Some details on two-phase variances

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This document explains the computation of variances for totals in two-phase designs. Variances for other statistics are computed by the delta-method from the variance of the total of the estimating functions.

The variance formulas come from conditioning on the sample selected in the first phase

$$\text{var}[\hat{T}] = E\left[\text{var}\left[\hat{T}|\text{phase 1}\right]\right] + \text{var}\left[E\left[\hat{T}|\text{phase 1}\right]\right]$$

The first term is estimated by the variance of \hat{T} considering the phase one sample as the fixed population, and so uses the same computations as any single-phase design. The second term is the variance of \hat{T} if complete data were available for the phase-one sample. This takes a little more work.

The variance computations for a stratified, clustered, multistage design involve recursively computing a within-stratum variance for the total over sampling units at the next stage. That is, we want to compute

$$s^{2} = \frac{1}{n-1} \sum_{i=1}^{n} (X_{i} - \bar{X})$$

where X_i are π -expanded observations, perhaps summed over sampling units. A natural estimator of s^2 when only some observations are present in the phase-two sample is

$$\hat{s}^2 = \frac{1}{n-1} \sum_{i=1}^n \frac{R_i}{\pi_i} (X_i - \hat{\bar{X}})$$

where π_i is the probability that X_i is available and R_i is the indicator that X_i is available. We also need an estimator for \bar{X} , and a natural one is

$$\hat{\bar{X}} = \frac{1}{n} \sum_{i=1}^{n} \frac{R_i}{\pi_i} X_i$$

This is not an unbiased estimator of s^2 unless $\hat{\bar{X}} = \bar{X}$, but the bias is of order $O(n_2^{-1})$ where $n_2 = \sum_i R_i$ is the number of phase-two observations.

If the phase-one design involves only a single stage of sampling then X_i is Y_i/p_i , where Y_i is the observed value and p_i is the phase-one sampling probability. For multistage phase-one designs (not yet implemented) X_i will be more complicated, but still feasible to automate.

This example shows the unbiased phase-one estimate (from Takahiro Tsuchiya) and the estimate I use, in a situation where the phase two sample is quite small.

First we read the data

```
rei<-read.table(textConnection(</pre>
        N n.a h n.ah n.h
1
   1 300
          20 1
                  12
                       5
                          TRUE 1
    2 300
           20 1
                  12
                       5
                          TRUE
3
   3 300
           20 1
                  12
                       5 TRUE 3
   4 300
           20 1
                  12
                       5 TRUE
5
   5 300
           20 1
                  12
                       5 TRUE 5
    6 300
           20 1
                  12
                       5 FALSE NA
7
   7 300
          20 1
                  12
                       5 FALSE NA
   8 300
          20 1
                  12
                       5 FALSE NA
9
   9 300
          20 1
                  12
                      5 FALSE NA
10 10 300
          20 1
                  12
                       5 FALSE NA
11 11 300
          20 1
                  12
                      5 FALSE NA
12 12 300
          20 1
                  12 5 FALSE NA
13 13 300
                      3 TRUE 6
          20 2
                   8
14 14 300
          20 2
                   8
                      3 TRUE 7
15 15 300
          20 2
                     3 TRUE 8
                   8
16 16 300
          20 2
                   8 3 FALSE NA
17 17 300
          20 2
                       3 FALSE NA
                   8
18 18 300
          20 2
                   8
                       3 FALSE NA
19 19 300 20 2
                       3 FALSE NA
20 20 300 20 2
                       3 FALSE NA
                   8
"), header=TRUE)
  Now, construct a two-phase design object and compute the total of y
> des.rei <- twophase(id = list(~id, ~id), strata = list(NULL,</pre>
      "h), fpc = list("N, NULL), subset = "sub, data = rei)
> tot <- svytotal(~y, des.rei)</pre>
  The unbiased estimator is given by equation 9.4.14 of Särndal, Swensson, & Wretman.
> rei$w.ah <- rei$n.ah/rei$n.a
> a.rei <- aggregate(rei, by = list(rei$h), mean, na.rm = TRUE)
> a.rei$S.ysh <- tapply(rei$y, rei$h, var, na.rm = TRUE)
> a.rei$y.u <- sum(a.rei$w.ah * a.rei$y)
> a.rei$f <- with(a.rei, n.a/N)</pre>
> a.rei$delta.h <- with(a.rei, (1/n.h) * (n.a - n.ah)/(n.a - 1))
> Vphase1 \leftarrow with(a.rei, sum(N * N * ((1 - f)/n.a) * (w.ah * (1 - f)/n.a))
      delta.h) * S.ysh + ((n.a)/(n.a - 1)) * w.ah * (y - y.u)^2))
  The phase-two contributions (not shown) are identical. The phase-one contributions are quite
> Vphase1
[1] 24072.63
> attr(vcov(tot), "phases")$phase1
         [,1]
[1,] 23461.05
```