

Adaptive designs for sequential experiments*

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Received May 6, 2002; revision accepted June 21, 2002

Abstract: Various adaptive designs have been proposed and applied to clinical trials, bioassay, psychophysics, etc. Adaptive designs are also useful in high cost engineering trials. More and more people have been paying attention to these design methods. This paper introduces several broad families of designs, such as the play-the-winner rule, randomized play-the-winner rule and its generalization to the multi-arm case, doubly biased coin adaptive design, Markov chain model.

Key words: Clinical trial, Adaptive designs, (Randomized) Play-the-winner rule, Biased coin design, Markov chain, Asymptotic properties, Urn model.

Document code: A

CLC number: O211.4

INTRODUCTION

Traditional designs for clinical trials use the balanced (or 50-50) allocation of patients to treatments. For example, in a trial to compare experimental therapy (drug) to placebo (control), a standard feature of most designs is to distribute half of patients to each arm. It is reasonable that one may want to reduce the total number of failure outcomes in a trial, and keep the capability of making a comparison between experimental therapy and placebo as well. Hence the idea of adaptive designs has been proposed to serve the purpose.

Adaptive design, an important subdivision of experimental designs, are designs in which the probability a treatment assigned to the coming patient depends upon the results of the previous patients in the study. The goal is to show assignment probabilities to favor better treatment performance. This kind of design has also been applied to bioassay, psychophysics, etc. They are also useful in high cost engineering trials.

PLAY-THE-WINNER RULE AND RANDOMIZED PLAY-THE-WINNER RULE

Consider a two-arm clinical trial: two treat-

ments (1 and 2) with dichotomous response (success and failure). Patients are recruited into the clinical trial sequentially and respond immediately to treatments. Zelen (1969) propose the following design, which is well known as the play-the-winner (PW) rule: A success on a particular treatment generates a future trial on the same treatment with a new patient. A failure on a treatment generates a future trial on the alternate treatment. Let N_{n1} and N_{n2} be the number of the patients assigned to the treatment 1 and 2, respectively, in the first n stages. And let $p_i = P\{\text{success} \mid \text{treatment } i\}$ be the success probability of a patient on the treatment i , $q_i = 1 - p_i$, $i = 1, 2$. Then

$$\frac{N_{n1}}{n} \rightarrow \frac{q_2}{q_1 + q_2} \text{ a.s.}$$

and

$$\sqrt{n} \left(\frac{N_{n1}}{n} - \frac{q_2}{q_1 + q_2} \right) \xrightarrow{D} N(0, \sigma_{PW}^2),$$

where $\sigma_{PW}^2 = q_1 q_2 (p_1 + p_2) / (q_1 + q_2)^3$. This is first discussed in Zelen (1969).

As pointed out in Wei et al. (1978) and Wei (1979), the PW rule is too deterministic and is not applicable when we have delayed responses from patients to treatments. Motivated as

* Project supported by the National Natural Science Foundation of China (Nos. 19571021 and 10071072).

an extension to Zelen’s (1969) idea, Wei et al. (1978) introduced the following randomized play-the-winner (RPW) rule: We start with (α, α) balls (type 1 and 2 respectively) in the urn. If a type 1 ball is drawn, a patient is assigned to the treatment 1; If a type 2 ball is drawn, a patient is assigned to the treatment 2. The ball is replaced and the patient response is observed. A success on the treatment 1 or a failure on the treatment 2 generates a type 1 ball in the urn; A success on the treatment 2 or a failure on the treatment 1 generates a type 2 ball in the urn. The RPW rule may be regarded as a generalized Polya urn (GPU) model (Wei, 1979). Further, let $Y_{n1}(Y_{n2})$ be number of balls of type 1 (2) after n stage. From the results of Athreya et al. (1968), we have

$$\frac{Y_{n1}}{Y_{n1} + Y_{n2}} \rightarrow \frac{q_2}{q_1 + q_2} \text{ a.s. and}$$

$$\frac{N_{n1}}{n} \rightarrow \frac{q_2}{q_1 + q_2} \text{ a.s.}$$

When $p_1 + p_2 < 1.5$ (or $q_1 + q_2 > 0.5$), we have the following asymptotic normality:

$$\sqrt{n} \left(\frac{Y_{n1}}{Y_{n1} + Y_{n2}} - \frac{q_2}{q_1 + q_2} \right) \xrightarrow{D} N \left(0, \frac{q_1 q_2}{(2(q_1 + q_2) - 1)(q_1 + q_2)^2} \right) \tag{1}$$

and

$$\sqrt{n} \left(\frac{N_{n1}}{n} - \frac{q_2}{q_1 + q_2} \right) \xrightarrow{D} N(0, \sigma_{\text{RPW}}^2), \tag{2}$$

where

$$\sigma_{\text{RPW}}^2 = \frac{q_1 q_2 [5 - 2(q_1 + q_2)]}{[2(q_1 + q_2) - 1](q_1 + q_2)^2}.$$

The asymptotic normality was first given in Smythe et al. (1995). When $q_1 + q_2 < 0.5$, the limiting distributions of both the urn composition and the proportions of patients assigned to each treatments are unknown. The RPW rule is not deterministic and allows delayed responses by the patients.

If treatment 1 is "doing better", both PW rule and RPW rule are shown to favor treatment 1.

MULTI-ARM CASE RPW RULE AND GENERALIZED POLYA URN

signs can be developed from the GPU model. Consider an urn containing balls of K types. Initially, the urn contains $Y_0 = (Y_{01}, \dots, Y_{0K})$ balls, where Y_{0i} denotes the number of balls of type i , $i = 1, \dots, K$. A ball is drawn at random from the urn. Its type is observed and the ball is then replaced. At the n th stage, following a type i drawn, $D_{ij}(n)$ balls of type j , for $j = 1, \dots, K$, are added to the urn. In the most general sense, $D_{ij}(n)$ can be random and can be some function of a random process outside the urn process (in the case of adaptive designs, $D_{ij}(n)$ will be a random function of patient response). A ball must always be added at each stage (in addition to the replacement), and the expectation of the total numbers of balls added in each stage is the same (say γ), so

$$P \{D_{ij}(n) = 0, \text{ for all } j = 1, \dots, K\} = 0,$$

$$\sum_{j=1}^K E \{D_{ij}(n)\} = \gamma, \quad i, j = 1, \dots, K,$$

$$n = 1, 2, \dots.$$

Without losing generality, we can assume $\gamma = 1$. Let H_n be the matrix comprising element $\{h_{ij}(n) = E \{D_{ij}(n)\}\}$. We refer to D_n as the rules and H_n as the design matrices. If $H_n = H$ for all n , the model is said to be homogenous. In general, it is assumed that $H_n = H$. Let $Y_n = (Y_{n1}, \dots, Y_{nK})$, where Y_{ni} represents the number of balls in the urn of type i after n th stage, and $N_n = (N_{n1}, \dots, N_{nK})$, where N_{ni} represents the number of times a type i ball drawn in the first n draws. In the clinical trials, N_{ni} is the number of patients assigned to treatment i in the first n stages. Let $v = (v_1, \dots, v_K)$ be the left eigenvector corresponding to the largest eigenvalue of H with $v_1 + \dots + v_K = 1$. Then v_i is just the limiting proportion of both the patients assigned to treatment i and the type i balls in the urn, i.e.,

$$\frac{N_{ni}}{n} \rightarrow v_i \text{ a.s. and } \frac{Y_{ni}}{\sum_{j=1}^K Y_{nj}} \rightarrow v_i \text{ a.s.} \tag{3}$$

Athreya et al. (1968), Smyth (1996) and Bai et al. (1999; 2000*) showed the normality of Y_n

* Bai, Z. D. and Hu, F., 2000. Strong consistency and asymptotic normality for urn models.

and N_n : let $\lambda_1 = \gamma = 1$, $\lambda_2, \dots, \lambda_K$ be the eigenvalues of \mathbf{H} , and $\lambda = \max \{Re(\lambda_2), \dots, Re(\lambda_K)\}$, if $\lambda < 1/2$ and

$$\sum_{n=1}^{\infty} \frac{\|\mathbf{H}_n - \mathbf{H}\|}{\sqrt{n}} < \infty, \tag{4}$$

then

$$\sqrt{n} \left(\frac{\mathbf{Y}_n}{K} - \mathbf{v} \right) \xrightarrow{D} N(0, \boldsymbol{\Sigma}) \tag{5}$$

and

$$\sqrt{n} \left(\frac{N_n}{n} - \mathbf{v} \right) \xrightarrow{D} N(0, \boldsymbol{\Sigma}^*). \tag{6}$$

Under similar conditions, we showed that $\mathbf{Y}_n - n\mathbf{v}$ and $N_n - n\mathbf{v}$ can be approximated by Gaussian processes:

$$\mathbf{G}_{n1}\mathbf{H} + \mathbf{G}_{n2} \text{ and } \mathbf{G}_{n1} + \int_0^n \frac{\mathbf{G}_{x2}}{x} dx (\mathbf{I} - \mathbf{1}'\mathbf{v})$$

respectively (Bai et al., 2002c; 2001*), where \mathbf{G}_{it} are independent Gaussian processes which are solutions of the following type equation:

$$\mathbf{G}_t = \mathbf{W}_t + \int_0^t \frac{\mathbf{G}_s}{s} (\mathbf{H} - \mathbf{1}'\mathbf{v}) ds, \quad t > 0, \\ \mathbf{G}_0 = 0.$$

Here $\{\mathbf{W}(t)\}$ is a d -dimensional Brownian motion. In particular, the limiting combining distribution of \mathbf{Y}_n and N_n was obtained. For more results, one can refer to Bai et al. (2001a; 2001b) and Rosenberger (1996).

Define $X_n = i$ if a ball of type i is drawn at n th stage; $T_n = 1$ if the response of the n th patient is a "success", 0 otherwise. Assume that $P\{T_n = 1 | X_j = i\}$ is independent of n , hence we can write

$$p_i = P\{T_n = 1 | X_n = i\}, \quad q_i = 1 - p_i. \tag{7}$$

Now we give some important special cases.

Case 1. Let $\mathbf{H}^{(1)} = (h_{ij}^{(1)})_{i,j=1}^K$, where $h_{ii}^{(1)} = p_i$ and $h_{ij}^{(1)} = q_i / (K - 1)$ ($i \neq j$). Then v_i in Eq.(3) is

$$v_i^{(1)} = \frac{1/q_K}{\sum_{j=1}^K q_j}.$$

This model was proposed by Wei (1979), which means that, at the n th stage, if a patient

is assigned to treatment i and cured, then a type i ball is added to the urn, otherwise, if treatment i for a patient fails, then $\frac{1}{K - 1}$ balls are added to each of the other $K - 1$ treatments.

Case 2. Let $\mathbf{H}^{(2)} = (h_{ij}^{(2)})_{i,j=1}^K$, where $h_{ii}^{(2)} = p_i$ and $h_{ij}^{(2)} = q_i p_j / (M - p_i)$ ($i \neq j$) and $M = \sum_{j=1}^K p_j$. Then v_i in Eq. (3) is

$$v_i^{(2)} = \frac{p_i(M - p_i)/q_i}{\sum_{j=1}^K p_j(M - p_j)/q_j}.$$

This model was proposed by Bai et al. (2002b), which means that, at the n th stage, if a patient is assigned to treatment i and cured, then a type i ball is added into the urn, otherwise, if treatment i for the patients fails, then a number of balls proportional to the success rates of the other $K - 1$ treatments are added into the urn. But this design is not practicable, since the success probabilities p_j are unknown. They should be replaced by their sample estimators. This leads to the following case.

Case 3. Let $\mathbf{S}_n = (S_{n1}, \dots, S_{nK})$, where S_{ni} denotes the number of successes of the i th treatment in the N_{ni} trials, $r_{ni} = (S_{ni} + 1) / (N_{ni} + 1)$ and $M_{n-1} = \sum_{j=1}^K r_{n-1,j}$. And let

$$\mathbf{H}^{(3)} = \mathbf{H}_n^{(3)} = \begin{pmatrix} p_1 & \frac{r_{n-1,2}}{M_{n-1} - r_{n-1,1}} q_1 & \dots & \frac{r_{n-1,K}}{M_{n-1} - r_{n-1,1}} q_1 \\ \frac{r_{n-1,1}}{M_{n-1} - r_{n-1,2}} q_2 & p_2 & \dots & \frac{r_{n-1,K}}{M_{n-1} - r_{n-1,2}} q_2 \\ \dots & \dots & \dots & \dots \\ \frac{r_{n-1,1}}{M_{n-1} - r_{n-1,K}} q_K & \frac{r_{n-1,2}}{M_{n-1} - r_{n-1,K}} q_K & \dots & p_K \end{pmatrix}$$

Case 4. The success probabilities defined in Eq.(7) are homogeneous for different stages (patients). But this assumption is not always realistic in certain situations, where patients may exhibit drift in characteristics over time, i.e.

$$p_i(n) = P\{T_n = 1 | X_n = i\}, \\ q_i(n) = 1 - p_i(n). \tag{8}$$

It is natural to assume that the treatments are sta-

* Bai, Z. D., Hu, F. and Zhang L. X., 2001. The weak and strong approximation for generalized Friedman's urn model.

ble in such case, i.e., it is assumed that $p_i(n) \rightarrow p_i, i = 1, \dots, K$. In such case, the design in Case 1 becomes $\mathbf{H}^{(4)} = \mathbf{H}_n^{(4)} = (h_{ij}^{(4)})_{i,j=1}^K$, where $h_{ii}^{(4)} = p_i(n)$ and $h_{ij}^{(4)} = q_i(n)/(K-1)(i \neq j)$.

Both $\mathbf{H}^{(1)}$ and $\mathbf{H}^{(2)}$ are independent of n , the stage number, and so the models are homogeneous. But $\mathbf{H}^{(3)}$ and $\mathbf{H}^{(4)}$ are dependent on n , and so the models are non-homogeneous.

Remark 1. When $K = 2$, the above three adaptive designs are all governed by the RPW rule. So all of these designs are extensions of the RPW rule for K treatments.

Remark 2. For designs $\mathbf{H}^{(1)}$ and $\mathbf{H}^{(2)}$, when $K > 2$, notice that for any $i \neq j, 1 \leq i, j \leq K$, if $p_i > p_j$, we have

$$\frac{v_i^{(2)}}{v_j^{(2)}} - \frac{v_i^{(1)}}{v_j^{(1)}} = \frac{p_i(M - p_i)/q_i}{p_j(M - p_j)/q_j} - \frac{1/q_i}{1/q_j} = \frac{(p_i - p_j)(M - p_i - p_j)/q_i}{p_j \sum_{k \neq j} p_k/q_k} > 0.$$

The design $\mathbf{H}^{(2)}$ is more reasonable than the design $\mathbf{H}^{(1)}$ because more patients will be assigned to a better treatment.

Remark 3. Let $a \geq 0$. An extension of the design $\mathbf{H}^{(3)}$ has $r_{n-1,j}^a$ and $M_{n-1}(a) = \sum_{j=1}^K r_{n-1,j}^a$ instead of $r_{n-1,j}$ and M_{n-1} respectively. Taking $a = 0$ and 1 yields designs $\mathbf{H}^{(1)}$ and $\mathbf{H}^{(3)}$ respectively.

Designs dependent on estimated unknown parameters: The asymptotic normality of Case 1 and Case 2 follows from Eqs. (5) and (6) immediately. In the Case 4, if

$$\sum_{n=1}^{\infty} \frac{|p_i(n) - p_i|}{\sqrt{n}} < \infty, i = 1, \dots, K, \quad (9)$$

then the condition (4) is satisfied, and so asymptotic properties also follow. But the model in Case 3 can not satisfy the condition (4), since the fastest convergence rate of \mathbf{H}_n is that

$\|\mathbf{H}_n - \mathbf{H}\| = O(\sqrt{n})$ in probability. The asymptotic normalities of such case were studied in Hu and Zhang*. We considered the general model with design matrices of the type $\mathbf{H}_n = \mathbf{H}(\hat{\Theta}_n)$ dependent on the estimated unknown parameter, where $\hat{\Theta}_n$ is the sample estimator of the unknown parameter Θ . Strong consistency and the asymptotic normalities are established for both the urn

composition and the number of patients assigned to each treatment. It should be mentioned that the condition (4) cannot be reduced to $\|\mathbf{H}_n - \mathbf{H}\| = O(\sqrt{n})$ in general.

Designs with delay responses: Typically, clinical trials do not result in immediate outcomes, i.e., individual patient outcomes may not be immediately available prior to the randomization of the next patient. Consequently, the urn cannot be updated immediately, but can be updated only when the outcomes become available. Fortunately, it is verified that stochastic staggered entry and delay mechanisms do not affect the asymptotic properties of both the urn composition \mathbf{Y}_n and the sample fractions \mathbf{N}_n for a wide class of designs defined by GPU (Bai et al., 2002a).

DOUBLY BIASED COIN ADAPTIVE DESIGNS

We come back to the PW rule and RPW rule. As it is known, the PW rule is too deterministic and is not applicable when we have delayed responses from patients of treatments. The RPW rule and its generalizations seem solve this problem. However, in using the RPW rule, when $q_1 + q_2 < 0.5$, the limiting distributions of the proportions of patients assigned to each treatments are unknown. But in practice, both q_1 and q_2 are usually very small. So, the RPW rule is not practical in such cases. The asymptotic variation of the proportion becomes a big problem in using the RPW rule (even the adaptive designs based on the GPU model) for $q_1 + q_2 < 0.5$. Even in the case that $q_1 + q_2 > 0.5$, if $q_1 + q_2$ is near 0.5, σ_{RPW} is much larger than σ_{PW} . That is to say, the RPW rule is too random so that the asymptotic variance of proportion of patients assigned to each treatment is very large when the cure rates are large and so it is much less stable than the PW rule. Also, in using the multi-arm RPW, the condition that $\lambda < 1/2$ is very hard to check. Even in the 3-arm case, it is very difficult to check this condition.

Now, with keeping the desired allocation pro-

* Hu, F. and Zhang, L. X., 2001. The asymptotic theorems of adaptive designs for clinical trials with generations depending on the estimated success rates.

portions $v_1 = q_2/(q_1 + q_2)$ and $v_2 = q_1/(q_1 + q_2)$, just as in the case of the PW rule and RPW rule, our goal is to reduce the asymptotic variance. A natural way is as follows. At the $(n + 1)$ th stage, we assign a patient to a certain treatment by comparing the value N_{n1}/n with v_1 , or N_{n2}/n with v_2 . If N_{n1}/n is larger than v_1 , then we assign a patient to the treatment 1 with a probability less than v_1 ; If N_{n1}/n is less than v_1 , then we assign a patient to the treatment 1 with a probability larger than v_1 ; If N_{n1}/n equals v_1 , then we assign a patient to the treatment 1 with probability v_1 and to the treatment 2 with probability v_2 . By choosing suitable function, we may minimize the asymptotic variance. However, v_1 and v_2 are unknown, so they should be replaced by their estimators based on the sample of the previous n stages. So, the following adaptive design of clinical trial is considered and proposed.

At the first stage, a patient is assigned to each treatment with the same probability 1/2. After m assignments, we let S_{mk} be the number of successes of all the N_{mk} patients on the treatment k in the first m assignments, $k = 1, 2$, as usual. And let $\hat{p}_{mk} = \frac{S_{mk} + 1/2}{N_{mk} + 1}$ be the sample estimation of p_k , and write $\hat{q}_{mk} = 1 - \hat{p}_{mk}$, $k = 1, 2$. At the $(m + 1)$ th stage, the $(m + 1)$ th patient is assigned to the treatment 1 with probability $g(\frac{N_{m1}}{m}, \hat{v}_{m1})$, and to the treatment 2 with probability $1 - g(\frac{N_{m1}}{m}, \hat{v}_{m1})$, where $\hat{v}_{m1} = \frac{\hat{q}_{m2}}{\hat{q}_{m1} + \hat{q}_{m2}}$ is the sample estimation of $v_1 = q_2/(q_1 + q_2)$. The functions $g(x, \rho)$ is called allocation rule. A large class of functions can be chosen as an allocation rule. If it is one of the following forms:

$$g(x, \rho) = 0 \vee \left(\frac{\rho \exp\{\alpha(\frac{\rho}{x} - 1)\} + 1 - (1 - \rho)\exp\{\alpha(\frac{1 - \rho}{1 - x} - 1)\}}{2} \right) \wedge 1,$$

$$g(x, \rho) = 0 \vee \left(\frac{\rho(\frac{\rho}{x})^\alpha + 1 - (1 - \rho)(\frac{1 - \rho}{1 - x})^\alpha}{2} \right) \wedge 1,$$

$$g(0, \rho) = 1, \quad g(1, \rho) = 0,$$

$$g(x, \rho) = \frac{\rho(\frac{\rho}{x})^\alpha}{\rho(\frac{\rho}{x})^\alpha + (1 - \rho)(\frac{1 - \rho}{1 - x})^\alpha},$$

where $\alpha \geq 0$, we have that

$$\frac{N_{n1}}{n} - v_1 = O\left(\sqrt{\frac{\log \log n}{n}}\right) \quad a.s. \quad \text{and}$$

$$\sqrt{n}\left(\frac{N_{n1}}{n} - v_1\right) \xrightarrow{D} N(0, \sigma_{DAD}^2)$$

where

$$\sigma_{DAD}^2 = \sigma_\alpha^2 := \frac{q_1 q_2 (p_1 + p_2)}{(q_1 + q_2)^3} + \frac{2q_1 q_2}{(1 + 2\alpha)(q_1 + q_2)^3}$$

It should be noted that the asymptotic normality holds for all $0 < p_1 < 1$ and $0 < p_2 < 1$ (Hu et al. 2003). It is easily seen that σ_α^2 is a strictly monotonous decreasing function of $\alpha \geq 0$ and $\sigma_\alpha^2 \rightarrow \sigma_{PW}^2$ as $\alpha \rightarrow +\infty$. Also, $\sigma_\alpha^2 < \sigma_{RPW}^2$ for all $\alpha > 1$, whenever $q_1 + q_2 > 1/2$. Furthermore, if $q_1 + q_2$ is near 1/2, then σ_α is much smaller than σ_{RPW} . So, this adaptive design is more stable than the RPW rule. The larger α is, the more stable and less random is the design. So, this design is a compromise between the stability in the PW rule and the randomization in the RPW rule. Even when α is very large, it keeps enough randomization to avoid determinism. Such design can keep the spirit of the RPW rule in that it assigns more patients to the better treatment and allows delayed responses by the patients.

In such kind of designs, the assignments are adapted by both the results of responses and the current proportions of patients assigned, and its original idea came from Efron's (1971) biased coin design. So, they are called doubly adaptive biased coin design, first introduced by Eisele (1994) and Eisele et al. (1995) in the two-arm case with the responses which are from standard exponential families. However, the results of Eisele (1994) and Eisele et al. (1995) are not applicable since their condition (v) is impossible to be satisfied, because, if q_1 and q_2 are small, both v_1 and $v_2 = 1 - v_1$ should be very large by their (v). But this condition is a key in their proving methods.

Hu et al. (2003) studied the general multi-arm case, where the condition that $\lambda < 1/2$ in using GPU model is no longer a problem.

MARKOV CHAIN ADAPTIVE DESIGNS

In this section, we propose another class of adaptive designs, the so-called Markov chain adaptive design. Consider the two-arm case first. Suppose that at the stage n , the treatment 1 is assigned to the n th patient. Then the $(n + 1)$ th patient will be assigned either treatment 1 or treatment 2 according certain probabilities, which depend on the response of the n th patient. Let α_s be the probability of assigning the $(n + 1)$ th patient to treatment 1, when the response of the n th patient to treatment 1 is "success", and let α_f be the probability of assigning the $(n + 1)$ th patient to treatment 1, when the response of the n th patient to treatment 1 is "failure". Similarly define β_s and β_f with treatment 2 instead of treatment 1 in the definitions of α_s and α_f . When $\alpha_s = 1, \alpha_f = 0, \beta_s = 1, \beta_f = 0$, we get Zelen's PW rule. We may choose the parameters $\alpha_s, \alpha_f, \beta_s$ and β_f for different goals.

Define $X_n = 1$ or 0 if n th patient is assigned to the treatment 1 or 2, respectively. Then $N_{n1} = \sum_{j=1}^n X_j$ is the number of patients assigned to treatment 1 in the first n stages. Let $p_1(n) = P\{\text{success} | X_n = 1\}$ and $p_2(n) = P\{\text{success} | X_n = 0\}$ and $\alpha_n = p_1(n)\alpha_s + (1 - p_1(n))\alpha_f, \beta_n = p_2(n)\beta_s + (1 - p_2(n))\beta_f$. Then $\{X_n\}$ is a Markov chain with the transition probability matrix

$$P_n = \begin{pmatrix} \alpha_n & 1 - \beta_n \\ 1 - \alpha_n & \beta_n \end{pmatrix}.$$

When $p_1(n) = p_1$ and $p_2(n) = p_2$ for all n , $\{X_n\}$ is homogeneous. For this case we can write $\alpha_n = \alpha, \beta_n = \beta$, and we have that

$$\frac{N_{n1}}{n} \xrightarrow{P} \mu \text{ and } \sqrt{n} \left(\frac{N_{n1}}{n} - \mu \right) \xrightarrow{D} N(0, \sigma^2) \quad (10)$$

where

$$\mu = \frac{1 - \beta}{2 - \alpha - \beta}, \sigma^2 = \frac{(1 - \alpha)(1 - \beta)(\alpha + \beta)}{(2 - \alpha - \beta)^2}$$

(Bai et al., 2001*). For non-homogeneous case we have similar results under the condition of type Eq. (9). Furthermore, we established the strong consistency and strong approximation of

N_{n1} :

$$N_{n1} - n\mu - \sigma W(n) = O((n \log \log n)^{1/4} (\log n)^{1/2}) \text{ a.s.},$$

where $\{W(t)\}$ is a standard Brownian motion (Lin et al. 2001**).

For the multi-arm case, we let $\mathbf{X}_n = (X_{n1}, \dots, X_{nK})$, where $X_{ni} = 1$ if the n th patient is assigned to treatment i , and let $\mathbf{N}_n = \sum_{k=1}^n \mathbf{X}_k$ be the number of patients assigned to each treatment at the first n stages, as usual. Let $p_i(n) = P\{\text{success} | X_{ni} = 1\}$. Assume the transition probability matrix of the Markov chain $\{\mathbf{X}_n\}$ is $\mathbf{H}_n = \{H_{ij}(n)\}$ which is a function of $p_i(n), i = 1, \dots, K$, i. e., $\mathbf{E}[\mathbf{X}_{n+1} | \mathbf{X}_n] = \mathbf{X}_n \mathbf{H}_n$. We showed that \mathbf{N}_n can be approximated by a multi-dimensional Brownian motion:

$$\mathbf{N}_n - n\mathbf{v} - \mathbf{W}(n)\mathbf{\Sigma} = o(n^{1/2}) + O\left(\sum_{k=1}^n \|\mathbf{H}_k - \mathbf{H}\|\right) \text{ a.s.}$$

where $\mathbf{v} = (v_1, \dots, v_K)$ is the left eigenvector corresponding to the largest eigenvalue of \mathbf{H} with $v_1 + \dots + v_K = 1, \{\mathbf{W}(t)\}$ is a K -dimensional standard Brownian motion. In particular, we have asymptotical normality, if

$$\sum_{k=1}^n \|\mathbf{H}_k - \mathbf{H}\| = o(n^{1/2}). \quad (11)$$

Condition (11) is usual satisfied if $\sum_{k=1}^n |p_i(k) - p_i| = o(n^{1/2}), i = 1, \dots, K$, which is implied by (9).

If \mathbf{H}_n depends on estimated unknown parameters, the problem becomes more complicated. Zhang*** studied such case.

Remark 4. In Section 2, we mentioned the normality in Eqs. (1) and (2), but the condition $p_1 + p_2 < 3/2$ was imposed. But for normality in Eq. (10), this restriction is cancelled. In the multi-arm case, the restriction that $\lambda < 1/2$ is also cancelled.

* Bai, Z. D., Chen, Y. M., Hu, F. and Lin, Z. Y., 2001. Adaptive designs based on Markov chains (I) (II).

** Lin, Z. Y. and Zhang L. X., 2001. Strogan approximation of a non-homogenous markov chain.

*** Zhang, L. X., 2002. A kind of multi-treatment adaptive designs with assignment probabilities depending on the estimated parameters.

Remark 5. Keeping μ we can choose α and β to minimize σ^2 so that we get more stable Markov chain adaptive designs. Also, if α and β was chosen suitably, the Markov chain adaptive designs can also keep both randomization and stability.

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