

## STATISTICAL SELECTION OF THE BEST SYSTEM

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### ABSTRACT

This tutorial discusses some statistical procedures for selecting the best of a number of competing systems. The term “best” may refer to that simulated system having, say, the largest expected value or the greatest likelihood of yielding a large observation. We describe various procedures for finding the best, some of which assume that the underlying observations arise from competing normal distributions, and some of which are essentially nonparametric in nature. In each case, we comment on how to apply the above procedures for use in simulations.

### 1 INTRODUCTION

Experiments are often performed to compare two or more system designs in order to determine which scenario is the best. The statistical methods of *screening*, *selection*, and *multiple comparisons* are applicable when we are interested in making comparisons among a finite, possibly large, number of scenarios. The particular method that is appropriate depends on the type of comparison desired and properties of the data under study. For instance, are we interested in comparing means or quantiles? Are the available data independent or correlated within and/or among systems?

In this review, the term “best” may refer to that simulated system having, say, the largest expected value or the greatest likelihood of yielding a large observation. We will typically, but not always, regard the best population as the one having the largest expected value.

We describe a number of procedures for finding the best, some of which assume that the underlying observations arise from competing normal distributions, and some of which are essentially nonparametric in nature. In each case, we comment on how to apply the above procedures for use in simulations.

To get things going, the next section will give some additional low-level background on screening, selection, and multiple comparisons procedures. Section 3 establishes relevant notation and ground rules, while Section 4 presents

some very basic methods for purposes of motivating the upcoming procedures. Section 5 discusses three normal means procedures for selecting the best (or nearly the best) scenario. Of the normal procedures, the first is a screen-and-select procedure for finding the population with the largest expected value; in this procedure, inferior competitors are screened out after an initial stage of sampling. The second is a sequential procedure that can eliminate inferior choices at any stage and uses the common random numbers variance reduction technique in which we intentionally induce positive correlation between scenarios. The third is an efficient two-stage procedure that also uses common random numbers. Section 6 deals with three nonparametric procedures. Of these procedures, the first is a single-stage procedure for finding the most probable multinomial cell, the second is sequential, and the third is a clever augmentation that makes more efficient use of the underlying observations. We give conclusions in Section 7.

There are a number of general references for the interested reader in this area of selection of the best. Gibbons, Olkin, and Sobel (1977) and Bechhofer, Santner, and Goldsman (1995) give presentations from a statistical point of view, while Goldsman and Nelson (1998) and Kim and Nelson (2005a) devote a great deal of effort to the simulation side of the story.

### 2 BACKGROUND

We will usually assume that the observations coming from a particular scenario are independent and identically distributed (i.i.d.). Since this is never the case when dealing with simulation output (which is, for instance, almost always serially correlated), we will make appropriate comments to show how to apply the above procedures for use in simulations.

What are screening, selection, and multiple comparisons procedures? Screening and selection procedures (SSPs) are statistical methods designed to find the “best” (or “nearly the best”) system from among a collection of competing alter-

natives. For example, such procedures could be efficacious in any of the following practical situations:

- Find the normal population with the largest mean: A manufacturer would like to know which of three potential plant layouts will maximize expected revenues.
- Find the most probable multinomial cell: A polling service wishes to determine the most popular candidate before a certain election.
- Find the Bernoulli population having the largest success parameter: A medical research team conducts a clinical study comparing the success rates of five different drug regimens for a particular disease.

Informally speaking, SSPs are used to

- *screen* the competitors in order to find a small subset of those systems that contains the best system (or at least a “good” one).
- *select* outright the best system.

In practice, we could invoke a screening procedure to pare down a large number of alternatives into a palatable number; at that point, we might use a selection procedure to make the more fine-tuned choice of the best. Provided that certain assumptions are met, SSPs usually guarantee a user-specified probability of advertised performance—i.e., with high probability, a screening procedure will choose a subset containing the best (or a good) alternative, and a selection procedure will pick the best.

Multiple-comparison procedures (MCPs) treat the comparison problem as an inference problem on the performance parameters of interest. MCPs account for the error that arises when making simultaneous inferences about differences in performance among the systems. Usually, MCPs report to the user simultaneous confidence intervals for the differences. Recent research has shown that MCPs can be combined with SSPs for a variety of problems—including the manufacturing, medical, and polling examples outlined above. In fact, the field has progressed steadily over the last twenty years, particularly in the simulation community.

What is especially nice about SSPs and MCPs is that they are relevant, easily adaptable, and statistically valid in the context of computer simulation because the assumptions behind the procedures can frequently be satisfied: For example, these procedures sometimes require normality of the observations, an assumption that can often be secured by batching large numbers of (cheaply generated) outputs. Independence can be obtained by controlling random-number assignments. And multiple-stage sampling—which is required by some methods—is feasible in computer simulation because a subsequent stage can be initialized simply by retaining the final random-number seeds from the preceding

stage. As a bonus, it is possible to enhance (in a theoretically rigorous way) the performance of some of the procedures through the use of common random numbers, a popular variance reduction technique sometimes used in simulation.

### 3 SOME NOTATION

To facilitate what follows we define some notation: Let  $Y_{ij}$  represent the  $j$ th simulation output from system design  $i$ , for  $i = 1, 2, \dots, k$  alternatives and  $j = 1, 2, \dots$ . For fixed  $i$ , we will always assume that the outputs from system  $i$ ,  $Y_{i1}, Y_{i2}, \dots$ , are i.i.d. These assumptions are plausible if  $Y_{i1}, Y_{i2}, \dots$  are outputs across independent replications, or if they are appropriately defined batch means from a single replication after accounting for initialization effects. Let  $\mu_i = E[Y_{ij}]$  denote the expected value of an output from the  $i$ th system, and let  $\sigma_i^2 = \text{Var}[Y_{ij}]$  denote its variance. Further, let

$$p_i = \Pr \left\{ Y_{ij} > \max_{\ell \neq i} Y_{\ell j} \right\}$$

be the probability that  $Y_{ij}$  is the largest of the  $j$ th outputs across all systems when  $Y_{1j}, Y_{2j}, \dots, Y_{kj}$  are mutually independent.

The methods we describe make comparisons based on either  $\mu_i$  or  $p_i$ . Although not a restriction on either SSPs or MCPs, we will only consider situations in which there is no known functional relationship among the  $\mu_i$  or  $p_i$  (other than  $\sum_{i=1}^k p_i = 1$ ). Therefore, there is no potential information to be gained about one system from simulating the others—such as might occur if the  $\mu_i$  were a function of some explanatory variables—and no potential efficiency to be gained from fractional-factorial experiment designs, group screening designs, etc.

### 4 MOTIVATION AND APPROACHES

In this section, we begin with a motivational example and then go over the approaches that are relevant to the goal of selecting the best. Sections 5 and 6 outline procedures that are appropriate for real-world application.

**Example 1** *Simulation models of 25 different inventory policies have been developed for potential implementation at a large distribution/warehouse center. The single measure of system performance is the expected profit achieved while a particular policy is in effect. Differences between different policies’ expected profits of less than about \$10,000 are considered practically equivalent.*

We discuss briefly the three approaches employed here—subset selection (screening), indifference-zone selection (choosing the single best), and multiple comparisons (inference). For purposes of motivation, the simple procedures outlined in this section will all assume that the underlying data are normally distributed.

#### 4.1 Subset Selection

The subset selection approach is a screening device that attempts to select a (random-size) *subset* of the  $k = 25$  competing designs of Example 1 that contains the design with the greatest expected profit. Gupta (1956, 1965) proposed a single-stage procedure for this problem that is applicable in cases when the data from the competing designs are balanced (i.e., have the same number of observations from each contender) and are normal with common (unknown) variance  $\sigma^2$ . Nelson, et al. (2001) handle more general cases—in particular, that in which the unknown variances  $\sigma_i^2$ ,  $i = 1, 2, \dots, k$ , are not necessarily equal.

It is worthwhile to illustrate the concept of subset selection on the simplest possible case—that in which the observations  $Y_{ij}$  are independent and normally distributed with known, common variance  $\sigma^2$ . For ease of exposition, suppose that (unknown to us),  $\mu_k \geq \mu_{k-1} \geq \dots \geq \mu_1$ , so that system  $k$  is the best system. Further suppose that we have  $n$  outputs from each of the systems. Our goal is to use this data to obtain a subset  $I \subseteq \{1, 2, \dots, k\}$  such that

$$\Pr\{k \in I\} \geq 1 - \alpha, \quad (1)$$

where  $1/k < 1 - \alpha < 1$  for some user-specified  $\alpha$ . Ideally  $|I|$  is small, the best case being  $|I| = 1$ . Gupta's solution was to include in the set  $I$  all systems  $\ell$  such that

$$\bar{Y}_\ell(n) \geq \max_{i \neq \ell} \bar{Y}_i(n) - h\sigma\sqrt{\frac{2}{n}}, \quad (2)$$

where  $\bar{Y}_i(n)$  is the sample mean of the (first)  $n$  outputs from system  $i$ , and  $h$  is a constant whose value will depend on  $k$  and  $1 - \alpha$ . The proof that rule (2) provides guarantee (1) is instructive and shows what the value of  $h$  should be:

$$\begin{aligned} \Pr\{k \in I\} &= \Pr\left\{\bar{Y}_k(n) \geq \max_{i \neq k} \bar{Y}_i(n) - h\sigma\sqrt{\frac{2}{n}}\right\} \\ &= \Pr\left\{\frac{\bar{D}_{ik}(n) - \mu_{ik}}{\sigma\sqrt{2/n}} \leq h - \frac{\mu_{ik}}{\sigma\sqrt{2/n}}, \forall i \neq k\right\} \\ &\geq \Pr\{Z_i \leq h, i = 1, 2, \dots, k-1\} = 1 - \alpha, \end{aligned}$$

where we define  $\bar{D}_{ik}(n) = \bar{Y}_i(n) - \bar{Y}_k(n)$  and  $\mu_{ik} = \mu_i - \mu_k$  for all  $i \neq k$ , and where  $(Z_1, Z_2, \dots, Z_{k-1})$  have a multivariate normal distribution with means 0, variances 1, and common pairwise correlations  $1/2$ . Therefore, to provide the guarantee (1),  $h$  needs to be the  $1 - \alpha$  quantile of the maximum of such a multivariate normal random vector, a quantile that turns out to be relatively easy to approximate numerically (see, e.g., Bechhofer, Santner, and Goldsman 1995) or via simulation. Notice the inequality in the final step where we make use of the fact that  $\mu_k \geq \mu_i$ .

A theme that runs throughout much of the literature is to first use appropriate standardization of estimators and then bound the resulting probability statements in such a way that a difficult multivariate probability statement becomes one that is readily solvable.

#### 4.2 Indifference-Zone Selection

A disadvantage of the subset-selection procedure in Section 4.1 is that the retained set  $I$  may, and likely will, contain more than one system. However, there is no procedure that can guarantee a subset of size 1 and satisfy (1) for arbitrary  $n$ . Even when  $n$  is under our control, as it is in computer simulation, the appropriate value will depend on the true differences  $\mu_k - \mu_i, \forall i \neq k$ . To attack this problem, Bechhofer (1954) suggested the following compromise: guarantee to select the single best system,  $k$ , whenever  $\mu_k - \mu_{k-1} \geq \delta$ , where  $\delta > 0$  is the smallest difference the experimenter feels is worth detecting—this “practically significant” difference is called the indifference parameter.

Specifically, the procedure should guarantee

$$\Pr\{\text{select system } k | \mu_k - \mu_{k-1} \geq \delta\} \geq 1 - \alpha, \quad (3)$$

where  $1/k < 1 - \alpha < 1$ . If there are systems whose means are within  $\delta$  of the best, then the experimenter is “indifferent” to which of these is selected, leading to the term *indifference-zone* (IZ) formulation. In the motivational example, the indifference parameter is  $\delta = \$10000$ . Law and Kelton (2000) describe a number of IZ procedures that have proven useful in simulation, while Bechhofer, Santner, and Goldsman (1995) provide a comprehensive review of SSPs to that date.

Bechhofer's procedure is as follows: From each system, take

$$n = \left\lceil \frac{2h^2\sigma^2}{\delta^2} \right\rceil \quad (4)$$

outputs, where  $h$  is an appropriate constant (determined below) and  $\lceil x \rceil$  means to round  $x$  up; then select the system with the largest sample mean as the best. Assuming  $\mu_k - \mu_{k-1} \geq \delta$  and recalling our definitions  $\bar{D}_{ik}(n) =$

$\bar{Y}_i(n) - \bar{Y}_k(n)$  and  $\mu_{ik} = \mu_i - \mu_k$  for all  $i \neq k$ , we have

$$\begin{aligned} \Pr\{\text{select } k\} &= \Pr\{\bar{Y}_k(n) > \bar{Y}_i(n), \forall i \neq k\} \\ &= \Pr\left\{\frac{\bar{D}_{ik}(n) - \mu_{ik}}{\sigma\sqrt{2/n}} < -\frac{\mu_{ik}}{\sigma\sqrt{2/n}}, \forall i \neq k\right\} \\ &\geq \Pr\left\{\frac{\bar{D}_{ik}(n) - \mu_{ik}}{\sigma\sqrt{2/n}} < \frac{\delta}{\sigma\sqrt{2/n}}, \forall i \neq k\right\} \\ &\geq \Pr\left\{\frac{\bar{D}_{ik}(n) - \mu_{ik}}{\sigma\sqrt{2/n}} < h, \forall i \neq k\right\} \\ &= \Pr\{Z_i < h, i = 1, 2, \dots, k-1\} = 1 - \alpha, \end{aligned}$$

where again  $(Z_1, Z_2, \dots, Z_{k-1})$  has a multivariate normal distribution with means 0, variances 1, and common pairwise correlations  $1/2$ , implying  $h$  needs to be the  $1 - \alpha$  quantile of the maximum of such a multivariate normal random vector. Notice that the first inequality results from the assumption that  $\mu_k - \mu_{k-1} \geq \delta$ , while the second occurs because  $\sqrt{n} \geq \sqrt{2}h\sigma/\delta$ . Both of these tricks are standard: the first frees the probability statement of dependence on the true means, while the second frees it of dependence on the value of the variance.

It is worth noting that, over all configurations of the true means such that  $\mu_k - \mu_{k-1} \geq \delta$ , the configuration  $\mu_i = \mu_k - \delta$ , for all  $i \neq k$ , minimizes the probability of correct selection; it is therefore known as the *least-favorable configuration* (LFC). In this paper we break from the statistics literature in that we will not be concerned with identifying the LFC; our only interest is insuring that (3) is met.

Bechhofer's procedure is essentially a power calculation: how large a sample is required to detect differences of at least  $\delta$ ? When true differences are greater than  $\delta$ , Bechhofer's  $n$  may be much larger than needed. By taking observations and making decisions *sequentially*, it is often possible to reach an earlier decision. Although the roots of sequential selection procedures can be traced back at least to Wald (1947), the first procedure directly designed for selection purposes is due to Paulson (1964). That procedure takes observations *fully sequentially*—meaning one at a time—and *eliminates* systems from continued sampling when it is statistically clear that they are inferior. Thus, a problem with a single dominant alternative may terminate very quickly. See Section 5.2.1 for a simple sequential procedure that is easily adopted for use in simulation.

### 4.3 Multiple Comparisons

MCPs approach the comparison problem by providing simultaneous confidence intervals on selected differences among the systems' parameters. Hochberg and Tamhane (1987) and Hsu (1994, 1996) are good references on the topic. As noted by Hsu (1996, pp. 100–102), the connection between SSPs

and MCPs comes through the problem of multiple comparisons with the best (MCB). MCB forms simultaneous confidence intervals for  $\mu_i - \max_{\ell \neq i} \mu_\ell$ ,  $i = 1, 2, \dots, k$ , the difference between each system and the best of the rest. Specialized to the known-variance case, the intervals take the form

$$\mu_i - \max_{\ell \neq i} \mu_\ell \in \left[ -\left(\bar{Y}_i(n) - \max_{\ell \neq i} \bar{Y}_\ell(n) - h\sigma\sqrt{\frac{2}{n}}\right)^-, \left(\bar{Y}_i(n) - \max_{\ell \neq i} \bar{Y}_\ell(n) + h\sigma\sqrt{\frac{2}{n}}\right)^+ \right], \quad (5)$$

where  $h$  is the same critical value used in Bechhofer's and Gupta's procedures,  $-x^- = \min\{0, x\}$  and  $x^+ = \max\{0, x\}$ . Under our assumptions these  $k$  confidence intervals are simultaneously correct with probability  $\geq 1 - \alpha$ .

Consider the set  $I$  containing the indices of all systems whose MCB upper confidence bound is greater than 0. Thus, for  $i \in I$ ,

$$\bar{Y}_i(n) > \max_{\ell \neq i} \bar{Y}_\ell(n) - h\sigma\sqrt{\frac{2}{n}},$$

meaning these are the same systems that would be retained by Gupta's subset-selection procedure. Since  $\mu_k - \max_{\ell \neq k} \mu_\ell > 0$ , and these intervals are simultaneously correct with probability  $\geq 1 - \alpha$ , system  $k$  will be in the subset identified by the MCB upper bounds with the required probability.

Now suppose that  $n$  has been selected such that  $n \geq 2h^2\sigma^2/\delta^2$ , implying that

$$h\sigma\sqrt{\frac{2}{n}} \leq \delta$$

as in Bechhofer's procedure. Let  $B$  be the index of the system with the largest sample mean. Then the MCB lower bounds guarantee with probability  $\geq 1 - \alpha$  that

$$\begin{aligned} \mu_B - \max_{\ell \neq B} \mu_\ell &\geq -\left(\bar{Y}_B(n) - \max_{\ell \neq B} \bar{Y}_\ell(n) - h\sigma\sqrt{\frac{2}{n}}\right)^- \\ &\geq -\delta. \end{aligned}$$

The final inequality follows because  $\bar{Y}_B(n) - \max_{\ell \neq B} \bar{Y}_\ell(n) \geq 0$  by the definition of  $B$ , and  $h\sigma\sqrt{2/n} \leq \delta$  because of our choice of  $n$ . As noted by Nelson and Goldsman (2001), this establishes that the system selected by Bechhofer's procedure is guaranteed to be within  $\delta$  of the true best *under any configuration of the means*. Further, if  $\mu_k - \mu_{k-1} > \delta$ , then  $\Pr\{B = k\} \geq 1 - \alpha$  as required.

As a consequence of this analysis both Bechhofer's and Gupta's procedures can be augmented with MCB confidence intervals "for free," and Bechhofer's procedure is guaranteed to select a system within  $\delta$  of the best. Nelson and Matejcik (1995) establish very mild conditions under which these results hold for far more general SSPs.

The procedure we display below in Section 5.1 is a combined subset, indifference-zone selection, and MCB procedure. The advantage of a combined procedure is that we not only select a system as best, we also gain information about how close each of the inferior systems is to being the best. This information is useful if secondary criteria that are not reflected in the performance measure (such as ease of installation, cost to maintain, etc.) may tempt us to choose an inferior system if it is not deficient by much.

### 5 NORMAL MEANS PROCEDURES

With the motivation from the previous section in mind, we now present three procedures for finding the best (largest mean) normal distribution. The first is a combined screen-select-infer procedure for finding the population with the largest expected value; in this procedure, which uses all three of the approaches outlined in Sections 4.1–4.3, inferior competitors are screened out after an initial stage of sampling. The second is a sequential procedure that can eliminate inferior choices at any stage and uses the common random numbers variance reduction technique in order to make more precise (and therefore efficient) comparisons among the competing populations. The third is an efficient two-stage procedure that also uses common random numbers.

#### 5.1 Subset + Rinott + MCB Procedure

The combined procedure that follows uses a sampling strategy in which the normal observations between scenarios are independent, i.e.,  $Y_{ij}$  is independent of  $Y_{i',j}$  for all  $i \neq i'$  and all  $j$ . Nelson, et al. (2001) show how to combine a simple subset (screening) procedure with a two-stage indifference-zone selection procedure due to Rinott (1978). After the fact, MCB confidence intervals are then provided for free. The procedure simultaneously guarantees a probability of correct selection and confidence-interval coverage probability of at least  $1 - \alpha$  under the stated assumptions. This combined procedure is of great utility when the experimenter is initially faced with a large number of alternatives—the idea is for the subset procedure to pare out non-contending systems, after which Rinott selects the best from the survivors.

##### Procedure Subset + Rinott + MCB

1. Specify the overall desired probability of correct selection  $1 - \alpha$ , the indifference-zone parameter

$\delta$ , a common initial sample size from each scenario  $n_0 \geq 2$ , and the initial number of competing systems  $k$ . Further, set

$$t = t_{1-(1-\alpha/2)^{\frac{1}{k-1}}, n_0-1},$$

where  $t_{\gamma, \nu}$  is the upper- $\gamma$  quantile of a  $t$ -distribution with  $\nu$  degrees of freedom, and let  $h$  solve the following integral,

$$1 - \frac{\alpha}{2} = \int_0^\infty \left[ \int_0^\infty \Phi \left( \frac{h}{\sqrt{\nu \left( \frac{1}{x} + \frac{1}{y} \right)}} \right) f_\nu(x) dx \right]^{k-1} f_\nu(y) dy,$$

where  $\Phi(\cdot)$  is the standard normal cumulative distribution function and  $f_\nu(\cdot)$  is the probability density function of the  $\chi^2$ -distribution with  $\nu = n_0 - 1$  degrees of freedom. The FORTRAN program `rinott` in Bechhofer, Santner, and Goldsman (1995) calculates values of  $h$ , or one can use the tables in Wilcox (1984) or Bechhofer, Santner, and Goldsman (1995).

2. Take an i.i.d. sample  $Y_{i1}, Y_{i2}, \dots, Y_{in_0}$  from each of the  $k$  normal populations, obtained independently.
3. Calculate the first-stage sample means  $\bar{Y}_i^{(1)} = \sum_{j=1}^{n_0} Y_{ij} / n_0$ , and marginal sample variances

$$S_i^2 = \frac{\sum_{j=1}^{n_0} (Y_{ij} - \bar{Y}_i^{(1)})^2}{n_0 - 1},$$

for  $i = 1, 2, \dots, k$ .

4. Calculate the quantity

$$W_{ij} = t \left( \frac{S_i^2 + S_j^2}{n_0} \right)^{1/2}$$

for all  $i \neq j$ . Form the screening subset  $I$ , containing every alternative  $i$  such that  $1 \leq i \leq k$  and

$$\bar{Y}_i^{(1)} \geq \bar{Y}_j^{(1)} - (W_{ij} - \delta)^+ \text{ for all } j \neq i.$$

- 
5. If  $I$  contains a single index, then stop and return that system as the best. Otherwise, for all  $i \in I$ , compute the second-stage sample sizes

$$N_i = \max \left\{ n_0, \left\lceil (hS_i/\delta)^2 \right\rceil \right\}.$$

6. Take  $N_i - n_0$  additional i.i.d. observations from all systems  $i \in I$ , independently of the first-stage sample and the other systems.
7. Compute the overall sample means  $\bar{Y}_i = \sum_{j=1}^{N_i} Y_{ij}/N_i$  for  $i \in I$ .
8. Select the system with the largest  $\bar{Y}_i$  as best.

- 
9. With probability at least  $1 - \alpha$ , we can claim that
    - For all  $i \in I^c$ , we have  $\mu_i < \max_{j \neq i} \mu_j$  (i.e., the systems excluded by the screening are not the best), and
    - If we define  $J_i = \{j : j \in I \text{ and } j \neq i\}$ , then for all  $i \in I$ ,

$$\mu_i - \max_{j \in J_i} \mu_j \in \left[ -\left(\bar{Y}_i - \max_{j \in J_i} \bar{Y}_j - \delta\right)^-, \left(\bar{Y}_i - \max_{j \in J_i} \bar{Y}_j + \delta\right)^+ \right].$$

(Thus, these confidence intervals bound the difference between each alternative and the best of the others in  $I$ .)

## 5.2 Common Random Numbers

A fundamental assumption of the previous Subset+Rinott+MCB procedure is that the  $k$  systems are simulated independently (see Step 2 in that procedure). In practice this means that different streams of pseudo-random numbers are assigned to the simulation of each system. However, under fairly general conditions, assigning common random numbers (CRN) to the simulation of each system decreases the variances of estimates of the pairwise differences in performance. Unfortunately, CRN also complicates the statistical analysis when  $k > 2$  systems are involved. The following procedures from Kim and Nelson (2001) and Nelson and Matejck (1995) provide (almost) the same guarantees as procedure Subset+Rinott+MCB under a more complex set of conditions, but have been shown to be quite robust to departures from those conditions. And unlike Subset+Rinott+MCB, they are designed to exploit the use of CRN to reduce the total number of observations required to make a correct selection.

### 5.2.1 Procedures that Allow Common Random Numbers

We next examine a sequential procedure due to Kim and Nelson (2001) that can eliminate inferior choices at any stage. This procedure uses a sampling strategy in which the normal observations may be dependent due to the use of common random numbers.

The KN procedure is a bit more complicated to implement than the vanilla Rinott (1978) procedure, but it

has several distinct advantages. First, once an initial set of  $n_0$  observations is collected from each treatment, KN is parsimonious in taking additional observations in that they are added one-at-a-time and the data are examined to determine if sufficient information has been collected to stop. In contrast, Rinott and its enhancements take potentially large groups of observations. Second, KN allows treatments to be discarded before the final decision; those treatments that appear inferior can legitimately be dropped from further consideration. See Kim and Nelson (2001, 2005b) and Goldsman, et al. (2002) for more details.

### Procedure KN

1. Specify the overall desired probability of correct selection  $1 - \alpha$ , the indifference-zone parameter  $\delta$ , a common initial sample size from each scenario  $n_0 \geq 2$ , and the initial number of competing systems  $k$ . Calculate the constant

$$\eta = \frac{1}{2} \left[ \left( \frac{2\alpha}{k-1} \right)^{-2/(n_0-1)} - 1 \right].$$

Further, set  $I = \{1, 2, \dots, k\}$  and let  $h^2 = 2\eta(n_0 - 1)$ .

2. Take a random sample of  $n_0$  observations  $Y_{ij}$  ( $1 \leq j \leq n_0$ ) from population  $i$ ,  $i = 1, 2, \dots, k$ . For treatment  $i$  compute the sample mean based on the  $n_0$  observations,  $\bar{Y}_i(n_0) = \sum_{j=1}^{n_0} Y_{ij}/n_0$ ,  $i = 1, 2, \dots, k$ . For all  $i \neq \ell$ , compute the sample variance of the difference between treatments  $i$  and  $\ell$ ,

$$S_{i\ell}^2 = \frac{1}{n_0 - 1} \sum_{j=1}^{n_0} (Y_{ij} - Y_{\ell j} - [\bar{Y}_i(n_0) - \bar{Y}_\ell(n_0)])^2$$

and set

$$N_{i\ell} = \left\lfloor h^2 S_{i\ell}^2 / \delta^2 \right\rfloor,$$

where  $\lfloor \cdot \rfloor$  is the floor (integer round-down) function. Finally, for all  $i$  set

$$N_i = \max_{\ell \neq i} N_{i\ell}.$$

If  $n_0 > \max_i N_i$ , then stop and select the population with the largest sample mean  $\bar{Y}_i(n_0)$  as the one having the largest mean. Otherwise, set the sequential counter  $r = n_0$  and go to the Screening phase of the procedure.

3. *Screening*: Set  $I^{\text{old}} = I$  and re-set

$$I = \{i : i \in I^{\text{old}} \text{ and } \bar{Y}_i(r) \geq \bar{Y}_\ell(r) - W_{i\ell}(r), \\ \text{for all } \ell \in I^{\text{old}}, \ell \neq i\},$$

where

$$W_{i\ell}(r) = \max \left\{ 0, \frac{\delta}{2r} \left( \frac{h^2 S_{i\ell}^2}{\delta^2} - r \right) \right\}.$$

4. *Stopping Rule*: If  $|I| = 1$ , then stop and select the treatment with index in  $I$  as having the largest mean. If  $|I| > 1$ , take one additional observation  $Y_{i,r+1}$  from each treatment  $i \in I$ . Increment  $r = r + 1$  and go to the screening stage if  $r < \max_i N_i + 1$ . If  $r = \max_i N_i + 1$ , then stop and select the treatment associated with the largest  $\bar{Y}_i(r)$  having index  $i \in I$ .

The following two-stage procedure due to Nelson and Matejcik (1995) is similar to Subset+Rinott+MCB, but different in that it purposefully sets out to exploit the use of common random numbers. This is accomplished via a common sample variance estimator that is calculated using observations between alternatives as well as within.

#### Procedure NM + MCB

- Specify the constants  $\delta$ ,  $\alpha$ , and  $n_0 \geq 2$ . Let  $g = T_{k-1, (k-1)(n_0-1), 0.5}^{(\alpha)}$ , an equicoordinate critical point of the equicorrelated multivariate central  $t$ -distribution; this constant can be found in Hochberg and Tamhane (1987), Appendix 3, Table 4; Bechhofer, Santner, and Goldsman (1995); or by using the FORTRAN program AS251 of Dunnett (1989).
- Take an i.i.d. sample  $Y_{i1}, Y_{i2}, \dots, Y_{in_0}$  from each of the  $k$  systems *using CRN across systems*.
- Compute the approximate sample variance of the difference of the sample means

$$S^2 = \frac{2 \sum_{i=1}^k \sum_{j=1}^{n_0} (Y_{ij} - \bar{Y}_i - \bar{Y}_{\cdot j} + \bar{Y}_{\cdot\cdot})^2}{(k-1)(n_0-1)},$$

where  $\bar{Y}_i = \sum_{j=1}^{n_0} Y_{ij}/n_0$ ,  $\bar{Y}_{\cdot j} = \sum_{i=1}^k Y_{ij}/k$ , and  $\bar{Y}_{\cdot\cdot} = \sum_{i=1}^k \sum_{j=1}^{n_0} Y_{ij}/kn_0$ .

4. Compute the final sample size

$$N = \max \left\{ n_0, \left\lceil (gS/\delta)^2 \right\rceil \right\}.$$

- Take  $N - n_0$  additional i.i.d. observations from each system, using CRN across systems.
- Compute the overall sample means  $\bar{\bar{Y}}_i = \sum_{j=1}^N Y_{ij}/N$  for  $i = 1, 2, \dots, k$ .

- Select the system with the largest  $\bar{\bar{Y}}_i$  as best.
- Simultaneously form the MCB confidence intervals

$$\mu_i - \max_{j \neq i} \mu_j \in$$

$$\left[ - \left( \bar{\bar{Y}}_i - \max_{j \neq i} \bar{\bar{Y}}_j - \delta \right)^-, \left( \bar{\bar{Y}}_i - \max_{j \neq i} \bar{\bar{Y}}_j + \delta \right)^+ \right]$$

for  $i = 1, 2, \dots, k$ .

### 5.3 Simulation Considerations

As we have already mentioned, the methods described in this section all rely on our ability to generate i.i.d. normal observations within each scenario. (We may or may not want the observations to be independent between systems, especially if we are thinking of using common random numbers.) Of course, data from a simulation are rarely i.i.d. normal. Luckily, we can achieve approximate normality by taking sample averages of contiguous observations (batching); and we can achieve independence by running independent replications. Moderate departures from normality do not really pose a problem, since all of the procedures appear to be robust in that sense (see Bechhofer, Santner, and Goldsman 1995 or Goldsman, et al. 2002). But one really ought to make sure that the  $Y_{ij}$  are indeed independent within each scenario, for procedure performance seriously deteriorates when that assumption fails.

## 6 MULTINOMIAL PROCEDURES

This section begins with a version of the motivational example from Section 4, but with a slightly different criterion for describing the best alternative. We then describe three procedures to achieve the new goal of finding the best.

**Example 2** *Simulation models of 25 different inventory policies have been developed for potential implementation at a large distribution/warehouse center. The goal now is to select the system that is most likely to have the largest actual profit (instead of the largest expected profit).*

### 6.1 Setup

We define  $p_i$  as the probability that design  $i$  will produce the largest profit from a given vector-observation  $\mathbf{Y}_j = (Y_{1j}, Y_{2j}, \dots, Y_{kj})$ . The goal now is to select the design associated with the largest  $p_i$ -value. This goal is equivalent to that of finding the multinomial category having the largest probability of occurrence; and there is a rich body of literature concerning such problems. In fact, we make almost no assumptions on the underlying distributions

of the competing populations—thus, the procedures to be discussed below are, in a sense, nonparametric.

More specifically, suppose that we want to select the correct category with probability  $1 - \alpha$  whenever the ratio of the largest to second largest  $p_i$  is greater than some user-specified constant, say  $\theta > 1$ . The indifference constant  $\theta$  can be regarded as the smallest ratio “worth detecting.”

### 6.2 The Procedures

This subsection describes three nonparametric procedures. The first is a single-stage procedure for finding the most probable multinomial cell, the second is a sequential procedure, and the third is a clever augmentation of the first that makes more efficient use of the underlying observations.

#### 6.2.1 Single-Stage Procedure

The following *single-stage* procedure was proposed by Bechhofer, Elmaghraby, and Morse (1959) to guarantee the above probability requirement.

##### Procedure BEM

1. For the given  $k$ , and  $(\alpha, \theta)$  specified prior to the start of sampling, find  $n$  from the tables in Bechhofer, Elmaghraby, and Morse (1959), Gibbons, Olkin, and Sobel (1977), or Bechhofer, Santner, and Goldsman (1995).
2. Take a random sample of  $n$  observations  $Y_{i1}, Y_{i2}, \dots, Y_{in}$  from each alternative  $i$ ,  $i = 1, 2, \dots, k$ . Turn these into  $n$  independent multinomial observations,  $X_j = (X_{1j}, X_{2j}, \dots, X_{kj})$ ,  $j = 1, 2, \dots, n$ , by setting

$$X_{ij} = \begin{cases} 1, & \text{if } Y_{ij} > \max_{\ell \neq i} \{Y_{\ell j}\} \\ 0, & \text{otherwise,} \end{cases}$$

where we assume (for notational convenience) that there are never ties for the maximum observation within a particular vector  $Y_j$ .

3. Let  $W_i = \sum_{j=1}^n X_{ij}$  for  $i = 1, 2, \dots, k$ . Select the design that yielded the largest  $W_i$  as the one associated with the largest  $p_i$  (where we simply randomize in the case of ties).

#### 6.2.2 Sequential Procedure

A more efficient procedure, due to Bechhofer and Goldsman (1986), uses *closed, sequential* sampling; that is, the procedure stops when one design is “sufficiently ahead” of the others.

##### Procedure BG

1. For the given  $k$ , and  $(\alpha, \theta)$  specified prior to the start of sampling, find the *truncation number* (i.e., an upper bound on the number of vector-observations)  $n_0$  from the tables in Bechhofer and Goldsman (1986) or Bechhofer, Santner, and Goldsman (1995).
2. At the  $m$ th stage of experimentation ( $m \geq 1$ ), take the random multinomial observation  $X_m = (X_{1m}, X_{2m}, \dots, X_{km})$  (defined above) and calculate the *ordered* category totals

$$W_{[1]m} \leq W_{[2]m} \leq \dots \leq W_{[k]m};$$

also calculate

$$Z_m = \sum_{i=1}^{k-1} (1/\theta)^{(W_{[k]m} - W_{[i]m})}.$$

3. Stop sampling at the first stage when either

$$Z_m \leq \alpha/(1 - \alpha)$$

$$\text{or } m = n_0$$

$$\text{or } W_{[k]m} - W_{[k-1]m} \geq n_0 - m,$$

whichever occurs first.

4. Let  $N$  (a random variable) denote the stage at which the procedure terminates. Select the design that yielded the largest  $W_{iN}$  as the one associated with the largest  $p_i$  (randomize in the case of ties).

#### 6.2.3 Augmentation of Procedure BEM

Miller, Nelson, and Reilly (1998) present a remarkably efficient procedure that directly uses the original  $Y_{ij}$  observations (instead of the 0-1  $X_{ij}$ , which lose information). Their procedure AVC, based on all possible vector comparisons of the observations, always results in an increased probability of correct selection when compared to the analogous implementation of the Bechhofer, Elmaghraby, and Morse (1959) procedure.

##### Procedure AVC

1. For the given  $k$ , and  $(\alpha, \theta)$  specified prior to the start of sampling, use the same  $n$  as in procedure BEM.
2. Take a random sample of  $n$  observations  $Y_{i1}, Y_{i2}, \dots, Y_{in}$  from each alternative  $i$ ,  $i = 1, 2, \dots, k$ . Consider all  $n^k$  vectors of the form  $Y'_j = (Y'_{1j}, Y'_{2j}, \dots, Y'_{kj})$ ,  $j = 1, 2, \dots, n^k$ , where



$Y'_{ij}$  is one of the  $n$  observations from alternative  $i$ . Turn these into  $n^k$  (non-independent) multinomial observations,  $\mathbf{X}'_j = (X'_{1j}, X'_{2j}, \dots, X'_{kj})$ ,  $j = 1, 2, \dots, n^k$ , by setting

$$X'_{ij} = \begin{cases} 1, & \text{if } Y'_{ij} > \max_{\ell \neq i} \{Y'_{\ell j}\} \\ 0, & \text{otherwise,} \end{cases}$$

where we again assume that there are never ties for the maximum observation within a particular vector  $\mathbf{Y}'_j$ .

- Let  $W'_i = \sum_{j=1}^{n^k} X'_{ij}$  for  $i = 1, 2, \dots, k$ . Select the design that yielded the largest  $W'_i$  as the one associated with the largest  $p_i$  (randomize in the case of ties).

## 7 FINAL THOUGHTS

Space limitations preclude detailed discussion, but we also mention a number of additional interesting topic areas that are presently enjoying a great deal of attention. For example,

- New sequential procedures have appeared recently in the “traditional” statistics literature.
- Decision-theoretic perspectives on the selection-of-the best problem.
- Combined selection + optimization procedures in the stochastic environment.
- Optimal sampling budget allocation strategies.
- Combined variance estimation + selection for stationary processes.

The bottom line is that SSPs and MCPs are useful and efficient tools for practical problems; and the potential benefits continue to accrue in this varied and active area of research.

## ACKNOWLEDGMENTS

Portions of this article were published previously in Goldsman and Nelson (2001) and Kim and Nelson (2003). Partial support for our research was provided by National Science Foundation Grant DMI-0400260.

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