatrix} 1 & 1 & 1 & 1 & 2 & 2 & 2 & 2 & 3 & 3 & 3 & 3 & 4 & 4 & 4 & 4 & 5 & 5 & 5 & 5 \\ 1 & 1 & 1 & 1 & 2 & 2 & 2 & 2 & 3 & 3 & 3 & 3 & 4 & 4 & 4 & 4 & 5 & 5 & 5 & 5 \\ 2 & 3 & 4 & 5 & 3 & 4 & 5 & 1 & 4 & 5 & 1 & 2 & 5 & 1 & 2 & 3 & 1 & 2 & 3 & 4 \\ 2 & 3 & 4 & 5 & 3 & 4 & 5 & 1 & 4 & 5 & 1 & 2 & 5 & 1 & 2 & 3 & 1 & 2 & 3 & 4 \\ 3 & 5 & 2 & 4 & 4 & 1 & 3 & 5 & 5 & 2 & 4 & 1 & 1 & 3 & 5 & 2 & 2 & 4 & 1 & 3 \\ 3 & 5 & 2 & 4 & 4 & 1 & 3 & 5 & 5 & 2 & 4 & 1 & 1 & 3 & 5 & 2 & 2 & 4 & 1 & 3 \\ 3 & 5 & 2 & 4 & 4 & 1 & 3 & 5 & 5 & 2 & 4 & 1 & 1 & 3 & 5 & 2 & 2 & 4 & 1 & 3 \end{pmatrix}$$



The following table displays the A-, D-, E-efficiency factors for designs with 6 periods and  $t(t - 1)$  subjects generated by the sequence [ 1 1 2 2 3 3 ] using the method described above. The efficiency factors are given relative to universally optimal approximate designs.

$t$	4	5	6	7	8	9	10
<i>A-efficiency</i>	0.951	0.977	0.973	0.978	0.974	0.970	0.968
<i>D-efficiency</i>	0.951	0.977	0.973	0.978	0.974	0.970	0.968
<i>E-efficiency</i>	0.951	0.977	0.951	0.978	0.950	0.950	0.949

We may note that this method is interesting only for  $t = 7$  or  $t = 8$ . For the other values of  $t$ , the symmetric design with  $t(t - 1)$  subjects generated by the sequence [ 1 1 1 2 2 2 ] is more efficient.

The following table displays the A-, D-, E-efficiency factors for designs with 7 periods and  $t(t - 1)$  subjects generated by the sequence [ 1 1 2 2 3 3 3 ] using the method described above.

$t$	4	5	6	7	8	9	10
<i>A-efficiency</i>	0.974	0.990	0.982	0.983	0.978	0.973	0.971
<i>D-efficiency</i>	0.974	0.990	0.982	0.983	0.978	0.973	0.971
<i>E-efficiency</i>	0.974	0.990	0.961	0.983	0.955	0.954	0.954

For  $t = 4, 5, 7$ , the information matrices are completely symmetric. For  $t \geq 4$  and when the number of subjects is  $t(t - 1)$ , these designs are preferable to symmetric designs generated by the sequence [ 1 1 1 2 2 2 2 ].

changed according to RAB13

If  $t = 4$  or  $t$  is an odd prime, this method always gives a design  $d$  for which  $G_d$  is doubly transitive and so  $\tilde{C}_d[\phi]$  is completely symmetric. If  $t$  is any prime power, there is a second method which gives a design  $d$  in  $t(t - 1)$  periods for which  $G_d$  is completely symmetric.

**Step 1** Identify the treatments with the elements of the finite field  $\text{GF}(t)$  of order  $t$ .

**Step 2** Form any triplet  $[x, y, z]$  of distinct treatments.

**Step 3** Use this to produce all triplets of the form  $[ax + b, ay + b, az + b]$  for which  $a$  and  $b$  are in  $\text{GF}(t)$  and  $a \neq 0$ .

**Step 4** Use these triplets to construct a design from the desired sequence just as in the previous method.

For example, when  $t = 8$ , one correspondence between  $\{1, \dots, 8\}$  and  $\text{GF}(8)$  gives the following starting design with three periods.

$$\begin{pmatrix} 8 & 7 & 1 & 3 & 2 & 6 & 4 & 5 & 8 & 1 & 2 & 4 & 3 & 7 & 5 & 6 & 8 & 2 & 3 & 5 & 4 & 1 & 6 & 7 & 8 & 3 & 4 & 6 & 5 & 2 & 7 & 1 \\ 7 & 8 & 3 & 1 & 6 & 2 & 5 & 4 & 1 & 8 & 4 & 2 & 7 & 3 & 6 & 5 & 2 & 8 & 5 & 3 & 1 & 4 & 7 & 6 & 3 & 8 & 6 & 4 & 2 & 5 & 1 & 7 \\ 1 & 3 & 8 & 7 & 4 & 5 & 2 & 6 & 2 & 4 & 8 & 1 & 5 & 6 & 3 & 7 & 3 & 5 & 8 & 2 & 6 & 7 & 4 & 1 & 4 & 6 & 8 & 3 & 7 & 1 & 5 & 2 \end{pmatrix}$$

$$\begin{pmatrix} 8 & 4 & 5 & 7 & 6 & 3 & 1 & 2 & 8 & 5 & 6 & 1 & 7 & 4 & 2 & 3 & 8 & 6 & 7 & 2 & 1 & 5 & 3 & 4 \\ 4 & 8 & 7 & 5 & 3 & 6 & 2 & 1 & 5 & 8 & 1 & 6 & 4 & 7 & 3 & 2 & 6 & 8 & 2 & 7 & 5 & 1 & 4 & 3 \\ 5 & 7 & 8 & 4 & 1 & 2 & 6 & 3 & 6 & 1 & 8 & 5 & 2 & 3 & 7 & 4 & 7 & 2 & 8 & 6 & 3 & 4 & 1 & 5 \end{pmatrix}$$

The design obtained from this starting design and the generating sequence [ 1 1 2 2 3 3 ], respectively [ 1 1 2 2 3 3 3 ], has efficiency factor equal to 0.977, respectively to 0.981.

For  $t = 9$ , we obtain the following starting design.

$$\begin{pmatrix} 1 & 1 & 2 & 2 & 3 & 3 & 4 & 4 & 5 & 5 & 6 & 6 & 7 & 7 & 8 & 8 & 9 & 9 & 1 & 1 & 4 & 4 & 7 & 7 & 2 & 2 & 5 & 5 & 8 & 8 & 3 & 3 & 6 & 6 & 9 & 9 \\ 2 & 3 & 1 & 3 & 1 & 2 & 5 & 6 & 4 & 6 & 4 & 5 & 8 & 9 & 7 & 9 & 7 & 8 & 4 & 7 & 1 & 7 & 1 & 4 & 5 & 8 & 2 & 8 & 2 & 5 & 6 & 9 & 3 & 9 & 3 & 6 \\ 3 & 2 & 3 & 1 & 2 & 1 & 6 & 5 & 6 & 4 & 5 & 4 & 9 & 8 & 9 & 7 & 8 & 7 & 7 & 4 & 7 & 1 & 4 & 1 & 8 & 5 & 8 & 2 & 5 & 2 & 9 & 6 & 9 & 3 & 6 & 3 \end{pmatrix}$$

$$\begin{pmatrix} 1 & 1 & 5 & 5 & 9 & 9 & 2 & 2 & 6 & 6 & 7 & 7 & 3 & 3 & 4 & 4 & 8 & 8 & 1 & 1 & 6 & 6 & 8 & 8 & 2 & 2 & 4 & 4 & 9 & 9 & 3 & 3 & 5 & 5 & 7 & 7 \\ 5 & 9 & 1 & 9 & 1 & 5 & 6 & 7 & 2 & 7 & 2 & 6 & 4 & 8 & 3 & 8 & 3 & 4 & 6 & 8 & 1 & 8 & 1 & 6 & 4 & 9 & 2 & 9 & 2 & 4 & 5 & 7 & 3 & 7 & 3 & 5 \\ 9 & 5 & 9 & 1 & 5 & 1 & 7 & 6 & 7 & 2 & 6 & 2 & 8 & 4 & 8 & 3 & 4 & 3 & 8 & 6 & 8 & 1 & 6 & 1 & 9 & 4 & 9 & 2 & 4 & 2 & 7 & 5 & 7 & 3 & 5 & 3 \end{pmatrix}$$

The design obtained from this starting design and the generating sequence [ 1 1 2 2 3 3 ], respectively [ 1 1 2 2 3 3 3 ], has efficiency factor equal to 0.950, respectively to 0.954.

*5.7. Comments.* Here we briefly discuss the performances of the optimal designs obtained in this paper when the true statistical model is simpler than the full interaction model.

Under the assumption that the true model is the self and mixed model proposed by Afsarnejad and Hedayat (2002), Druilhet and Tinsson (2014) obtained optimal approximate designs for the estimation of total effects. So, we can compute the efficiency factors of our designs as defined in (21) for several values of  $k$  and for all  $t$  with  $2 \leq t \leq 30$ . For  $k = 3$ , our designs have efficiency factors greater than 0.67. For  $k = 4$ , the optimal designs are the same under both models. For  $k = 5$ , our designs have efficiency factors greater than 0.98. For  $k = 6$ , our designs have efficiency factors greater than 0.97.

We cannot make the analogous comparison under the the assumption that the additive model is the true one, because in this case there are no optimal designs for total effects available in the literature (Bailey and Druilhet, 2004, considered only circular designs).

We now compare our designs to complete-block neighbour-balanced designs (CBNBDs) such as the column-complete latin squares widely used in practice.

Under the self and mixed model, CBNBDs give non-estimable total effects but are optimal for the estimation of direct treatment effects (Kunert and Stufken, 2002). The efficiency factors of our designs for the direct treatment effects are 0.39 for  $k = t = 3$ ; 0.33 for  $k = t = 4$ ; 0.25 for  $k = t = 5$ ; 0.33 for  $k = t = 6$ ; and 0.36 for  $k = t = 7$ .

Under the additive model, the efficiency factors of our designs for the estimation of total effects relative to CBNBDs are 1.15 for  $k = t = 3$ ; 1.31 for  $k = t = 4$ ; 1.24 for  $k = t = 5$ ; 1.33 for  $k = t = 6$ ; and 1.38 for  $k = t = 7$ . For the estimation of direct effects, CBNBDs are optimal (Kunert, 1984, Kushner 1997) and the efficiency factors of our designs are 0.82 for  $k = t = 3$ ; 0.67 for  $k = t = 4$ ; 0.52 for  $k = t = 5$ ; 0.59 for  $k = t = 6$  and 0.61 for  $k = t = 7$ .

**Acknowledgements.** Most of this work was carried out in the autumn of 2011 at the Isaac Newton Institute for Mathematical Sciences in Cambridge, UK, during the Design and Analysis of Experiments programme.

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