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The Pursuit of Balance in Sequential Randomized Trials

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Abstract

In many randomized trials, subjects enter the sample sequentially. Because the covariates for all units are not known in advance, standard methods of stratification do not apply. We describe and assess the method of D_A -optimal sequential allocation (Atkinson, 1982) for balancing stratification covariates across treatment arms. We provide simulation evidence that the method can provide substantial improvements in precision over commonly-employed alternatives. We also describe our experience implementing the method in a field trial of a clean water and handwashing intervention in Dhaka, Bangladesh, the first time the method has been used. We provide advice and software for future researchers.

KEYWORDS: Stratification; Sequential Randomization; Design of Experiments JEL CODES: C93, O12

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Introduction

Randomized-controlled trials (RCTs) are an increasingly important tool for policy evaluation and estimation of economic parameters. However, they are expensive, and efficient use of limited resources (funding, inputs from implementation partners, and researchers' time) requires that they be designed carefully. In an important contribution, Bruhn and McKenzie (2009) reviewed stratification methods that were common in economics RCTs at the time, and showed that large gains in precision were available by adopting more sophisticated stratification methods from the clinical trials literature. These stratification methods require researchers to obtain stratification covariates from all subjects prior to randomization. However, this is not always feasible. In clinical trials, subjects are often allocated to treatment as they arrive. In field trials, operational constraints may prevent defining and surveying the full sample frame in advance. In such situations, subjects must be assigned *sequentially*, with the researcher only learning the value of the stratification variables for that subject's at the time of enrollment and assignment.¹

In this paper, we propose the use of D_A -optimal sequential allocation (Atkinson, 1982) to improve balance and power when subjects are enrolled sequentially. The D_A -optimal method minimizes imbalance given the constraint of not knowing covariate values in advance. We describe the method and its properties, and provide an algorithm for its implementation. We conduct a set of simulations, based on Bruhn and McKenzie (2009), and show that the D_A -optimal method offers clear benefits relative to commonly-used sequential alternatives. In fact, surprisingly, optimal sequential designs are almost as well-balanced as stratifications performed with full knowledge of covariates in advance. In spite of these practical advantages, the method had not, to our knowledge and according to three survey articles, ever been employed in the field.² We describe our experience implementing the method in a water

¹Examples of sequential randomization in economics include Beaman and Magruder (2012), which randomized without stratification, and Bronchetti et al. (2013), which stratified using the block randomization method we describe in Section 4.2.

²See McEntegart (2003), Table 1 in Taves (2010), and Ciolino et al. (2011). Confirmed by personal communication with J. Cicolino, Northwestern University, January 17, 2014.

treatment and hygiene intervention in Dhaka, Bangladesh (Guiteras et al., 2014), and offer practical advice on its implementation under field conditions. Implementation was feasible with standard software (Stata), and produced an allocation that was well-balanced both on the stratification variables chosen *ex ante* and, *ex post*, on other important variables that were not included in the stratification.

1 Theory

Our exposition follows Atkinson (2002), with some changes in notation. First, we lay out the model and notation. Second, we develop the theory for the traditional situation of a fixed population of N subjects, for whom covariates X have been collected in advance. Third, we introduce sequential designs using a simplified case where the researcher is concerned with the precision of all estimated parameters, both treatment effects and nuisance parameters (coefficients on stratification variables). Finally, we adapt the sequential design to the standard situation where only precisely estimated treatment effects are of interest.

1.1 Model and Notation

Suppose the researcher is conducting an individual-level trial with J treatments, including the control treatment. We first consider a linear model with homogeneous treatment effects and i.i.d. errors. In Section 2, we discuss extensions, inclusing heteroscedasticity, nonlinear models, and cluster designs. The model for unit i is

$$y_i = d'_i \alpha + x'_i \beta + \varepsilon_i = w'_i \theta + \varepsilon_i, \tag{1}$$

where d_i is a $J \times 1$ vector of indicator variables assigning unit *i* to a single treatment (i.e., exactly one element of d_i is equal to one), x_i is a $K \times 1$ vector of covariates, and ε_i is an error term. Without loss of generality, we order the treatments with the control condition first. Let $d_i(j)$ indicate assignment to the *j*th treatment; that is, $d_i(1) = (10 \cdots 0)'$, $d_i(2) = (0\ 1\ 0\ \cdots\ 0)'$, etc. We are interested in estimating *contrasts* between the elements of α ; that is, $\alpha_1 - \alpha_2$, $\alpha_1 - \alpha_3$, etc. The control group mean is a nuisance parameter,³ as are the K elements of β (the coefficients on the covariates), so we have K + 1 nuisance parameters and J - 1 parameters of interest.⁴

1.2 Optimal designs with baseline covariates

First, consider a population of N subjects, for whom the researcher has obtained baseline covariates X prior to randomization. The population regression model is given by

$$E[Y] = D\alpha + X\beta = W\theta, \tag{2}$$

where D is the $N \times J$ matrix assigning all subjects to treatment (i.e., $D = \begin{pmatrix} d_1 & \cdots & d_n \end{pmatrix}'$). X is the $N \times K$ matrix of covariates, and α and β are as before. Given the covariates X, our goal is to choose D to minimize the variance of our estimated treatment effect. As a simple example, with one treatment plus a control condition, J = 2, we are interested in the contrast $\alpha_1 - \alpha_2$ and wish to minimize $V [\hat{\alpha}_1 - \hat{\alpha}_2]$.

A useful matrix to create contrasts is

$$L'_{(J-1)\times J} = \begin{bmatrix} 1 & -1 & 0 & 0 \\ 1 & 0 & -1 & 0 \\ & & & \\ 1 & 0 & & -1 \end{bmatrix}.$$

³We are not interested in α_1 per se, but a precise estimate $\hat{\alpha}_1$ is necessary to estimate contrasts precisely.

⁴A more familiar setup for economics readers would include an intercept term as a covariate, so $\alpha_{\sim 1}$ would have J - 1 elements (corresponding to the J - 1 treatment conditions excluding the control) and the augmented covariate vector (1, x')' would have K + 1 elements including the intercept. This turns out to be less convenient for some of the matrix algebra below.

Now we can create a vector of contrasts by premultiplying α by L':

$$L'\alpha = \begin{bmatrix} \alpha_1 - \alpha_2 \\ \vdots \\ \alpha_1 - \alpha_J \end{bmatrix}.$$

To annihiliate the nuisance parameters, we augment L' with a $(J-1) \times K$ matrix of zeros, and define

$$A' = \begin{bmatrix} L' & 0 \end{bmatrix}.$$

The variance of $\hat{\alpha}$ is is proportional to square root of the determinant of the generalized variance:⁵

$$\left|A'(W'W)^{-1}A\right| = \left|L'\left\{D'D - D'X(X'X)^{-1}X'D\right\}^{-1}L\right|.$$
(3)

This quantity is minimized when D'X = 0; that is, when the treatment assignment is *orthogonal* to the covariates, which is to say the treatments are *balanced* across the covariates. When D'X = 0, the generalized variance simplifies, and the determinant is

$$|A'(W'W)^{-1}A| = |L'(D'D)^{-1}L| = J^J/N^{J-1}$$

This minimum possible value is the standard against we any other treatment assignment D. Note that this value is increasing in J and decreasing in N, which matches our intuition that the variance will increase with the number of treatments and decrease with the number of observations.

The relative efficiency of a design D is the ratio of the determinant of the generalized ⁵Recall that $W = \begin{bmatrix} D & X \end{bmatrix}$, so $W'W = \begin{bmatrix} D' \\ X' \end{bmatrix} \begin{bmatrix} D & X \end{bmatrix} = \begin{bmatrix} D'D & D'X \\ X'D & X'X \end{bmatrix}$. Then use results on inverses of partitioned matrices and use the zero block of the matrix A to zero out several terms. variance to this minimal value:

$$\mathcal{E} = \left\{ \frac{J^J / N^{J-1}}{A' \left(W'W \right)^{-1} A} \right\}^{1/(J-1)},$$

where 1/(J-1) is a scale factor. A smaller denominator $|A'(W'W)^{-1}A|$ leads to higher \mathcal{E} , implying a more efficient design. Note that $\mathcal{E} = 1$ for an exactly balanced design. A useful representation is the *loss*

$$\mathcal{L} = N\left(1 - \mathcal{E}\right),$$

which is expressed as the effective loss of observations relative to an optimal design. That is, a non-optimal design D with N units is as precise as an optimal design with $N(1 - \mathcal{E})$ fewer units. For an exactly balanced design, $\mathcal{L} = 0$.

Although not the focus of this paper, this framework can be used for near-optimal randomization in cases where a researcher can collect baseline data prior to randomization. Specifically, create a large number S of random allocations $\{D^1, \ldots, D^s, \ldots, D^S\}$ and choose the allocation D^s with lowest associated loss.⁶ Kasy (2013) considers a more general Bayesian framework, and provides a search algorithm to find an optimal allocation.⁷

1.3 Sequential *D*-optimality

To extend to sequential randomized trials, we first consider the simple case where all elements of $\theta = (\alpha', \beta')'$ are of interest. Our goal is to minimize the variance of $\hat{\theta}$. The variance of $\hat{\theta}$ is proportional to the inverse of the design matrix $(W'W)^{-1}$, so we want to minimize $(W'W)^{-1}$ or, equivalently, maximize |W'W|, which will give us a *D*-optimum design.

Suppose the first n units have been allocated, with the resulting design given by $W_n =$

⁶To conduct randomization inference, rather than choose the allocation with minimum \mathcal{L} , the researcher can instead specify an acceptable maximum $\overline{\mathcal{L}}$, retain R + 1 draws with loss less than $\overline{\mathcal{L}}$, select one of these R+1 at random, and retain the remaining R for randomization inference. Code is available from the authors on request.

⁷This optimal allocation is unique if any element of x is continuous, and may be unique (in finite samples) even for discrete x with a large number of treatments and covariate cells.

 $\begin{bmatrix} D_n & X_n \end{bmatrix}$. Suppose that unit n + 1 arrives, with covariate vector x_{n+1} . By the Generalized Equivalence Theorem of Kiefer and Wolfowitz (1960), maximizing |W'W| is equivalent to minimizing the maximum variance of the predicted response, where this maximum is taken over the space of w. That is, given the current allocation W_n , the variance of \hat{y} at a point w = (d', x')' is

$$V\left[\hat{y}\left(w\right)\right] \propto s\left(w, W_{n}\right) = w'\left(W_{n}'W_{n}\right)^{-1}w,\tag{4}$$

where $s(w, W_n)$ is the standarized variance at w given the allocation W_n . Because minimizing $V\left[\hat{\theta}\right]$ is equivalent to minimizing the maximum variance of the predicted mean, we can restate our objective as minimizing

$$\sup_{w}V\left(\hat{y}\left(w\right)\right).$$

That is, given the existing design W_n and the covariate values x_{n+1} , the optimal assignment for unit n+1 is

$$d_{n+1}^{*}(x_{n+1}, W_{n}) = \arg\min_{d} \left\{ \sup_{w} V(\hat{y}(w)) \right\}.$$

In other words, we want to allocate this unit to the treatment where the variance is greatest. To accomplish this, for unit n + 1, write the set of possible values for w_{n+1} as $w_{n+1}(1) = (d(1)', x'_{n+1})', \ldots, w_{n+1}(J) = (d(J)', x'_{n+1})'$, where d(j) denotes allocation to treatment j. For each $w_{n+1}(j)$, we calculate

$$s_{j} = s \left(w_{n+1}(j), W_{n} \right) = w_{n+1}(j)' \left(W_{n}' W_{n} \right)^{-1} w_{n+1}(j).$$
(5)

The best allocation for unit n + 1 is the d(j) with the largest value of s_j . In this simplest case, we mechanically assign person n + 1 to this treatment. The intuition is that the unit is being assigned to the treatment where it is most needed, which is where the variance is highest. In Section 2.7, we discuss non-deterministic or "biased coin" assignment.

1.4 Sequential D_A -optimality

The *D*-optimal procedure in the previous subsection minimizes $V\left[\hat{\theta}\right]$. That is, it maximizes precision for estimates of both treatment effects and the coefficients on baseline characteristics. However, in most cases our goal is to maximize precision for the estimated treatment effects. That is, our objective is to minimize $V\left[\hat{\alpha}\right]$, and we are not *per se* interested in minimizing $V\left[\hat{\beta}\right]$. Atkinson calls this problem D_A -optimality. The intuition and the basic procedures are the same, but the formula for the standardized variance $s_A\left(w_{n+1}\left(j\right), W_n\right)$ is slightly more complicated:

$$s_A(w_{n+1}(j), W_n) = w_{n+1}(j)' (W'_n W_n)^{-1} A \left\{ A' (W'_n W_n)^{-1} A \right\}^{-1} A' (W'_n W_n)^{-1} w_{n+1}(j), \quad (6)$$

where $A' = \begin{bmatrix} L' & 0 \end{bmatrix}$, as above.

The assignment algorithm follows the logic of Section 1.3. Suppose that n units have been allocated, and the current matrix of assignments and covariates is $W_n = \begin{bmatrix} D_n & X_n \end{bmatrix}$. When unit n + 1 arrives with covariates x_{n+1} , check the value of $s_j = s_A (w_{n+1}(j), W_n)$ for each possible assignment j. The optimal allocation of unit n+1 is where where s_j is greatest.

1.5 Algorithm

The procedure described above cannot be used for the first units, because W'W is singular as long as the number of observations, n, is less than the number of incidental parameters, J. These first units could be assigned randomly, or the W'W matrix could be made invertible by adding a small amount of random noise to the diagonal.

Having allocated n units, allocate unit n + 1 as follows:

- 1. Subject n+1 arrives with x_{n+1}
- 2. For each treatment j, calculate $s_A(w_{n+1}(j), W_n)$ using 6.
- 3. Assign treatment to the study arm where $s_A(w_{n+1}(j), W_n)$ is greatest.

4. Update W_n to W_{n+1} .

2 Extensions

2.1 Unequal allocations

The basic exposition assumes that the researcher places equal weight on the precision of each element of α (that is, on the coefficient of each treatment effect). However, the researcher may wish to weight these unequally – if, for example, the cost of treatments vary, or if external constraints require unequal numbers of treated units. To create unequal allocations, inflate or deflate the corresponding values of $s_j = s_A(w_{n+1}(j), W_n)$. For example, to overweight treatment j, premultiply s_j by an appropriate weight m_j . These weights $\{m_j\}$ can be calculated analytically in some simple cases, or the researcher can conduct simulations to tune the weights. See Section 6 for an example.

2.2 Heteroscedasticity

If the variance of the outcome of interest is a function of treatment, equal allocations may be inefficient because coefficients corresponding to treatments that increase variance will be less precisely estimated. If the researcher has a strong prior that the variance of the outcome of interest is likely to be greater under certain treatment conditions, she can allocate more units to that treatment following the weighting strategy described in the previous subsection.

2.3 Unequal penalties for imbalance

The researcher may wish to emphasize balance in one covariate or set of covariates over others. Suppose one set of K_1 covariates x_1 have very strong predictive power for the outcome variable, while the remaining K_2 covariates x_2 have some predictive power but less than x_1 . The researcher wants to balance against both covariates, but imbalance in x_1 will cause greater efficiency loss than imbalance in x_2 . To assign a greater penalty to imbalance in x_1 , note from Equation (3) that efficiency is maximized when D'X is zero, and decreases as $D'X(X'X)^{-1}X'D$ becomes larger. Partition X into $X = \begin{bmatrix} X_1 & X_2 \end{bmatrix}$. Now

$$D'X (X'X)^{-1} X'D = D' \begin{bmatrix} X_1 & X_2 \end{bmatrix} (X'X)^{-1} \begin{bmatrix} X_1 & X_2 \end{bmatrix}' D$$
(7)
= $\begin{bmatrix} D'X_1 & D'X_2 \end{bmatrix} (X'X)^{-1} \begin{bmatrix} D'X_1 & D'X_2 \end{bmatrix}',$

so to penalize imbalance in x_1 simply multiply $D'X_1$ – but not $D'X_2$ – by a scalar weight m > 1. The same logic applies in the sequential algorithm: in the standardized variance formula (6), replace $D'_n X_{1n}$ with $mD'_n X_{1n}$.⁸ Again, simulations can help choose the appropriate weights.

2.4 Subgroups and Interactions

As described, the algorithm maximizes marginal balance. That is, it minimizes the variance of the estimates of the treatment effects overall, not for any particular value of the covariates. To optimize estimates for subgroups, augment the vector containing the covariates of interest, α , with the relevant interaction coefficients. Similarly, if the interaction of two treatments is of interest, augment α and d appropriately. See Section 6 for an example.

2.5 Nonlinear models

The algorithm is motivated by linear regression, so the allocation it produces may not be optimal for nonlinear designs. In particular, for linear regression, the optimal design does not depend on the value of the unknown parameters, but it does for nonlinear or generalized linear models (Atkinson and Haines, 1996). Therefore, the optimal design, for a

⁸Note that, as with (3), the middle term $A' (W'_n W_n)^{-1} A$ of (6) reduces to $L' \left\{ D'_n D_n - D'_n X_n (X'_n X_n)^{-1} X'_n D_n \right\}^{-1} L$, with the loss coming from $D'_n X_n \neq 0$. Partitioning as above, the second term in the inverse becomes $\left[D'_n X_{1n} \quad D'_n X_{2n} \right] (X'_n X_n)^{-1} \left[D'_n X_{1n} \quad D'_n X_{2n} \right]'$, so we increase the penalty for imbalance in x_1 by replacing $D'_n X_{1n}$ with $mD'_n X_{1n}$.

non-sequential or sequential trial, in a nonlinear model requires that the researcher specify her prior belief distribution about the parameter of interest. While the intuition is similar (maximizing the log of the determinant of the information matrix), the calculation can be quite difficult. However, it is difficult to imagine a scenario where balancing to minimize the variance of OLS estimates would severely worsen the precision of nonlinear estimators. Therefore, we speculate that such concerns are of second order, and that, while the method may not produce the optimal design for a nonlinear model, it is likely to produce a good approximation. In highly specialized situations, there may be some efficiency gains to more specialized solutions.⁹

2.6 Time trends

Because the algorithm seeks to maintain balance at each point in the sequence, it is robust to trends or fluctuations in potential outcomes that occur as sample enrollment proceeds. For example, neither a geographic pattern to enrollment nor a change in recruitment methods would cause bias, even if these were correlated with potential outcomes (e.g., moving from richer to poorer neighborhoods, or making a greater effort to recruit poor subjects).

2.7 Biased coin methods

The D_A -optimal method will produce unbiased estimates as long as each unit's exact place in the sequence is uncorrelated with potential outcomes. This assumption could be violated if, for example, an intake nurse in a clincial trial knows the algorithm and current allocation. The nurse could then manipulate the order in which subjects are processed to ensure that a particular subject receives a particular treatment.

To reduce the possibility of gaming, a "biased coin" version of the sequential allocation algorithm allocates a subject probabilistically, putting highest probability on the study arm

⁹For an example, see Zocchi and Atkinson (1999), who derive the optimal design to study the relationship between a discrete treatment (dose of gamma radiation) and a multinomial, ordered outcome (whether housefly pupae die before opening, die during emergence, or survive past emergence).

that would reduce the variance of estimated treatment effect, $s_A(w_{n+1}(j), W_n)$, the most. Following the logic of Efron (1971) and of Atkinson (1982) studying a related problem, suggests this formula for the probability for allocation to study arm:

$$\pi(j) = \frac{s_A(w_{n+1}(j), W_n)}{\sum_{j'} s_A(w_{n+1}(j'), W_n)}$$

Randomization may also be useful for expositional purposes, to explain to subjects that the process is fair and to users of the research who are used to hearing about "randomized," rather than "optimally allocated," trials.

2.8 Cluster designs

The development above is based on individual-level treatments, but can be adapted for cluster designs. The first application is to interventions where the treatment is assigned at a cluster level. In this case, the D_A -optimal method applies directly, using cluster-level covariates.

A second application is when treatments are assigned to individuals, but individuals belong to clusters and for logistical reasons these clusters are enrolled sequentially. For example, consider a study of the demand for water filters among N households in V villages.¹⁰ The researcher wishes to vary a sales treatment at the household level. However, villages are enrolled – and stratification covariates collected – sequentially, so at the time of assigning treatments in village v, the researcher only knows covariates for households in villages $1, \ldots, v$, and the history of assignments in villages $1, \ldots, v-1$. This is not a purely sequential allocation, since the researcher knows the values of the covariates for all households in village v, and can assign treatments simultaneously within village v. However, the researcher does not know the value of covariates for households in future villages $v+1, \ldots V$. The tools of sequential allocation can be usefully applied in this situation. Suppose that the matrix

¹⁰This example is inspired by Berry, Fischer, and Guiteras (2015), who, regrettably, were not aware of the D_A -optimal method at the time of implementation, and used a complicated version of the Block method described in Section 4.2.

of assignments and covariates through village v - 1 is $W_{v-1} = [D_{v-1} X_{v-1}]$. The researcher obtains covariate data for households in village v, resulting in the covariate matrix X_v for all v villages. The researcher then creates a large number S of random treatment allocations for village v, resulting in a set of assignment matrices $\{D_v^1, \ldots, D_v^s, \ldots, D_v^S\}$ and corresponding design matrices $\{W_v^1, \ldots, W_v^s, \ldots, W_v^S\}$. For each s, we can calculate the associated determinant $A' (W_v^{s'} W_v^s)^{-1} A$, where, as in Section 1.2, the matrix A allows us to focus on the parameters of interest. Since this determinant is proportional to the expected variance of $\hat{\alpha}$, we select the allocation that minimizes this determinant.

3 Inference

Confidence intervals can be constructed from the usual regression-based methods, and the standard covariance matrices can also be used for t-tests of hypotheses. Shao, Yu, and Zhong (2010) prove that controlling for balancing variables will yield tests of the correct size. As emphasized by Bruhn and McKenzie (2009), researchers should commit ex-ante to controlling for the balancing variables, since this increases power on average, but retaining the option to analyze without controlling for the balancing variables gives the researcher a degree of freedom that can distort the size of a test.

Randomization inference can be conducted by following the "reasoned basis for inference" logic of Fisher (1935): because the design assumption is that the precise order of arrival of units is arbitrary, one can construct counterfactual distributions by reshuffling this order in which subjects arrive. If the study also uses a biased-coin approach, then one can rerandomize the biased coin flips. Shao and Yu (2013) also propose a covariate-augmented bootstrap method and show that it provides valid tests for generalized linear models.

4 Simulation Methods

Using the four panel datasets and simulation structure of Bruhn and McKenzie (2009), we compare the D_A -optimal method to two sequential methods, simple randomization and block randomization (permutations within block), which are commonly used in clinical and other trials (McEntegart, 2003). We describe these methods in greater detail in Section 4.2. For each dataset and each allocation method, we simulate 10,000 randomizations. In each iteration, the enrollment order is randomized, treatments are assigned according to the given method, a simulated output variable is created by adding a true treatment effect (possibly zero) to those assigned treatment, and we obtain an estimated treatment effect with associated standard errors and p-values.

4.1 Data

We use the four panel datasets used in Bruhn and McKenzie (2009), who provide further detail on these data. The first dataset covers microenterprises in Sri Lanka and contains information on firms' profits, assets and other firm and owner characteristics. The second dataset is a subsample from the Mexican employment survey (ENE) containing data on heads of household between 20 and 65 years of age who were first interviewed in the second quarter of 2002. The third dataset comes from the 1997 and 2000 waves of the Indonesian Family Life Survey (IFLS) and contains child schooling outcomes as well as household level data such as weekly expenditure. The final dataset comes from the Learning and Educational Achievement Project (LEAPS) in Pakistan and includes math and height z-scores as well as other covariates for children aged 8 to 12 at baseline. From each dataset, we draw subsamples of 30, 100 and 300 observations to allow a comparison of the methods over small, medium and large samples.

4.2 Allocation Methods

In the simulations, we use three allocation methods: a simple, unstratified randomization; block randomization (permutations within strata); and the D_A -optimal method.

In simple randomization, each subject is randomly assigned to treatment as she arrives, with each treatment given the desired probability. In our simulations, there are only two arms (treatment and control), and each is given probability 0.5.

In Block randomization, the researcher creates a separate randomization list for each stratum (unique combination of balancing covariates). For example, when balancing on 3 binary variables, there are 8 (=2³) strata.¹¹ Since the number of subjects who will fall in each stratum is unknown ex ante, a list should maximize overall balance at each point in the sequence while keeping predictability of each allocation low. The Block method uses lists that are sequences of blocks of random but balanced permutations of the treatments. The size of the blocks can be as low as the number of treatments — 2 if there is one treatment and a control (e.g. (T,C), (C,T), (C,T), (T,C) ...) — or higher to reduce predictability (e.g. 2 treatments, block size 4: (T,C,C,T), (C,T,C,T), (T,C,T,C), (C,T,T,C), ...). The size of the blocks may also be varied randomly to minimize predictability. In our simulations, we abstract from concerns about allocation manipulation and set the block size equal to the number of treatments, which achieves the greatest balance and therefore provides the most conservative test of the D_A -optimal method.

Our third method is the D_A -optimal method, for which we follow the algorithm laid out in section 1.5. For the first J units, we added random noise to the diagonal of the matrix. We did not use the biased coin variant.

For the Block and D_A -optimal methods, we use the same balancing variables as Bruhn

¹¹When using the the Block method, the researcher must discretize continuous variables, e.g. above and below the median. However, the median (or other sensible cut point) may not be known in advance. One advantage of the D_A -optimal method is that it allows for continuous covariates. In our application, we included the number of households in each compound as a balancing variable. Although we did not know the distribution of this variable in advance, the D_A -optimal method produced an allocation that was wellbalanced both on the mean of the continuous variable and on the proportion above / below the median.

and McKenzie (2009). The outcome at baseline is always included. Other variables are chosen based on their likelihood of being good predictors of the follow-up outcome.

5 Simulation Results

5.1 Balance at Baseline

We first examine balance on the baseline outcome variable. After each randomization we regress the baseline outcome on the assigned treatment. Since the treatment is fictitious, the distribution of these coefficients gives an idea of the potential for imbalance under each method. Figure 1 presents the distributions for the three methods over a number of datasets at all three sample sizes. Table 1 gives the average coefficient, the 95th percentile of the distribution, and the proportion of iterations with a p-value less than 0.1 for each dataset and method, for the sample size 100.

Panel A of Table 1 shows that all methods, even simple randomization, achieve balance on average at baseline since all have means close to zero. However Panel B shows the methods that balance over the baseline outcome do significantly better than simple randomization at avoiding extremes and that the D_A -optimal method clearly dominates the Block method in this respect. The ninety-fifth percentile of the distribution drops from around 0.4 for simple randomization to around 0.2 for the Block method and to below 0.08 for the D_A -optimal method. This point is well illustrated by the kernel density plots in figure 1 where we see the the distribution of coefficients for D_A -optimal as a sharp spike around zero compared to the more rounded bell curves for Block and simple randomization.

We also see that including additional balancing variables in either method reduces balance on existing balancing variables, although only slightly.

Table 1 also shows how the methods may affect balance on "unobservables." Of course, it is not possible in practice to assess balance on true unobservables, but in a simulation we can mimic unobservables with observed variables we expect to be correlated with the outcome of interest, but which we intentionally exclude from the set of stratification variables. Standard stratification methods should not, on average, lead to more imbalance on unobservables than unstratified randomization (Aickin, 2001). Here, we obtain a similar result for the sequential balancing methods, which do no worse than simple randomization.

5.2 Balance at Follow-up

We now address balance on the outcome of interest at follow-up. Just as for the baseline case, we have regressed the follow-up outcome on the fictitious treatment variable in each iteration. Figure 1 shows the distribution of coefficients. Tables 2 and 3 present the same statistics as Table 1 but for the follow-up case, for sample sizes 30 and 300 and with the addition of Panel D. In all cases, the p-values are based on regressions that take into account the method of randomization: for Block randomization, stratum dummies are added to the regression, for D_A -optimal, the balancing covariates are included as controls.

Panel A from tables 2 and 3 show that balance on the follow-up outcome is achieved on average by all methods. We expect to see the likelihood of extreme imbalance reduced only when the balancing covariates used are good predictors of the outcome. Bruhn and McKenzie note that covariates in the LEAPS datasets (specifically, lagged values for both outcomes, math and height z-scores) have high predictive power, with 43% or more of the variation in the outcome explained by the balancing covariates. The Sri Lankan microenterprise data and the IFLS schooling data fall at the other end of the spectrum with 17% or less explained. The results in panel B from table 2 confirm this hypothesis. For both of the LEAPS outcomes, we see substantial improvements in balance when using Block or D_A optimal methods compared to simple randomization. However, for outcomes where balancing covariates have low predictive power, we see no improvement, and perhaps even slightly worse balance in the case of the Mexican employment data. D_A -optimal does better than Block more often than not, but the advantage is not very large.

Additionally, as with Bruhn and McKenzie (2009), we see that the benefit of using

covariates to balance attenuates as the sample size increases (compare Panel B of Table 2 to Table 3). We also confirm that test statistics are incorrect when not controlling for the method of randomization (see Panel C vs Panel D in Tables 2 and 3).

5.3 Power to detect a given treatment effect

Lastly, we turn to perhaps the most important question: how do the methods compare in terms of power for detecting a given treatment effect? To answer this question, we add a constant treatment effect to the output of all subjects assigned to treatment,¹² and then regress the modified outcome on treatment. Tables 4 and 5 presents the proportion of estimated treatment effects that are significant at the 0.10 level, with and without proper controls for the method of randomization. As with Bruhn and McKenzie (2009), we see an increase in power in nearly all cases when proper controls for the method of randomization are included.

In Table 4, Panel B we see the D_A -optimal method improves power for all six datasets. The size of the improvements are modest with the given balancing variables, but improvement is consistent. The Block method on the other hand decreases power as often as it increases it. Table 5 shows that all methods, including simple randomization are indistinguishable as N grows large.

6 Experience from the field

We implemented the D_A -optimal method in a randomized trial of safe water and handwashing interventions among 435 compounds in slums of Dhaka, Bangladesh. To our knowledge, this study was the first to apply the D_A -optimal method. Details of the study and results are provided in Guiteras et al. (2014).

All participants received behavior change communication and a free trial of a "chlorine 12 See the notes to Tables 4 and 5 for the values of these imposed treatment effects.

dispenser," a compound-level device for treating water with chlorine (Evidence Action, 2014). At the end of the free trial, we measured the compound's willingness to pay for a one-year subscription for use and maintenance of the chlorine dispenser. We had 8 study arms in a 2-by-2-by-2 interaction of

- Behavior change message was a standard health message vs. messages based on disgust and shame;
- Included handwashing messages and soapy water bottle vs. not included; and
- Measured collective wilingness to pay for the compound vs. individual household willingness to pay.

Because we were especially interested in the effect of the disgust and shame treatment on handwashing in the handwashing arm, we gave handwashing 2/3 weight. The other treatments were equally weighted. Our balancing variables were the number of households in the compound and an indicator for the presence of gas burners connected to the municipal supply.

To implement the allocation, the enumerator collected baseline data on compound size and gas status from each eligible compound and transmitted these data to the field office, either by SMS or a phone call. The field supervisor then entered the covariates into a Stata program that assigned the compound to a treatment cell using the D_A -optimal method.

Manipulation was not likely in our context, because the enumerators could not plausibly have anticipated which assignment any given compound would receive. First, the enumerators collected several baseline variables, but did not know which would be used for stratification. Second, they did not have access to the in-progress list of compounds, covariates and assignments. Third, they were unfamiliar with the D_A -optimal algorithm. If manipulation is a concern, the biased-coin variant (Section 2.7) may be appealing.

The resulting sample was well-balanced on both the balancing variables and other plausibly important covariates that were not explicitly included in the randomization. See Tables A1-A4 of Guiteras et al. (2014) for detailed results. Compounds were well-balanced both on the mean of the (continuous) number of households in the compound and on an indicator for whether the compound had the median (8) or fewer number of households, even though we did not know in advance what the median number would be.

These results were obtained in spite of choosing one balancing variable poorly. In our piloting, we observed that water treatment practices varied importantly by whether the compound had a connection to the municipal gas supply. A gas connection also appeared to be a useful proxy for better overall socio-economic status. However, gas coverage in our study area turned out to be much higher than in the pilot area (even though the pilot area was nearby and similar in many other respects), and in fact was nearly universal (> 95%). Despite this unhelpful balancing variable, the algorithm produced a sample that was well-balanced on the other balancing variable and on our main SES variable, household monthly income. We view the robustness of the method as encouraging.

7 Discussion and Conclusion

The D_A -optimal method is a useful tool for improving balance in situations when a full ex-ante stratification is not feasible. The benefits are most pronounced when sample sizes are modest. Our field experience demonstrates that implementation is feasible and yields a well-balanced sample.

Our implementation required a qualified field supervisor to run the treatment assignment. Future trials could improve on our implementation by coding the algorithm into tablets or smartphones. With such an app, enumerators in the field would then be able to allocate treatments optimally in real time.

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Figure 1: Distribution of Differences in Means between Treatment and Control, at Baseline and Follow-up



(Sample size of 100)							
	Simple	Block	Block	Da Optimal	Da Optimal		
	Random	(2 variables)	(4 variables)	(2 variables)	(4 variables)		
Panel A. Average difference in BASELINE between treatment and control means (in SD)							
Microenterprise profits (Sri Lanka)	-0.003	0.001	-0.001	0.000	0.000		
Housing expenditure (Indonesia)	0.000	0.000	-0.001	0.000	0.000		
Labor income (Mexico)	-0.002	-0.002	0.000	0.000	0.000		
Height z-score (Pakistan)	0.000	-0.002	0.001	0.000	0.000		
Math test score (Pakistan)	-0.005	0.001	0.000	0.000	0.000		
Baseline unobservables (Sri Lanka)	0.000	0.000	0.000	0.000	0.000		
Baseline unobservables (Mexico)	0.000	-0.001	0.000	0.001	0.000		
Panel B. Ninety-fifth percentile of difference in BASELINE between treatment and control means (in SD)							
Microenterprise profits (Sri Lanka)	0.390	0.194	0.244	0.055	0.064		
Housing expenditure (Indonesia)	0.391	0.148	0.193	0.038	0.058		
Labor income (Mexico)	0.389	0.283	0.305	0.074	0.078		
Height z-score (Pakistan)	0.389	0.160	0.203	0.038	0.055		
Math test score (Pakistan)	0.396	0.164	0.237	0.039	0.055		
Baseline unobservables (Sri Lanka)	0.434	0.417	0.414	0.434	0.427		
Baseline unobservables (Mexico)	0.457	0.448	0.439	0.448	0.429		
Panel C. Proportion of p-values < 0.1 for testing difference in BASELINE means							
Microenterprise profits (Sri Lanka)	0.101	0.000	0.005	0.000	0.000		
Housing expenditure (Indonesia)	0.100	0.000	0.001	0.000	0.000		
Labor income (Mexico)	0.103	0.017	0.031	0.000	0.000		
Height z-score (Pakistan)	0.098	0.000	0.001	0.000	0.000		
Math test score (Pakistan)	0.106	0.000	0.006	0.000	0.000		
Baseline unobservables (Sri Lanka)	0.101	0.096	0.094	0.096	0.092		
Baseline unobservables (Mexico)	0.108	0.094	0.095	0.102	0.096		

Table 1--How Do The Different Methods Compare in Terms of Baseline Balance?

Notes: Statistics are based on 10000 simulations of each method.

(Sample size of 30)							
	Simple	Block	Block	Da Optimal	Da Optimal		
	Random	(2 variables)	(4 variables)	(2 variables)	(4 variables)		
Panel A. Average difference in FOLLON	W-UP betwe	en treatment an	d control means	(in SD)			
Microenterprise profits (Sri Lanka)	-0.001	0.004	-0.001	-0.001	-0.001		
Child schooling (Indonesia)	-0.003	0.001	0.000	0.008	-0.002		
Housing expenditure (Indonesia)	0.000	-0.004	0.000	-0.002	0.005		
Labor income (Mexico)	0.001	0.006	0.000	0.000	-0.001		
Height z-score (Pakistan)	-0.004	0.005	-0.005	-0.002	0.001		
Math test score (Pakistan)	-0.003	0.000	0.001	-0.001	-0.004		
Panel B. Ninety-fifth percentile of diffe	erence in FO	LLOW-UP betwe	en treatment and	d control means	(in SD)		
Microenterprise profits (Sri Lanka)	0.713	0.610	0.717	0.700	0.700		
Child schooling (Indonesia)	0.556	0.745	0.699	0.652	0.652		
Housing expenditure (Indonesia)	0.722	0.655	0.673	0.687	0.652		
Labor income (Mexico)	0.703	0.713	0.696	0.731	0.754		
Height z-score (Pakistan)	0.712	0.619	0.663	0.497	0.504		
Math test score (Pakistan)	0.716	0.441	0.641	0.515	0.526		
Panel C. Proportion of p-values < 0.1 for testing difference in FOLLOW-UP means with inference as if pure							
randomization was used (e.g., no adjustment for strata or balancing variables)							
Microenterprise profits (Sri Lanka)	0.103	0.054	0.105	0.101	0.105		
Child schooling (Indonesia)	0.050	0.110	0.088	0.084	0.059		
Housing expenditure (Indonesia)	0.105	0.074	0.080	0.087	0.071		
Labor income (Mexico)	0.101	0.103	0.092	0.117	0.128		
Height z-score (Pakistan)	0.102	0.056	0.073	0.015	0.016		
Math test score (Pakistan)	0.098	0.005	0.064	0.019	0.023		
Panel D. Proportion of p-values < 0.1 for testing difference in FOLLOW-UP means with inference which takes							
account of randomization method (i.e., controls for stratum, or balancing variables)							
Microenterprise profits (Sri Lanka)	0.103	0.084	0.085	0.111	0.109		
Child schooling (Indonesia)	0.095	0.120	0.087	0.087	0.101		
Housing expenditure (Indonesia)	0.104	0.099	0.126	0.102	0.100		
Labor income (Mexico)	0.101	0.105	0.082	0.115	0.122		
Height z-score (Pakistan)	0.105	0.093	0.148	0.101	0.095		
Math test score (Pakistan)	0.095	0.095	0.105	0.114	0.102		

Table 2--How Do The Different Methods Compare in Terms of Balance on Future Outcomes?

Notes: The coefficients in panels A and B are for specifications without controls for balancing

variables or stratum dummies. Statistics are based on 10000 simulations of each method.

(Sample size of 300)						
	Simple	Block	Block	Da Optimal	Da Optimal	
	Random	(2 variables)	(4 variables)	(2 variables)	(4 variables)	
Panel A. Average difference in FOLLO	W-UP betwe	een treatment an	d control means	(in SD)		
Microenterprise profits (Sri Lanka)	0.001	0.000	0.003	-0.002	0.000	
Child schooling (Indonesia)	0.000	0.002	0.001	-0.002	-0.001	
Housing expenditure (Indonesia)	0.001	0.000	0.001	-0.001	0.001	
Labor income (Mexico)	0.000	0.000	-0.001	-0.001	-0.002	
Height z-score (Pakistan)	0.001	0.001	0.000	-0.001	-0.001	
Math test score (Pakistan)	0.001	0.000	0.001	0.000	0.000	
Panel B. Ninety-fifth percentile of diffe	erence in FO	LLOW-UP betwe	en treatment and	d control means	(in SD)	
Microenterprise profits (Sri Lanka)	0.219	0.211	0.211	0.210	0.213	
Child schooling (Indonesia)	0.227	0.219	0.212	0.227	0.216	
Housing expenditure (Indonesia)	0.226	0.195	0.190	0.199	0.193	
Labor income (Mexico)	0.226	0.196	0.195	0.198	0.195	
Height z-score (Pakistan)	0.227	0.188	0.193	0.187	0.188	
Math test score (Pakistan)	0.224	0.182	0.183	0.175	0.174	
Panel C. Proportion of p-values < 0.1 for testing difference in FOLLOW-UP means with inference as if pure						
randomization was used (e.g., no adjustment for strata or balancing variables)						
Microenterprise profits (Sri Lanka)	0.095	0.086	0.086	0.085	0.091	
Child schooling (Indonesia)	0.115	0.087	0.083	0.111	0.088	
Housing expenditure (Indonesia)	0.099	0.059	0.050	0.060	0.054	
Labor income (Mexico)	0.101	0.058	0.058	0.062	0.058	
Height z-score (Pakistan)	0.099	0.048	0.054	0.046	0.047	
Math test score (Pakistan)	0.097	0.042	0.043	0.033	0.034	
Panel D. Proportion of p-values < 0.1 for testing difference in FOLLOW-UP means with inference which takes						
account of randomization method (i.e., controls for stratum, or balancing variables)						
Microenterprise profits (Sri Lanka)	0.095	0.109	0.139	0.103	0.112	
Child schooling (Indonesia)	0.103	0.101	0.107	0.113	0.098	
Housing expenditure (Indonesia)	0.098	0.100	0.100	0.099	0.096	
Labor income (Mexico)	0.099	0.100	0.094	0.119	0.117	
Height z-score (Pakistan)	0.097	0.098	0.100	0.103	0.101	
Math test score (Pakistan)	0.102	0.100	0.096	0.097	0.096	

Table 3--How Do The Different Methods Compare in Terms of Balance on Future Outcomes?

Notes: The coefficients in panels A and B are for specifications without controls for balancing

variables or stratum dummies. Statistics are based on 10000 simulations of each method.

(cample size of sec 7						
		Block	Da Optimal			
	Simple Random	(2 variables)	(2 variables)			
Panel A. Proportion of of p-values < 0.10 when no adjustment is made for method of randomization						
Microenterprise profits (Sri Lanka)	0.144	0.105	0.145			
Child schooling (Indonesia)	0.124	0.149	0.133			
Housing expenditure (Indonesia)	0.387	0.375	0.378			
Labor income (Mexico)	0.187	0.178	0.196			
Height z-score (Pakistan)	0.174	0.132	0.075			
Math test score (Pakistan)	0.155	0.050	0.081			
Panel B. Proportion of p-values < 0.10 when adjustment is made for randomization method						
Microenterprise profits (Sri Lanka)	0.144	0.142	0.154			
Child schooling (Indonesia)	0.122	0.134	0.135			
Housing expenditure (Indonesia)	0.396	0.415	0.407			
Labor income (Mexico)	0.176	0.171	0.191			
Height z-score (Pakistan)	0.248	0.199	0.257			
Math test score (Pakistan)	0.217	0.315	0.234			

Table 4--How Do The Different Methods Compare in Terms of Power in Detecting a Given Treatment Effect? (Sample size of 30)

Notes: Statistics are based on 10000 simulations of each method. Simulated treatment effects are as follows: Microenterprise profits—an Rs. 1,000 (LKR) increase in profits (about 25 percent of average baseline profits); child schooling—one in three randomly selected children in the treatment group who would have dropped out don't; household expenditure—an increase of 0.4 in In household expenditure per capita, which corresponds to about one-half of a standard deviation or moving a household from the twenty-fifth to the fiftieth percentile; labor income— a MEX\$920 increase in income (about 20 percent of average baseline income); height z-score—an increase of one quarter of a standard deviation in the z-score, where the z-score is defined as standard deviations from mean US height for age; math test score—an increase of one quarter of a standard deviation in the test score.

(
		Block	Da Optimal			
	Simple Random	(4 variables)	(4 variables)			
Panel A. Proportion of of p-values < 0.10 when no adjustment is made for method of randomization						
Microenterprise profits (Sri Lanka)	0.286	0.283	0.279			
Child schooling (Indonesia)	0.609	0.573	0.593			
Housing expenditure (Indonesia)	0.998	1.000	1.000			
Labor income (Mexico)	0.488	0.482	0.481			
Height z-score (Pakistan)	0.729	0.756	0.759			
Math test score (Pakistan)	0.622	0.656	0.653			
Panel B. Proportion of p-values < 0.10 when adjustment is made for randomization method						
Microenterprise profits (Sri Lanka)	0.303	0.346	0.305			
Child schooling (Indonesia)	0.602	0.599	0.607			
Housing expenditure (Indonesia)	1.000	1.000	1.000			
Labor income (Mexico)	0.589	0.541	0.581			
Height z-score (Pakistan)	0.862	0.849	0.853			
Math test score (Pakistan)	0.807	0.783	0.807			

Table 5--How Do The Different Methods Compare in Terms of Power in Detecting a Given Treatment Effect? (Sample size of 300)

Notes: Statistics are based on 10000 simulations of each method. Simulated treatment effects are as follows: Microenterprise profits—an Rs. 1,000 (LKR) increase in profits (about 25 percent of average baseline profits); child schooling—one in three randomly selected children in the treatment group who would have dropped out don't; household expenditure—an increase of 0.4 in In household expenditure per capita, which corresponds to about one-half of a standard deviation or moving a household from the twenty-fifth to the fiftieth percentile; labor income— a MEX\$920 increase in income (about 20 percent of average baseline income); height z-score—an increase of one quarter of a standard deviation in the z-score, where the z-score is defined as standard deviations from mean US height for age; math test score—an increase of one quarter of a standard deviation in the test score.