

Chapter 9

On a Partially Sequential Ranked Set Sampling Paradigm

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Abstract In a two-sample setting it is important to design statistical procedures that can take advantage of additional information to minimize the sample sizes required to reach reliable inferences about possible differences between the two populations. This is particularly true when it is difficult and/or costly to obtain sample observations from one or both of the populations. One class of procedures designed with this goal in mind uses the partially sequential sampling (*PS*) approach, first introduced by Wolfe (Journal of the American Statistical Association 72:202–205, 1977a). The use of ranked set sampling (*RSS*), first introduced by McIntyre (Australian Journal of Agricultural Research 3:385–390, 1952, reprinted in 2005), offers another approach for minimizing required sample sizes through the mechanism of obtaining more representative samples than can be achieved using simple random samples. In this paper we provide a review of these two sampling techniques and discuss options for melding the two methodologies to obtain partially sequential ranked set sample (*PSRSS*) two-sample test procedures that take advantage of the sample saving properties of both the *PS* and *RSS* approaches. To illustrate this combination, we consider *PSRSS* procedures where the fixed (control) sample is obtained via simple random sampling and the sequential (treatment) sample is obtained via ranked set sampling. Properties of the associated tests are discussed, including the limiting distributions as the fixed sample size tends to infinity.

Keywords Distribution free tests • Judgment ranked order statistics • Minimizing sample sizes • Negative binomial distribution • Using auxiliary sampling information

9.1 Introduction

Minimizing the cost associated with collection of the sample data is a critical feature of most statistical analyses. As a result, it is important to develop statistical approaches to sampling that minimize the sample sizes necessary to achieve desired

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properties, whether it be precision of estimators, length of confidence intervals, or power of statistical tests. One technique that has been shown to be useful in this regard is ranked set sampling (*RSS*), first introduced by McIntyre (1952, reprinted in 2005) in the context of sampling from pasture and crop plots. This sampling approach uses readily available auxiliary information from individual units in a population to aid in the selection of more representative units for measurement than are typically generated by simple random sampling (*SRS*). Development of statistical procedures using this *RSS* approach remains an active area of research. [See, for example, the recent survey article by Wolfe (2012).] A second approach to data collection designed to reduce the sample size in a treatment versus control two-sample setting is the partially sequential (*PS*) paradigm introduced by Wolfe (1977a,b). This approach uses a negative binomial sampling framework to minimize the number of treatment observations necessary for reaching satisfactory statistical conclusions regarding the treatment's efficacy.

In this paper we review the basic tenets of both the *RSS* and *PS* methodologies and discuss how to combine these approaches to develop partially sequential ranked set sample (*PSRSS*) two-sample test procedures. In Sect. 9.2 we present the *PS* two-sample framework and review previous work in this area. We describe the basic *RSS* approach in Sect. 9.3 and discuss a number of options available within this structure. We propose a class of melded *PSRSS* two-sample test procedures in Sect. 9.4 and develop their basic small sample and asymptotic properties as the control sample size becomes large. Section 9.5 is devoted to a general discussion of the opportunities presented by this new methodology as well as extensions for future research.

9.2 Partially Sequential Two-Sample Procedures

The partially sequential approach to data collection in the two-sample setting was first introduced by Wolfe (1977a). It is particularly appropriate for data collection settings such as the following:

1. A sample from the first population (e.g., control) has already been collected and we do not wish to collect any more observations from the second population (e.g., new treatment) than are necessary for reaching a decision.
2. Neither sample has been collected, but one of the samples (say the 'standard' procedure observations) is relatively easy and inexpensive to collect, while the other sample observations (corresponding to the 'new treatment') are costly and/or difficult to collect. In such situations our goal would be to collect a sample (usually large) of standard observations and then collect only enough difficult-to-obtain new treatment observations necessary to reach statistically valid conclusions about potential differences between the two populations.

We first describe a general *PS* procedure to test for differences between two distributions. Let X_1, \dots, X_m be a random sample from a continuous probability

distribution with p.d.f. $f(x)$ and c.d.f. $F(x)$, where m is a fixed positive integer, and let $G(y)$ be a second continuous distribution function with associated p.d.f. $g(y)$. Let (x_1, \dots, x_m) be an arbitrary m -tuple of real numbers and let $A(x_1, \dots, x_m)$ be a subset of the real line R depending on the m -tuple (x_1, \dots, x_m) . For example, $A(\cdot)$ could be the portion of R below the minimum x -value or the portion of R above the maximum x -value. Define the indicator function $\Psi(\cdot)$ by

$$\Psi(y) = \begin{cases} 1, & \text{if } y \in A(x_1, \dots, x_m), \\ 0, & \text{if } y \notin A(x_1, \dots, x_m). \end{cases} \tag{9.1}$$

Now let Y be a random variable (independent of X_1, \dots, X_m) from the second distribution $G(y)$. Applying $\Psi(y)$ to these random variables X_1, \dots, X_m and Y , we obtain the following

$$\Psi(Y) = \begin{cases} 1, & \text{if } Y \in A(X_1, \dots, X_m), \\ 0, & \text{if } Y \notin A(X_1, \dots, X_m). \end{cases} \tag{9.2}$$

Thus, $\Psi(Y)$ is the random indicator variable for the random set $A(X_1, \dots, X_m)$.

With (9.2) in mind we sequentially sample mutually independent Y 's from the distribution $G(y)$ until a preset number, say r , of these Y 's are in the set $A(x_1, \dots, x_m)$, where (x_1, \dots, x_m) is the observed value of the previously collected random vector (X_1, \dots, X_m) . Define the statistic N_m (having random contributions from both the X and Y samples) by

$$N_m = \{\text{number of } Y \text{ observations required to get } r \text{ } Y\text{'s in } A(x_1, \dots, x_m)\}. \tag{9.3}$$

Wolfe (1977a) discussed how to use N_m to test the null hypothesis $H_0 : F(x) \equiv G(x)$ against appropriate alternatives [depending on the nature of the set $A(x_1, \dots, x_m)$]. The decision rule he proposed is to reject H_0 when $N_m \leq N_0(\alpha, r, m, A)$, where $N_0(\alpha, r, m, A)$ is the lower α th percentile point for the null (H_0) distribution of N_m . Note that with this approach we will never need to collect more than $N_0(\alpha, r, m, A)$ Y observations. In fact, we would stop even sooner with an even smaller Y sample size and (1) reject H_0 as soon as we obtain r Y observations in $A(x_1, \dots, x_m)$ or (2) fail to reject H_0 as soon as we obtain $\{N_0(\alpha, r, m, A) - r + 1\}$ Y observations not in $A(x_1, \dots, x_m)$.

9.2.1 Properties of Partially Sequential Procedures

For given $X_1 = x_1, \dots, X_m = x_m$, let

$$p_m = p_m(x_1, \dots, x_m) = P_G\{Y \in A(x_1, \dots, x_m)\}. \tag{9.4}$$

Thus, p_m is the conditional probability that an observation from the distribution G falls in the set $A(x_1, \dots, x_m)$ prescribed by the observed values from the F distribution. Then, conditional on $X_1 = x_1, \dots, X_m = x_m$, N_m has a negative binomial distribution with parameters r and p_m ; that is,

$$\begin{aligned}
 P(N_m = n | X_1 = x_1, \dots, X_m = x_m) \\
 = \binom{n-1}{r-1} [p_m(x_1, \dots, x_m)]^r [1 - p_m(x_1, \dots, x_m)]^{n-r} I_{\{r, r+1, r+2, \dots\}}(n).
 \end{aligned}
 \tag{9.5}$$

The unconditional distribution of N_m is obtained from the result in (9.5) by integrating over the distribution of the X 's, namely,

$$P(N_m = n) = E_F \left\{ \binom{n-1}{r-1} [p_m(X_1, \dots, X_m)]^r [1 - p_m(X_1, \dots, X_m)]^{n-r} \right\} I_{\{r, r+1, r+2, \dots\}}(n).
 \tag{9.6}$$

Since the investigator has flexibility in setting the sample size m for the X observations, it is of interest to know how N_m behaves as m becomes large, that is, as $m \rightarrow \infty$. If, for given F and G , $p_m(X_1, \dots, X_m)$ converges in probability to a fixed number $p = p(F, G)$, $0 < p \leq 1$, as $m \rightarrow \infty$, then the limiting distribution ($m \rightarrow \infty$) of N_m is negative binomial with parameters r and p ; that is, the asymptotic distribution ($m \rightarrow \infty$) of N_m is

$$P^*(N_m = n) = \binom{n-1}{r-1} p^r (1-p)^{n-r} I_{\{r, r+1, r+2, \dots\}}(n).
 \tag{9.7}$$

(Note: A limiting value of $p = p(F, G) = 0$ does not satisfy the conditions for this result. If a pair (F, G) produces a limiting value of $p = 0$, the statistic N_m does not possess a limiting distribution as $m \rightarrow \infty$, since in such cases N_m increases stochastically without limit as $m \rightarrow \infty$.)

When m is fixed and large we can use the limiting distribution in (9.7) to select r to guarantee asymptotic ($m \rightarrow \infty$) power against an alternative to H_0 of interest. Let H_a be an alternative to H_0 against which we require an approximate power β , where $0 < \beta < 1$ is arbitrary. Let p^* be the value of p in (9.7) that corresponds to the alternative H_a . Then from the definition of $N_0(\alpha, r, \infty, A)$ (i.e., the approximate α -level critical value for the asymptotic, $m \rightarrow \infty$, distribution of N_m), this approximate power requirement corresponds to

$$\sum_{n=r}^{N_0(\alpha, r, \infty, A)} \binom{n-1}{r-1} (p^*)^r (1-p^*)^{n-r} \geq \beta.
 \tag{9.8}$$

For many partially sequential procedures, the left side of the inequality in (9.8) is a non-decreasing function of r . In this case, to satisfy our asymptotic power requirements with the fewest Y observations, we can preset r to be $r = r^*$, where r^* is the smallest integer for which (9.8) is satisfied.

9.2.2 Examples

The *PS* approach can be used equally well in parametric or nonparametric settings. We briefly discuss two such examples.

Example 1: Parametric Setting

Let $F(x) = \Phi\{\frac{x-\mu_1}{\sigma}\}$ and $G(y) = \Phi\{\frac{y-\mu_2}{\sigma}\}$, where $\Phi(t)$ is the standard normal distribution function. The null hypothesis of interest is $H_0 : \mu_1 = \mu_2$ and we consider here the alternative $H_a : \mu_2 > \mu_1$.

One method for selecting the set $A(x_1, \dots, x_m)$ would be to view the indicator $\Psi(\cdot)$ in (9.1) as a critical function for testing H_0 against H_a for random samples of sizes m and 1 from the F and G distributions, respectively. For example, we know that the uniformly most powerful level α^* test of H_0 against H_a for m X observations and a single Y observation has critical region

$$C(y, x_1, \dots, x_m) = \left\{ (y, x_1, \dots, x_m) : \frac{m^{1/2}}{(m+1)^{1/2}} \frac{(y-\bar{x})}{s} \geq t_{\alpha^*}(m-1) \right\},$$

where $\bar{x} = \sum_{i=1}^m x_i/m$, $s^2 = \sum_{i=1}^m (x_i - \bar{x})^2/(m-1)$ and $t_{\alpha^*}(m-1)$ is the upper α^* percentile point for the t distribution with $m-1$ degrees of freedom.

Thus, in this setting it is natural to take the set $A(x_1, \dots, x_m)$ to be

$$A(x_1, \dots, x_m) = \{y : y \geq \bar{x} + t_{\alpha^*}(m-1)(\{m+1\}s^2/m)^{\frac{1}{2}}\}. \tag{9.9}$$

In fact, Orban and Wolfe (1978) showed that this choice of $A(x_1, \dots, x_m)$ leads to the asymptotically ($m \rightarrow \infty$) most powerful level α^* partially sequential procedure for testing H_0 against H_a .

With $A(x_1, \dots, x_m)$ given by (9.9), we have

$$p_m = 1 - \Phi(\{\bar{x} + t_{\alpha^*}(m-1)(\{m+1\}s^2/m)^{\frac{1}{2}} - \mu_2\}/\sigma)$$

and the limiting distribution of N_m as $m \rightarrow \infty$ is negative binomial (9.7) with parameters r and $p = \lim_{m \rightarrow \infty} p_m = 1 - \Phi\{\frac{\mu_1 - \mu_2}{\sigma} + z_{\alpha^*}\}$.

Example 2: Nonparametric Setting

Let F and G be arbitrary, continuous distribution functions. We wish to test $H_0 : F \equiv G$ against the alternative $H_a : \xi_2 > \xi_1$, where ξ_1 and ξ_2 are the medians of the F and G distributions, respectively. Assume that m is an odd integer (more complicated, but tractable for m even) and define $A(x_1, \dots, x_m)$ by

$$A(x_1, \dots, x_m) = \{y : y > m_x\}, \tag{9.10}$$

where $m_x = \text{median}(x_1, \dots, x_m)$. Then the *PS* two-sample median test associated with N_m (9.3) has the following properties:

(a) $p_m = 1 - G(m_x)$ and the exact null (H_0) distribution of N_m is given by

$$P_0(N_m = n) = \begin{cases} \binom{n-1}{r-1} \frac{m!}{[\{(m-1)/2\}!]^2} \frac{\Gamma\left(\frac{m+2n-2r+1}{2}\right)\Gamma\left(\frac{2r+m+1}{2}\right)}{(m+n)!}, & n = r, r+1, \dots \\ 0, & \text{elsewhere.} \end{cases} \quad (9.11)$$

(b) The limiting distribution of N_m as $m \rightarrow \infty$ is negative binomial (9.7) with parameters r and $p = \lim_{m \rightarrow \infty} p_m = 1 - G(\xi_1)$, with $p = 1/2$ or $> 1/2$ depending on whether H_0 or H_a is true, respectively.

Wolfe (1977b) initially proposed this *PS* two-sample median procedure and Orban and Wolfe (1982) studied its properties, including the expected number of Y observations required to conduct the test. They also provided the necessary tables for selecting r so that the approximate power requirement in (9.8) can be attained.

9.3 Ranked Set Sampling

The goal of *RSS* is to collect observations that are more likely to be representative of the full range of values in a population than the same number of observations obtained via *SRS*. To obtain a balanced *RSS* of k observations from a population, we proceed as follows. First, an initial *SRS* of k units is selected from the population and rank ordered on the attribute of interest. This ranking can be obtained through a variety of mechanisms, including visual comparisons, expert opinion, or through the use of correlated concomitant variables, but it cannot involve actual measurements of the attribute of interest on the selected units. The unit that is judged to be the smallest in this ranking is taken as the first item in the *RSS* and the attribute of interest is formally measured for the unit and denoted by $X_{[1]}$. Note that square brackets are used instead of the usual round brackets for the smallest order statistic since $X_{[1]}$ may or may not actually have the smallest attribute measurement among the k units in the *SRS*, even though our ranking judged it to be the smallest. The other remaining $k - 1$ units in our initial *SRS* are not considered further in making inferences about the population—they were used solely to assist in the selection of the smallest ordered ranked unit for measurement.

Following the selection of $X_{[1]}$, a second *SRS* (independent of the first *SRS*) of size k is selected from the population and ranked in the same manner as the first *SRS*. From this second *SRS* we select the item ranked as the second smallest of the k units (i.e., the second judgment order statistic) and add its attribute measurement, $X_{[2]}$, to the *RSS*. From a third *SRS* (independent of both previous *SRS*'s) of size k we

select the unit ranked to be the third smallest (i.e., the third judgment order statistic) and include its attribute measurement, $X_{[3]}$, in the *RSS*. This process continues until we have selected the unit ranked to be the largest of the k units in the k th independent *SRS* and included its attribute measurement, $X_{[k]}$, in our *RSS*.

This process results in the k measured observations $X_{[1]}, X_{[2]}, \dots, X_{[k]}$ and is called a *cycle*. The number of units, k , in each *SRS* is called the set size. To complete a single ranked set cycle, we need to access a total of k^2 units from the population to separately rank k independent simple random samples of size k each. The measured observations, $X_{[1]}, X_{[2]}, \dots, X_{[k]}$, constitute a *balanced ranked set sample of size k* , where the descriptor “balanced” refers to the fact that we have collected one judgment order statistic for each of the ranks $1, 2, \dots, k$. To obtain a final balanced *RSS* with a desired total number of measured observations $n = qk$, we repeat the entire process for q independent cycles, yielding the balanced *RSS* of size n : $X_{[1]_j}, X_{[2]_j}, \dots, X_{[k]_j}$, for $j = 1, \dots, q$.

Note that a balanced *RSS* of size n differs from an *SRS* of size n in a number of important ways. An *SRS* is designed so that the n observations in the sample are mutually independent and identically distributed. This means that, probabilistically speaking, each of the individual sample items can be viewed as representative of a typical value from the underlying population. That is certainly not the case for a balanced *RSS* of size n . While the individual observations in a balanced *RSS* are also mutually independent, they are clearly not identically distributed. As such, it is not the case that each of the individual observations in a balanced *RSS* represents a typical value from the underlying population. On the contrary, the individual judgment order statistics represent very distinctly different portions of the underlying population. It is, however, precisely this additional structure on the items in the balanced *RSS* that enables it to provide greater assurance that the entire range of population values are represented in the sample data.

There have been numerous papers in the literature demonstrating the advantages that balanced *RSS* provides relative to *SRS*, both in terms of precision accuracy and in terms of reducing required sample sizes. Dell and Clutter (1972) showed that the estimator of the population mean μ based on a balanced *RSS* is unbiased and it has a variance that is never larger than the variance of the estimator of μ based on a *SRS* of the same size. The remarkable thing is that this result is true even if the judgment ranking for the balanced *RSS* is not perfect. The better the judgment ranking, of course, the greater the improvement from using a balanced *RSS* instead of a *SRS*. Stokes and Sager (1988) obtained similar results for the *RSS* estimator of the distribution function of the population and Terpstra (2004) did the same for the *RSS* maximum likelihood estimator for a population proportion.

While a balanced *RSS* is the most commonly occurring form of ranked set sampling data, there are situations where it is not optimal to collect the same number of measured observations for each of the judgment order statistics. For example, suppose we are interested primarily in making inferences about the median ξ of a distribution based on an odd number of observations $k = 2d + 1$. It is well known that among all the order statistics the sample median, $X_{(d+1)}$, contains the most information about ξ when the underlying distribution is unimodal and symmetric.

Thus, to make inferences about ξ , it is natural to measure the same judgment order statistic, $X_{[d+1]}$, in each set so that it is measured all k times in each of the q cycles. The resulting *RSS* consists of qk measured observations, each of which is a judgment median from a set of size k . This is the most efficient *RSS* for making inferences about the population median ξ for a distribution that is both unimodal and symmetric, and it is clearly as unbalanced as possible. (A similar argument calls for a distinctly different unbalanced *RSS* for estimating the median of an asymmetric unimodal population or a multimodal population. See, for example, Ozturk and Wolfe 2000, and Chen et al. 2006.) We should point out, however, that such a median unbalanced *RSS* would not necessarily be a good idea if we wanted to make inference about other features of the population, such as its distribution function or the population variance.

RSS and related methodology has an active and rich literature. The interested reader is referred to the recent survey and review articles in Wolfe (2004) and Wolfe (2012) for more comprehensive discussions.

9.4 A Class of *PSRSS* Two-Sample Percentile Test Procedures

There are three approaches that can be taken to incorporate *RSS* into partially sequential procedures:

1. Use *RSS* for the X sample data and *SRS* for the sequentially obtained Y sample data.
2. Use *RSS* for both the X sample and Y sample data.
3. Use *SRS* for the X sample data and *RSS* for the Y sample data.

All three of these options are worthy of consideration, although the first approach is probably the least interesting in the context where partially sequential procedures would be most useful. In this paper we concentrate on the most natural third option to provide an illustration of how to introduce *RSS* into the partially sequential process. To facilitate the discussion we consider the particular unbalanced *RSS* corresponding to all of the observations being collected at a single judgment order statistic and we assume that the judgment ranking is perfect, so that the various judgment order statistics can be viewed as true order statistics.

As before, let X_1, \dots, X_m be a random sample from a probability distribution with p.d.f. $f(x)$ and c.d.f. $F(x)$, where m is an odd integer, and let $G(y)$ be a second distribution function with associated p.d.f. $g(y)$. Let M_X be the X sample median and let m_x be the observed value of M_X . Once again we wish to test $H_0 : F \equiv G$ against the alternative $H_a : \xi_2 > \xi_1$, where ξ_1 and ξ_2 are the medians of the F and G distributions, respectively.

For illustrative purposes, we consider collecting unbalanced *RSS* data from G using a single cycle ($q = 1$) with set size k and measuring the j th order statistic, $Y_{(j)}$,

at each step of the sequential sampling, for fixed $j \in \{1, \dots, k\}$. With this RSS Y -sampling scheme and the indicator set $A(x_1, \dots, x_m) = \{y : y > m_x\}$, the associated PSRSS test of $H_0 : F \equiv G$ against the alternative $H_a : \xi_2 > \xi_1$ has the following properties:

(a) The unconditional exact distribution of N_m still has the form

$$P(N_m = n) = E_F \left\{ \binom{n-1}{r-1} [p_m(X_1, \dots, X_m)]^r [1 - p_m(X_1, \dots, X_m)]^{n-r} \right\} I_{\{r, r+1, r+2, \dots\}}(n), \quad (9.12)$$

but the parameter $p_m = p_m(x_1, \dots, x_m)$ is now given by

$$p_m = P\{Y_{(j)} > m_x\} = 1 - Q_j(m_x),$$

where $Q_j(\cdot)$ is the c.d.f. for the j^{th} order statistic for a random sample of size k from G , given by

$$Q_j(t) = \sum_{u=j}^k \binom{k}{u} [G(t)]^u [1 - G(t)]^{k-u}. \quad (9.13)$$

Combining (9.12) and (9.13), the unconditional distribution of N_m becomes

$$P(N_m = n) = E_{F_{M_X}} \left\{ \binom{n-1}{r-1} [1 - Q_j(M_X)]^r [Q_j(M_X)]^{n-r} \right\} I_{\{r, r+1, r+2, \dots\}}(n), \quad (9.14)$$

where F_{M_X} is the c.d.f. of the sample median for a random sample of size m from F . Using the standard form of F_{M_X} for an odd sample size m in expression (9.14), it follows that

$$\begin{aligned} P(N_m = n) &= \int_{-\infty}^{\infty} \binom{n-1}{r-1} [1 - Q_j(t)]^r [Q_j(t)]^{n-r} \\ &\quad \times \frac{m!}{[(\frac{m-1}{2})!]^2} \{F(t)[1 - F(t)]\}^{\frac{m-1}{2}} f(t) dt I_{\{r, r+1, r+2, \dots\}}(n), \end{aligned} \quad (9.15)$$

Under $H_0 : F \equiv G$ it follows from the change of variable $v = F(t)$ in (9.15) that the null distribution for N_m “simplifies” to

$$\begin{aligned} P(N_m = n) &= \int_0^1 \binom{n-1}{r-1} \left\{ 1 - \sum_{u=j}^k \binom{k}{u} [v]^u [1 - v]^{k-u} \right\}^r \left[\sum_{u=j}^k \binom{k}{u} [v]^u [1 - v]^{k-u} \right]^{n-r} \\ &\quad \times \frac{m!}{[(\frac{m-1}{2})!]^2} \{v(1 - v)\}^{\frac{m-1}{2}} dv I_{\{r, r+1, r+2, \dots\}}(n), \end{aligned} \quad (9.16)$$

This expression clearly does not depend on the form of the continuous F , so that the test based on N_m is distribution-free and the exact critical values for the test can be evaluated from (9.16) without knowledge of F .

- (b) The limiting distribution of N_m as $m \rightarrow \infty$ is negative binomial with parameters r and $p_j^* = \lim_{m \rightarrow \infty} p_m = 1 - Q_j(\xi_1)$.

Using the expression for $Q_j(t)$ in (9.13), we see that

$$p_j^* = 1 - Q_j(\xi_1) = 1 - \sum_{u=j}^k \binom{k}{u} [G(\xi_1)]^u [1 - G(\xi_1)]^{k-u},$$

which simplifies under the null hypothesis to

$$p_{0j}^* = 1 - Q_j(\xi_1) = 1 - \sum_{u=j}^k \binom{k}{u} [F(\xi_1)]^u [1 - F(\xi_1)]^{k-u} = 1 - \sum_{u=j}^k \binom{k}{u} [0.5]^u [0.5]^{k-u}, \quad (9.17)$$

9.4.1 Special Cases

1. $j = k$ —here we are measuring the maximum judgment order statistic in each set and

$$p_{0k}^* = 1 - \sum_{u=k}^k \binom{k}{u} [0.5]^u [0.5]^{k-u} = 1 - \binom{k}{k} [0.5]^k [0.5]^{k-k} = 1 - (0.5)^k, \quad (9.18)$$

which converges to 1 as $k \rightarrow \infty$.

2. $j = 1$ —here we are measuring the minimum judgment order statistic in each set and

$$\begin{aligned} p_{01}^* &= 1 - \sum_{u=1}^k \binom{k}{u} [0.5]^u [0.5]^{k-u} \\ &= 1 - \left[\sum_{u=0}^k \binom{k}{u} [0.5]^u [0.5]^{k-u} - \binom{k}{0} [0.5]^0 [0.5]^{k-0} \right] \\ &= 1 - [1 - (0.5)^k] = (0.5)^k, \end{aligned} \quad (9.19)$$

which converges to 0 as $k \rightarrow \infty$. (Remember that this is not a viable option for PSRSS.)

3. $j = d + 1$, where $k = 2d + 1$ is an odd integer—here we are measuring the median judgment order statistic, $Y_{(d+1)}$, in each set and

$$\begin{aligned}
 p_{0(d+1)} &= 1 - \sum_{u=d+1}^{2d+1} \binom{2d+1}{u} [0.5]^u [0.5]^{(2d+1)-u} \\
 &= \sum_{u=0}^d \binom{2d+1}{u} [0.5]^u [0.5]^{(2d+1)-u} \\
 &= \frac{1}{2} \sum_{u=0}^{2d+1} \binom{2d+1}{u} [0.5]^u [0.5]^{(2d+1)-u} = \frac{1}{2}(1) = 0.5. \quad (9.20)
 \end{aligned}$$

The fact that the limiting distribution under the general alternative $F \neq G$ depends on both the negative binomial stopping parameter r and the set size k provides us with even greater flexibility in designing a study with the idea of guaranteeing prescribed power against specific alternatives. Both increasing r and increasing k will lead to increased power for the *PSRSS* median procedure, but increasing r will also lead to a larger number of measured observations from the Y distribution, something that we are trying to avoid. Increasing k and/or increasing the initial sample size m from the X distribution can be used as effective alternatives for increasing the power without increasing the number of measured Y observations.

9.5 Discussion and Future Research

Small sample and asymptotic properties of the *PSRSS* two-sample median test procedure (corresponding to special case 3) based on measuring the Y sample median (for an odd set size k) in every ranked set have been investigated extensively by Matthews et al. (2016). They found that taking the *RSS* approach for collection of the Y sample observations leads to both increased power and decreased expected Y sample size relative to the *PSRSS* version studied by Orban and Wolfe (1982). This is due to both the intrinsic structure inherent in the partially sequential approach to the two-sample problem and the ranked set sampling methodology employed in obtaining the Y sample. As noted in Sect. 9.4, further improvements in both power and reduced Y sample size can likely be obtained by utilizing *RSS* to collect both the X and Y sample observations. The basic formulation of this dual *RSS* approach would be analogous to what we utilized in this paper using *SRS* to collect the X sample items, although the mathematical properties would be more complicated. Another intriguing possibility would be to develop *PSRSS* methodology that utilized a fully balanced *RSS* approach to the collection of the Y observations, rather than relying solely on the use of the medians of the ranked sets. This could also include a fully balanced *RSS* approach to collection of the initial X sample, leading to natural partially sequential analogues to the two-sample balanced *RSS* procedures considered by Bohn and Wolfe (1992) and Fligner and MacEachern (2006).

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