

Best (but oft-forgotten) practices: sample size and power calculation for a dietary intervention trial with episodically consumed foods

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ABSTRACT

Dietary interventions often target foods that are underconsumed relative to dietary guidelines, such as vegetables, fruits, and whole grains. Because these foods are only consumed episodically for some participants, data from such a study often contains a disproportionately large number of zeros due to study participants who do not consume any of the target foods on the days that dietary intake is assessed, thus generating semicontinuous data. These zeros need to be properly accounted for when calculating sample sizes to ensure that the study is adequately powered to detect a meaningful intervention effect size. Nonetheless, this issue has not been well addressed in the literature. Instead, methods that are common for continuous outcomes are typically used to compute the sample sizes, resulting in a substantially under- or overpowered study. We propose proper approaches to calculating the sample size needed for dietary intervention studies that target episodically consumed foods. Sample size formulae are derived for detecting the mean difference in the amount of intake of an episodically consumed food between an intervention and a control group. Numerical studies are conducted to investigate the accuracy of the sample size formulae as compared with the ad hoc methods. The simulation results show that the proposed formulae are appropriate for estimating the sample sizes needed to achieve the desired power for the study. The proposed method for sample size is recommended for designing dietary intervention studies targeting episodically consumed foods. *Am J Clin Nutr* 2020;112:920–925.

Keywords: dietary intervention trials, episodic consumption, power, sample size, type I error

Introduction

Intake of whole plant foods is associated with reduced risk of numerous adverse health outcomes including obesity, cardiovascular disease, diabetes, and certain cancers (1–18). These foods are consistently underconsumed in the US diet, and as such, are often targeted in dietary intervention studies (19–25).

Because these foods may only be episodically consumed by some participants, data from 24-h recalls may contain a sizable number of participants who do not consume any of a target food group during the recall period (26, 27). The resulting distribution, which contains a disproportionately large number of zeros yields a so-called semicontinuous data structure (28). For example, data for children aged 2–8 y from the NHANES, 2001–2004 showed nonconsumption of total fruit, whole fruit, whole grains, total vegetables, dark green or orange vegetables or legumes, and milk on any single day among 17%, 40%, 42%, 3%, 50%, and 12% of participants, respectively (26).

The excessive zeros influence both the design and analysis of the trial and they need to be dealt with appropriately; otherwise the trial may be poorly powered, and the study findings may be biased and misleading. In addition to the excessive zeros, a further complication which also needs to be considered is that the intervention can change the percent of zeros in the intervention arm compared with the control arm.

There is a considerable body of work in the statistical literature on modeling and analysis of semicontinuous data (29–34). However, limited attention has been paid to the design of an intervention study that involves such data, and appropriate statistical methods for estimating sample sizes and power are lacking for comparison concerning the overall mean of the semicontinuous data. An extensive review of relevant literature

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has found a sizable number of interventional trials targeting vegetables, fruits, and whole grains (35–40). To the best of our knowledge, however, few published studies have provided details on sample size determination and power analysis, let alone consideration of accounting for excessive zeros in the outcome. Among those that discussed sample size calculation and power analysis, ad hoc methods suitable for continuous outcomes are often cited for sample size calculation.

In this commentary, we confine our attention to a 2-arm trial investigating the efficacy of an intervention to increase the consumption of an episodically consumed food, such as whole fruits or vegetables. The dietary intake data are collected using an assessment tool such as The Automated Self-Administered 24-h dietary recall developed by the National Cancer Institute (41). We assume that the primary objective of the trial is to compare the average amount of intake between the intervention and the control. Because a dietary intervention targeting an episodically consumed food can increase (or decrease) not only the average amount of intake of the food if consumed, but also the rate of consumption of the food, the sample size calculation should take both aspects into account.

In what follows, we present the appropriate approaches to computing the sample sizes and evaluate their performance using Monte Carlo simulation as compared with the ad hoc methods for comparison of 2 means. The sample size formulae are also provided (**Supplemental Technical Details**) for 1-arm trials against a historical control and prepost trials comparing intervention results with that prior to intervention.

The Common (but Incorrect) Practice

It is common in 2-arm trials with continuous outcomes to use the 2 sample size formulae for comparing the means of 2 independent populations. One assumes equal SD and the other assumes unequal SDs for the control and intervention arms (42).

Consider a 2-arm trial with equal allocation. For a study participant in the k th arm ($k = 1$ if control and 2 if intervention), let X_k denote the amount of food consumed, taking values of either 0 if the food was not consumed or a positive numerical value if the food was consumed. The mean of X_k is denoted by θ_k and the SD is denoted by ν_k . The null hypothesis to be tested is $H_0 : \theta_2 - \theta_1 = 0$, that is, the mean amount of intake in the intervention arm is the same as in the control arm. We would like to test the null hypothesis at a significance level α , and to determine the sample size per arm n so that the trial is powered at $(1 - \beta)$ level to detect a prespecified meaningful difference of $H_1 : \theta_2 - \theta_1 = \delta (> 0)$, that is, the intervention increases the mean amount of intake by δ as compared with the control arm (alternative hypothesis). To calculate the sample size using the formula for 2 means, 1 specifies the values for the 2 SDs, say ν_1 for control and ν_2 for intervention, and then compute the sample size per arm from the following formula:

$$n = \frac{(Z_{1-\alpha/2} + Z_{1-\beta})^2}{\delta^2} (\nu_1^2 + \nu_2^2), \quad (1)$$

where Z_γ is the 100 γ % quantile of the standard normal distribution.

If we assume the 2 SDs are equal, that is, $\nu_2 = \nu_1$, then the sample size formula (1) becomes:

$$n = \frac{2(Z_{1-\alpha/2} + Z_{1-\beta})^2}{\delta^2} \nu_1^2, \quad (2)$$

which is also commonly used in practice when designing a clinical trial, particularly when empirical data are scarce, making it difficult to specify the SD for the intervention arm of the study.

With the prespecified values of the 2 SDs, the targeted mean difference (effect size), level of significance, and power to detect the effect size, the sample sizes needed for the trial are then conveniently obtained from statistical software such as R, SAS, and PASS (43–45). In practice, these 2 formulae are frequently used when the endpoints, such as the amount of intake of whole fruits, are semicontinuous. As we demonstrate below, specifying the SDs for a semicontinuous outcome is more complicated than for a continuous outcome, especially under the alternative hypothesis needed for power analysis. We numerically show that using these ad hoc formulae yield inadequate estimation of the sample sizes, resulting in a study either seriously underpowered (thus compromising the ability of the trial to detect meaningful intervention effects) or unnecessarily overpowered (thus increasing the financial and administrative burden to conduct the trial).

The Appropriate Approach: Accounting for Nonconsumption

The amount of intake of an episodically consumed food from an interventional trial is semicontinuous with observations characterized by many zeros due to nonconsumption of the food. In general, the trial data in each arm can be divided into 2 parts (26): nonconsumption during assessment (consisting of all the zeros) and any consumption during assessment (consisting of the reported amount of intake). To properly compute the sample sizes, we need to take the distributions of both parts into consideration.

Suppose that, for the k th arm, the probability that the food is consumed is p_k , and the mean and SD of the amount of intake given the food is consumed are respectively, μ_k and σ_k . The probability p_k fully captures the distribution of the nonconsumption part of the data and μ_k and σ_k are 2 important parameters characterizing the distribution of the consumption part of the data.

To link the above parameters with the mean θ_k and SD ν_k of the amount of intake X_k , we have $\theta_k = p_k \mu_k$, and:

$$\nu_k^2 = p_k \sigma_k^2 + p_k (1 - p_k) \mu_k^2; \quad (3)$$

see Supplementary Equation (S1) in the Supplemental Technical Details for more detail. Hence the SD of X_k depends on the mean μ_k and SD σ_k of the amount of intake when the food is consumed, as well as the probability p_k that the food is consumed. The null hypothesis to be tested becomes $H_0 : \theta_2 - \theta_1 = p_2 \mu_2 - p_1 \mu_1 = 0$ and the alternative hypothesis to be powered is $H_1 : \theta_2 - \theta_1 = p_2 \mu_2 - p_1 \mu_1 = \delta (> 0)$. Furthermore, we denote the SD of X_k by ν_{0k} when the null hypothesis

H_0 is true and by v_{1k} when the alternative hypothesis H_1 is true. Then, the sample size n needed for each arm is given by:

$$n = \frac{\left(Z_{1-\alpha/2} \sqrt{v_{01}^2 + v_{02}^2} + Z_{1-\beta} \sqrt{v_{11}^2 + v_{12}^2} \right)^2}{\delta^2}. \quad (4)$$

See Supplementary Equation (S4) in the Supplemental Technical Details for more detail. To use formula (4) to compute the sample size, the key prerequisite is to specify the 4 SDs (v_{01} , v_{02} , v_{11} , v_{12}), each depending on the corresponding percent consuming and the means and SDs of the amount of intake when the food is consumed, as given by equation (3).

To compute the sample size appropriately, we recommend the following steps.

Step 1. Specify the level of significance α , usually set to be at 5% for a randomized trial; for a historical trial, it may be at 10%.

Step 2. Specify the desired power ($1-\beta$) for the study, usually set to be between 80 and 95%.

Step 3. Specify the intervention effect size δ to be detected at power $1-\beta$. Note that the effect size is the difference between $p_2\mu_2$ and $p_1\mu_1$, not μ_2 and μ_1 .

Step 4. For the control arm, specify the probability that the food is consumed, as well as the mean and SD of the amount of intake when the food is consumed, that is, p_{01} , μ_{01} , and σ_{01} under H_0 and p_{11} , μ_{11} , and σ_{11} under H_1 .

Step 5. Similarly for the intervention arm, specify the probability that the food is consumed, as well as the mean and SD of the amount of intake when the food is consumed, that is, p_{02} , μ_{02} , and σ_{02} under H_0 and p_{12} , μ_{12} , and σ_{12} under H_1 .

When specifying the values of the design parameters, the 2 constraints under the null and the alternative hypotheses, $p_{02}\mu_{02} - p_{01}\mu_{01} = 0$ and $p_{12}\mu_{12} - p_{11}\mu_{11} = \delta$, must be satisfied. Values of the parameters in Steps 4 and 5 can be derived based on historical/existing data from subjects similar to those in the present study. Otherwise, some plausible educational guesses are required. With these prespecified values of the design parameters, the 4 SDs in (4) can then be derived using (3), and subsequently the sample size can be computed using formula (4).

We next demonstrate that using the ad hoc sample size formulae (1–2) for comparing 2 means is problematic. Under the null hypothesis H_0 of no intervention effect on the average amount of intake, it can be further assumed that $(p_{01}, \mu_{01}, \sigma_{01}) = (p_{02}, \mu_{02}, \sigma_{02})$, that is, the intervention has no effect on either the probability of consumption of the food or the mean amount of intake if the food is consumed. This implies that $v_{01} = v_{02}$, that is, the amount of intake X_k has a common SD for both arms under the null hypothesis. Moreover, for the control group, the assumption that $(p_{01}, \mu_{01}, \sigma_{01}) = (p_{11}, \mu_{11}, \sigma_{11})$, leading to $v_{01} = v_{11}$, is also reasonable under the alternative hypothesis H_1 . For this simple case, the sample size formula reduces to:

$$n = \frac{\left(\sqrt{2}Z_{1-\alpha/2}v_{01} + Z_{1-\beta} \sqrt{v_{01}^2 + v_{12}^2} \right)^2}{\delta^2}. \quad (5)$$

Note that under the alternative hypothesis, the 2 rates p_{11} and p_{12} and the 2 means μ_{11} and μ_{12} are subject to the constraint

that $p_{12}\mu_{12} - p_{11}\mu_{11} = \delta (> 0)$. Such constraint makes v_{12} always different from the other 3 SDs; see **Supplemental Figure 1**. Therefore, the appropriate sample size formula (5) yields much different results from the ad hoc sample size formulae (1–2) for 2 means with either equal or unequal SDs.

It is also worth noting that the mean θ_k and SD v_k of the amount of intake X_k differ considerably from their counterpart μ_k and deviation σ_k of the amount of intake given the food is consumed, especially in the presence of the relatively high likelihood that the food is not consumed. Hence the SDs σ_1 and σ_2 of the amount of consumption given the food is consumed cannot be used in lieu of v_1 and v_2 when computing the sample size.

Numerical Results and Comparisons

Numerical studies via direct calculation or Monte Carlo simulation were conducted to investigate the performance of the sample size formula (4), as compared with that of the ad hoc formula (1) for comparing 2 means, which is obtained from (4) by replacing v_{11} by v_{01} and v_{12} by v_{02} . In all settings, the 2-sided nominal significance level α is set to be 5%, and the power is set to be 90%. The 3 distributional parameters, the probability that the food is consumed, and the mean and SD of the amount of intake given the food is consumed, are set to be 0.6, 0.5, and 1 for both arms when the null hypothesis is true. We vary these parameters under the alternative hypotheses to compute the required sample sizes. For all alternative hypotheses under consideration, we assume that 1) the SD of the amount of intake given the food is consumed in the intervention arm is 0.1 larger than that in the control arm, that is, $\sigma_{12} = \sigma_{11} + 0.1$; 2) the mean of the amount of intake given the food is consumed is 0.4 for the control arm, that is, $\mu_{11} = 0.4$; and 3) the intervention increases the probability that the food is consumed by 5%, i.e., $p_{12} = p_{11} + 0.05$. The intervention effect size is then derived from $\delta = p_{12}\mu_{12} - p_{11}\mu_{11}$ by plugging in the above alternative distributional parameters.

With each configuration of the design parameters, the sample sizes were calculated from equations (1) and (4) and are presented in **Table 1**, in the row labeled as “Ad hoc” and “Appropriate,” respectively. As demonstrated from **Table 1**, the sample sizes calculated using the ad hoc sample size formula differ substantially from that using the appropriate approach. Nevertheless, there is no clear pattern between them; in many cases the former is much larger and many others much smaller.

For each sample size n in **Table 1**, the power of the test was estimated based on 10,000 Monte Carlo replicates. For each replicate, the number of nonzeros (i.e., number of participants who consumed the food) in the control and intervention arm was generated from the Bernoulli distribution with success probability p_{11} and p_{12} , respectively. Subsequently, the corresponding amount of intake was generated from the normal distribution with mean μ_{11} (SD σ_{11}) and mean μ_{12} (SD σ_{12}), respectively.

The empirical power results are presented in **Figure 1**. It is evident that the proposed formula yields sample sizes that satisfactorily achieve the 90% nominal power (blue line), whereas the ad hoc formula fails to do so. As shown in **Figure 1**,

TABLE 1 Sample sizes per arm from the proposed (“Appropriate”) and ad hoc approach

| μ_{12} | p_{11} | $\sigma_{11} = 0.6$ | | | | $\sigma_{11} = 1.0$ | | | | $\sigma_{11} = 1.4$ | | | |
|------------|-------------|---------------------|------|------|------|---------------------|------|------|------|---------------------|------|------|------|
| | | 0.6 | 0.7 | 0.8 | 0.9 | 0.6 | 0.7 | 0.8 | 0.9 | 0.6 | 0.7 | 0.8 | 0.9 |
| 1.0 | δ | 0.41 | 0.47 | 0.53 | 0.59 | 0.41 | 0.47 | 0.53 | 0.59 | 0.41 | 0.47 | 0.53 | 0.59 |
| | Appropriate | 69 | 53 | 42 | 34 | 90 | 72 | 59 | 49 | 117 | 95 | 78 | 66 |
| | Ad hoc | 83 | 63 | 49 | 40 | 83 | 63 | 49 | 40 | 83 | 63 | 49 | 40 |
| 0.8 | δ | 0.28 | 0.32 | 0.36 | 0.40 | 0.28 | 0.32 | 0.36 | 0.40 | 0.28 | 0.32 | 0.36 | 0.40 |
| | Appropriate | 142 | 111 | 90 | 73 | 190 | 153 | 126 | 105 | 247 | 202 | 169 | 144 |
| | Ad hoc | 177 | 135 | 107 | 87 | 177 | 135 | 107 | 87 | 177 | 135 | 107 | 87 |

σ_{11} and σ_{12} are the SDs in the control and intervention arm of the amount of intake, given the food is consumed, under the alternative hypothesis; μ_{12} is the mean under the alternative hypothesis of the amount of intake given the food is consumed in the intervention arm; p_{11} is the probability that the food is consumed in the control arm; δ is the intervention effect size. The test statistic T_2 in the Supplemental Technical Details was used to obtain the results.

the sample sizes from the ad hoc formula (red line) can either seriously underpower the study (thus compromising the ability of the trial to detect meaningful intervention effects) or unnecessarily overpower the study (thus increasing the financial and administrative burden to conduct the trial). In addition, we also compared the type I error rates of the tests based on the 2 formulae; see **Supplemental Table 1**. The results show that the proposed approach adequately controls, whereas the ad hoc approach tends to inflate, the type I error rates when

the probabilities that the food is consumed differ between the 2 arms.

Discussion

For 2-arm parallel dietary intervention trials targeting episodically consumed foods, we have demonstrated that sample sizes computed using ad hoc methods are not adequate and proposed an appropriate approach for sample size calculation. As the

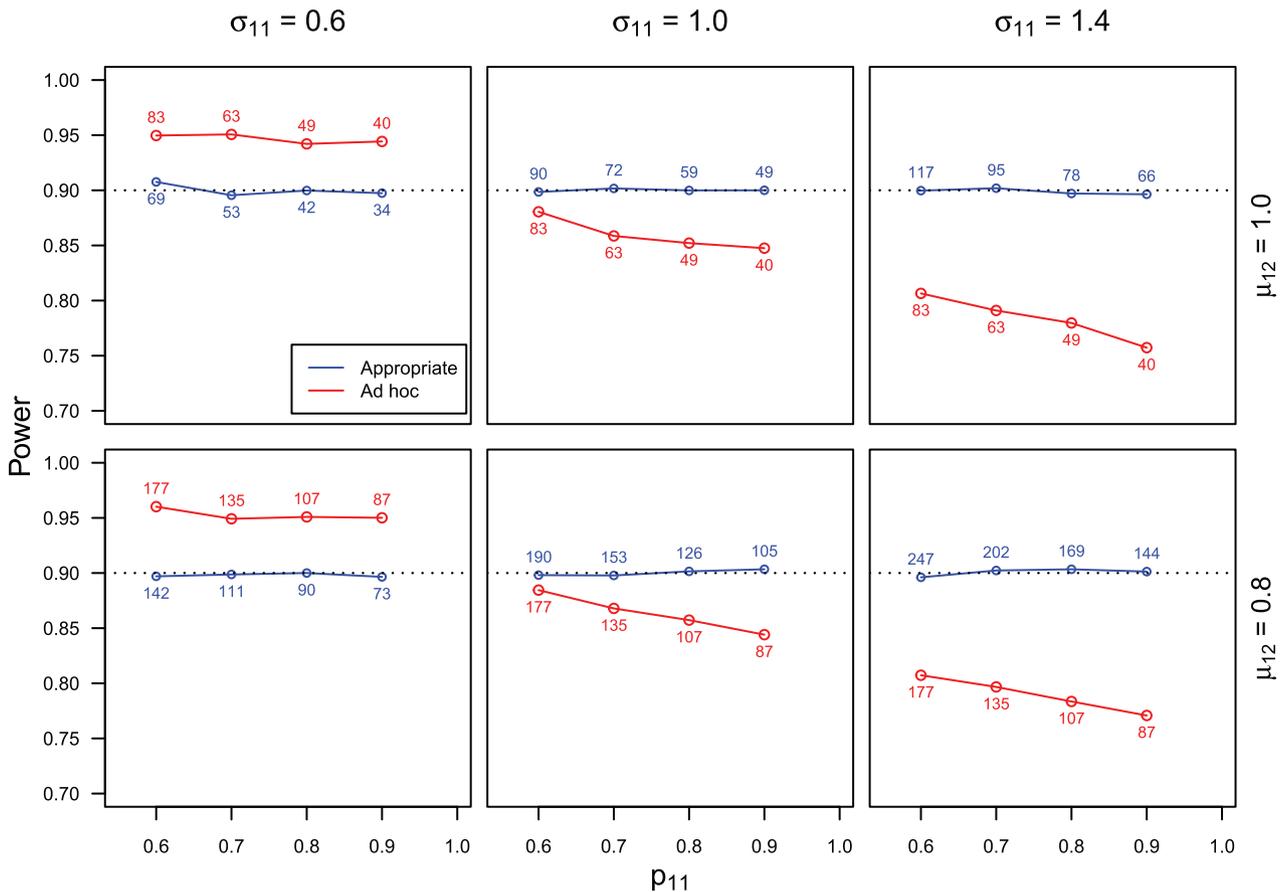


FIGURE 1 Empirical powers of the test for the sample sizes (annotated as text label) calculated using the proposed and ad hoc approach based on the nominal significance level of 0.05 and power of 90%. The blue lines are for the proposed approach and the red lines are for the ad hoc approach. The results are obtained based on the test statistic T_2 in the Supplemental Technical Details.

numerical results showed, the proposed approach adequately controls the type I error rates (see Supplemental Table 1) and achieves the desired power for the study. Hence, we recommend that investigators use our proposed method rather than the ad hoc methods for designing intervention studies targeting episodically consumed foods. It is worth pointing out that the difference between the proposed formula and the ad hoc formula will decrease as the probability that the food is consumed increases. When the probability is close to 1, the ad hoc formula can serve as a good approximation to the proposed one.

We have also considered 2 other types of trial designs, the 1-arm trial and prepost trial. The former evaluates the efficacy of an intervention scheme by comparing the dietary intake of the study participants under intervention with the available dietary records in a comparable general population. The latter does so by comparing dietary intake collected before and after the intervention; for such a prepost trial, the correlation between the dietary intake before and after intervention must be accounted for when computing the sample sizes, resulting in more complex formulae. In the Supplemental Technical Details, we have provided detailed derivation of the sample size formulae for the 2 types of trials.

Longitudinal dietary intake data collected from diet recalls at multiple time points across the study duration are common in dietary intervention studies. Sample size calculations technically become more complicated since the proportion of participants who consumed the food and the average amount of intake when the food is consumed could differ between treatment arms and between time points. Moreover, dietary data at any 2 time points from the same study participant are correlated. Other complex dietary intervention studies such as crossover trials and community-based trials also yield correlated data. The sample size calculations and power analysis need to properly incorporate all these aspects; thus, further research expanding upon these methods is needed.

In addition to the mean difference, the median difference of the amount of intake between the 2 arms can be used as a measure of intervention effect. In this case, nonparametric tests which are robust against distributional assumptions are preferred. However, for the food with a disproportionately large number of zeros, common nonparametric measures such as the Wilcoxon rank test would lead to many ties, which may result in substantial loss of power. To address this issue, Hallstrom (46) proposed a truncated Wilcoxon test by removing an equal (and maximal) number of zeros from each arm, but its performance depends on the respective difference direction of the proportion of consumption of the food and the average amount of intake of the food if consumed between the 2 arms. Further research is needed along this line.

The sample size required for a clinical trial depends on the null hypothesis and its corresponding test statistic. In the present article, the sample size calculations are based on testing the equality of the overall mean intake, i.e., $p_1\mu_1 = p_2\mu_2$. To the best of our knowledge, most dietary intervention studies published in the literature used the overall mean intake as the measure of intervention effect and success. Alternatively, success of the intervention can be defined as an increase in either the proportion of the food consumed or the mean intake when the food is consumed. Accordingly, the null hypothesis becomes

$p_1 = p_2$ and $\mu_1 = \mu_2$, and a global test such as the χ^2 test (47) can be used. However, the global test is less informative upon rejection of the null hypothesis because it only tells that ≥ 1 equality is rejected, whereas in contrast, the proposed test provides additional information on whether the overall mean intake differs between the 2 arms. In practice, when designing a dietary intervention trial with episodically consumed foods, the choice of the null hypothesis to be tested should depend on the primary scientific interest which uniquely defines the measure of effect and success of the intervention.

Semicontinuous data are frequently encountered in other research areas such as health expenditures, hospital length of stay, physical activities, and daily alcohol consumption (28). Our proposed methods for sample size calculation and power analysis are also recommended for designing an intervention study in these areas. It is worth noting that the power calculation formula is derived based on large sample theory. Studying the exact distribution for small samples of semicontinuous data is an important direction for future research.

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