

On Small Area Estimation under Informative Sampling

Danny Pfeffermann¹ and Michael Sverchkov²

¹Hebrew University and University of Southampton, ²Bureau of Labor Statistics and User Technology Associates, Inc.

²Bureau of Labor Statistics, Room 1950, 2 Massachusetts Avenue, NE, Washington, DC 20212

SUMMARY

Classical small area estimation techniques assume either that all the areas are represented in the sample or that the selection of the areas to the sample is noninformative. When the areas are sampled with unequal selection probabilities that are related to the values of the response variable, the classical estimators are biased; the magnitude of the bias depends on the sampling fraction and the covariance between the sampling weights and the response variable. We illustrate this point using very simple models employing the notions of the sample distribution and sample-complement distribution. We suggest simple unbiased estimators based on these distributions.

Key words: Sample distribution, Sample-complement distribution

1. The sample and sample-complement distributions

Consider a finite population U consisting of N units belonging to M areas, with N_i units in area i ,

$\sum_{i=1}^M N_i = N$. Let y define the study variable with value y_{ij} for unit j in area i and denote by x_{ij} the values of auxiliary (covariate) variables that are possibly known for that unit. In what follows we consider the population y -values as random realizations of the following two level stochastic process:

First level- values (*random effects*) $\{u_1 \dots u_M\}$ are generated independently from some distribution with probability density function (*pdf*) $f_p(u_i)$ for which $E_p(u_i) = 0$; $E_p(u_i^2) = \sigma_u^2$, where E_p

defines the expectation operator; *Second level-* values $\{y_{i1} \dots y_{iN_i}\}$ are generated from some conditional distribution with *pdf* $f_p(y_{ij} | x_{ij}, u_i)$, for $i = 1 \dots M$. We assume a two-stage sampling

scheme which in the first stage selects m areas with inclusion probabilities $\pi_i = \Pr(i \in s)$ and in the second step n_i units are sampled from area i selected in the first step with inclusion

probabilities $\pi_{ji} = \Pr(j \in s_i | i \in s)$. Note that the sample inclusion probabilities at both stages may depend in general on all the population or area values of y , x and possibly design variables z , used for the sample selection but not included in the working model. Denote by I_i and I_{ij} the sample indicator

variables at the two stages ($I_i = 1$ iff $i \in s$ and similarly for I_{ij}) and by $w_i = 1/\pi_i$ and $w_{ji} = 1/\pi_{ji}$ the corresponding first and second stage sampling weights.

Following Pfeffermann *et. al* (1998), we define the conditional *sample pdf* of u_i , i.e., the first level conditional *pdf* of u_i for area $i \in s$ as,

$$f_s(u_i) \stackrel{def}{=} f(u_i | I_i = 1) \stackrel{Bayes}{=} \frac{\Pr(I_i = 1 | u_i) f_p(u_i)}{\Pr(I_i = 1)} \quad (1.1)$$

Similarly, the conditional *sample-complement pdf*, i.e., the conditional *pdf* of u_i for area $i \notin s$ is defined in Sverchkov and Pfeffermann (2001) as,

$$f_c(u_i) \stackrel{def}{=} f(u_i | I_i = 0) \stackrel{Bayes}{=} \frac{\Pr(I_i = 0 | u_i) f_p(u_i)}{\Pr(I_i = 0)} \quad (1.2)$$

Notice that the *population*, *sample* and *sample-complement pdfs* of u_i are the same iff $\Pr(I_i = 1 | u_i) = \Pr(I_i = 1) \forall i$, in which case the sampling of areas is *noninformative*.

The second level *sample pdf* and *sample-complement pdf* of y_{ij} are defined similarly to (1.1) and (1.2) as,

$$f_s(y_{ij} | x_{ij}, u_i) \stackrel{def}{=} f(y_{ij} | x_{ij}, u_i, I_{ij} = 1) = \frac{\Pr(I_{ij} = 1 | y_{ij}, \mathbf{x}_{ij}, u_i) f_p(y_{ij} | \mathbf{x}_{ij}, u_i)}{\Pr(I_{ij} = 1 | \mathbf{x}_{ij}, u_i)} \quad (1.3)$$

$$f_c(y_{ij} | x_{ij}, u_i) \stackrel{def}{=} f(y_{ij} | x_{ij}, u_i, I_{ij} = 0) = \frac{\Pr(I_{ij} = 0 | y_{ij}, \mathbf{x}_{ij}, u_i) f_p(y_{ij} | \mathbf{x}_{ij}, u_i)}{\Pr(I_{ij} = 0 | \mathbf{x}_{ij}, u_i)} \quad (1.4)$$

The model defined by (1.1) and (1.3) defines the two-level *sample model* analogue of the population model defined by $f_p(u_i | z_i)$ and $f_p(y_{ij} | x_{ij}, u_i)$; see also Pfeffermann *et. al* (2001).

The following relationships are established in Pfeffermann and Sverchkov (1999) and Sverchkov and Pfeffermann (2001) for general pairs of random variables v_1, v_2 measured for elements $i \in U$ where E_p, E_s and E_c denote expectations under the *population, sample* and *sample-complement* distributions and (π_i, w_i) define the sample inclusion probability and the sampling weight.

$$f_s(v_{1i} | v_{2i}) = f(v_{1i} | v_{2i}, i \in s) = \frac{E_p(\pi_i | v_{1i}, v_{2i}) f_p(v_{1i} | v_{2i})}{E_p(\pi_i | v_{2i})} \quad (1.5)$$

$$E_p(v_{1i} | v_{2i}) = \frac{E_s(w_i v_{1i} | v_{2i})}{E_s(w_i | v_{2i})} ; E_p(\pi_i | v_{2i}) = \frac{1}{E_s(w_i | v_{2i})} \quad (1.6)$$

$$\begin{aligned} f_c(v_{1i} | v_{2i}) &= f(v_{1i} | v_{2i}, i \notin s) = \frac{E_p[(1 - \pi_i) | v_{1i}, v_{2i}] f_p(v_{1i} | v_{2i})}{E_p[(1 - \pi_i) | v_{2i}]} \\ &= \frac{E_s[(w_i - 1) | v_{1i}, v_{2i}] f_s(v_{1i} | v_{2i})}{E_s[(w_i - 1) | v_{2i}]} \end{aligned} \quad (1.7)$$

$$E_c(v_{1i} | v_{2i}) = \frac{E_p[(1 - \pi_i) v_{1i} | v_{2i}]}{E_p[(1 - \pi_i) | v_{2i}]} = \frac{E_s[(w_i - 1) v_{1i} | v_{2i}]}{E_s[(w_i - 1) | v_{2i}]} \quad (1.8)$$

Defining $v_{1i} = u_i, v_{2i} = \text{constant}$ yields the relationships holding for the random area effects u_i . Defining $v_{1ij} = y_{ij}; v_{2ij} = (x_{ij}, u_i)$ and substituting π_{ji} and w_{ji} for π_i and w_i respectively yields the relationships holding for the observations y_{ij} .

2. Optimal Small Area Predictors

The target estimated population parameters are the small area means $\bar{Y}_i = \sum_{j=1}^{N_i} y_{ij} / N_i$ for $i = 1 \dots M$. Let $D_s = \{(y_{ij}, \pi_{ji}, \pi_i), (i, j) \in s; (I_{kl}, I_k, x_{kl}), (k, l) \in U\}$ define the known data.

The MSE of a predictor \hat{Y}_i given D_s with respect to the *population pdf* is,

$$\begin{aligned} MSE(\hat{Y}_i | D_s) &= E_p[(\hat{Y}_i - \bar{Y}_i)^2 | D_s] = E_p\{[\hat{Y}_i - E_p(\bar{Y}_i | D_s)]^2 | D_s\} + V_p(\bar{Y}_i | D_s) \\ &= [\hat{Y}_i - E_p(\bar{Y}_i | D_s)]^2 + V_p(\bar{Y}_i | D_s) \end{aligned} \quad (2.1)$$

The variance $V_p(\bar{Y}_i | D_s)$ does not depend on the form of the predictor and hence the MSE is minimized when $\hat{Y}_i = E_p(\bar{Y}_i | D_s)$. In what follows we distinguish between *sampled areas* ($I_i = 1$) and *nonsampled areas* ($I_i = 0$). Denote by s_i the sample of units in sampled area i . Then, for the sampled areas,

$$\begin{aligned}
E_p(\bar{Y}_i | D_s, I_i = 1) &= \frac{1}{N_i} \left\{ \sum_{j \in s_i} E_p(y_{ij} | D_s) + \sum_{l \notin s_i} E_p(y_{il} | D_s, I_{il} = 0) \right\} \\
&= \frac{1}{N_i} \left\{ \sum_{j \in s_i} y_{ij} + \sum_{l \notin s_i} E_c(y_{il} | D_s) \right\}
\end{aligned} \tag{2.2}$$

For areas i not in the sample,

$$\begin{aligned}
E_p(\bar{Y}_i | D_s, I_i = 0) &= \frac{1}{N_i} \sum_{k=1}^{N_i} E_p(y_{ik} | D_s, I_{ik} = 0) \\
&= \frac{1}{N_i} \sum_{k=1}^{N_i} E_c(y_{ik} | D_s)
\end{aligned} \tag{2.3}$$

The predictors in (2.2) and (2.3) can be written in a single equation as,

$$\begin{aligned}
E_p(\bar{Y}_i | D_s) &= \frac{1}{N_i} \left\{ \sum_{k=1}^{N_i} y_{ik} I_{ik} + \sum_{k=1}^{N_i} E_c[y_{ik} (1 - I_{ik}) | D_s] \right\} I_i \\
&\quad + \frac{1}{N_i} \left\{ \sum_{k=1}^{N_i} E_c[y_{ik} | D_s] \right\} (1 - I_i)
\end{aligned} \tag{2.4}$$

3. Bias of Small Area Predictors when ignoring the Sampling Scheme

Consider for convenience the case of a sampled area. Ignoring the sampling scheme implies an implicit assumption that the *sample-complement* model and the *sample* model are the same such that

$\hat{\bar{Y}}_{i,IGN} = \sum_{j \in s_i} y_{ij} + \sum_{l \notin s_i} E_s(y_{il} | D_s)$. Hence,

$$\begin{aligned}
E_p[(\hat{\bar{Y}}_{i,IGN} - \bar{Y}_i) | D_s, I_i = 1] &= \frac{1}{N_i} \sum_{l \notin s_i} [E_s(y_{il} | D_s) - E_c(y_{il} | D_s)] \\
&= -\frac{1}{N_i} \sum_{l \notin s_i} \frac{Cov_s(y_{il}, w_{li} | D_s)}{E_s[(w_{li} - 1) | D_s]}
\end{aligned} \tag{3.1}$$

with the second equality following from (1.8). Thus, unless the response values y_{il} and the ‘within’ sampling weights w_{li} are uncorrelated, ignoring the sampling scheme results in biased predictors (see also the empirical results). A similar expression for the bias can be obtained for the nonsampled areas.

A simple Example. Let the *population model* be the “unit level random effects model”

$$y_{ij} = \mu + u_i + e_{ij}; \quad u_i \sim N(0, \sigma_u^2), \quad e_{ij} \sim N(0, \sigma_e^2) \tag{3.2}$$

with all the random effects and residual terms being mutually independent.

Let $\pi_i = c \times N_i$ where c is some constant and $\pi_{ji} = n_0 / N_i$ (fixed sample size n_0 within the selected areas), such that $\pi_{ij} = \Pr[(i, j) \in s] = \pi_i \pi_{ji} = \text{const}$. Note that the sample selection within the selected areas is *noninformative* in this case but if the area sizes N_i are correlated with the random effects u_i (say, the areas are *schools*, the study variable measures children’s attainment, the large schools are in the poor areas), the selection of the areas is *informative*.

Suppose that the areas sizes can be modeled as $\log(N_i) \sim N(Au_i, \sigma_M^2)$, implying that

$E_p(\pi_i | u_i) \propto \exp(Au_i + \frac{\sigma_M^2}{2})$ by familiar properties of the lognormal distribution. It follows that (see Pfeffermann *et al.* 1998, example 4.3),

$$f_s(u_i) = \frac{E_p(\pi_i | u_i) f_p(u_i)}{E_p(\pi_i)} = N(A\sigma_u^2, \sigma_u^2) \tag{3.3}$$

so that $E_s(u_i) = \gamma\sigma_u^2 \neq E_p(u_i) = 0$. The fact that the random effects in the sample have in this case a positive expectation is easily explained by the fact that the sampling scheme considered tends to select the areas with large positive random effects. Note, however, that by defining $\mu^* = \mu + A\sigma_u^2$ and $u_i^* = u_i - A\sigma_u^2$, the model holding for the sample data in sampled areas is $y_{ij} = \mu^* + u_i^* + e_{ij}$, $u_i^* \sim N(0, \sigma_u^2)$, $e_{ij} \sim N(0, \sigma_e^2)$, which is the same as the population model. Thus, the optimal predictors under the *population model* for the area means $\theta_i = \mu + u_i$ of the sampled areas ($I_i = 1$) are still optimal under the *sample model*

Next consider *nonsampled areas*. By (1.7),

$$f_c(u_i) = \frac{E_p[(1 - \pi_i) | u_i] f_p(u_i)}{E_p(1 - \pi_i)} = \frac{f_p(u_i)}{E_p(1 - \pi_i)} - \frac{E_p(\pi_i | u_i) f_p(u_i)}{E_p(1 - \pi_i)} \quad (3.4)$$

Let $E_p(m) = E_p[\sum_{l=1}^M I_l] = E_p[E_p(\sum_{l=1}^M I_l | \{N_i\})] = E_p[\sum_{l=1}^M \pi_i] = ME_p(\pi_i)$ define the expected number of sampled areas, such that $E_p(\pi_i) = E_p(m) / M$. If the number of sampled areas is fixed, $E_p(m) = m$. By (3.4) and (1.5),

$$f_c(u_i) = [Mf_p(u_i) - E_p(m)f_s(u_i)] / [M - E_p(m)] \text{ and hence,}$$

$$E_c(u_i) = -\frac{E_p(m)E_s(u_i)}{M - E_p(m)} = -\frac{E_p(m)A\sigma_u^2}{M - E_p(m)} \quad (3.5)$$

Here again, the negative expectation of the random effects pertaining to *nonsampled areas* is easily explained by the tendency of the sampling scheme to select the areas with the large positive random effects. Thus, ignoring the sampling scheme underlying the selection of the areas and predicting the sample means in nonsampled areas by, say, the average of the predictors in the sampled areas yields in general biased predictors with a positive bias defined by the absolute value of the right hand side of (3.5).

4. On Small Area Estimation based on Sample Distribution

In order to illustrate the proposed approach, we suppose that the area level random effects model defined by (4.1) holds for the sampled areas, i.e., for $j \in S_i$

$$y_{ij} = \mu + u_i + e_{ij}; u_i | I = 1_i \sim N(0, \sigma_u^2), e_{ij} | I_{ij} = 1 \sim N(0, \sigma_e^2) \quad (4.1)$$

We mention in this respect that the sample model can be identified using conventional techniques, see, e.g., Rao (2003).

Suppose that in the first stage m areas are selected with inclusion probabilities π_i (m is fixed) and in the second stage n_i units are sampled from area i selected in the first stage with inclusion probabilities π_{ji} where again, we assume for convenience that the sample sizes n_i are fixed. Assume that,

$$E_s(w_{ji} | y_{ij}, u_i) = E_s(w_{ji} | y_{ij}) = c_i \exp(by_{ij}) \quad (4.2)$$

where $\theta_i = \mu + u_i$ and $c_i > 0$ and b are fixed parameters. Notice that since n_i is fixed, $E_s(w_{ji} | u_i) = N_i / n_i$.

Comment: As with the sample model (4.1), the expectation in (4.2) refers to the *sample distribution* within the areas. The relationship between the sampling weights and the observed data holding in the sample can be identified and estimated therefore from the sample data. See Pfeffermann and Sverchkov (1999, 2003) for discussion and examples. On the other hand, the relationship between the sampling weights w_i and the small area means $\theta_i = \mu + u_i$ is more difficult to detect since the area means are not observable and in what follows we do not model this relationship. See Pfeffermann *et al.* (2001) for an example of modeling the selection probabilities at both stages. As established in Section 2, the optimal predictor for areas *in the sample* is,

$E_p(\bar{Y}_i | D_s, I_i = 1) = [\sum_{j \in s_i} y_{ij} + \sum_{l \notin s_i} E_c(y_{il} | D_s, I_i = 1)] / N_i$. In order to compute the expectations $E_c(y_{il} | D_s, I_i = 1)$ we follow the following steps. First, by (1.7), (4.1) and (4.2),

$$\begin{aligned} f_c(y_{il} | \theta_i, I_i = 1) &= \frac{[E_s(w_{li} | y_{il}, \theta_i) - 1] f_s(y_{il} | \theta_i)}{E_s(w_{li} | \theta_i) - 1} \\ &= [c_i \exp(by_{il}) - 1] \frac{1}{\sigma_e} \phi\left[\frac{y_{il} - \theta_i}{\sigma_e}\right] / \left[\frac{N_i}{n_i} - 1\right] \\ &= \frac{n_i}{N_i - n_i} \left\{ c_i \exp(\theta_i b + \frac{\sigma_e^2 b^2}{2}) \frac{1}{\sigma_e} \phi\left[\frac{y_{il} - (\theta_i + b\sigma_e^2)}{\sigma_e}\right] - \frac{1}{\sigma_e} \phi\left[\frac{y_{il} - \theta_i}{\sigma_e}\right] \right\} \end{aligned} \quad (4.3)$$

where ϕ is the standard normal *pdf*. Notice that if $b = 0$ (noninformative selection within the sampled areas with equal inclusion probabilities), $c_i = N_i / n_i$ and the *pdf* in (4.3) reduces to the conditional normal density defined by (4.1). Second, by (4.3),

$$E_c(y_{il} | \theta_i, I_i = 1) = \frac{n_i}{N_i - n_i} \left\{ c_i \exp(\theta_i b + \frac{\sigma_e^2 b^2}{2}) [\theta_i + b\sigma_e^2] - \theta_i \right\} \quad (4.4)$$

Finally,

$$E_c(y_{il} | D_s, I_i = 1) = E_s[E_c(y_{il} | D_s, I_i = 1, \theta_i)] = E_s[E_c(y_{il} | I_i = 1, \theta_i)] \quad (4.5)$$

where the exterior expectation is with respect to the distribution of $\theta_i | D_s, I_i = 1$. Under the model (4.1), the latter distribution is normal with mean $\hat{\theta}_i = \gamma_i \bar{y}_i + (1 - \gamma_i) \bar{y}$ and variance $v_i = \gamma_i \sigma_i^2 + (1 - \gamma_i)^2 (\sum_{i=1}^m \gamma_i / \sigma_i^2)^{-1}$ where $\bar{y}_i = \sum_{j=1}^{n_i} y_{ij} / n_i$ is the sample mean in sampled area i , $\bar{y} = \sum_{i=1}^m n_i \bar{y}_i / \sum_{i=1}^m n_i$, $\sigma_i^2 = \sigma_e^2 / n_i = \text{Var}(\bar{y}_i | u_i)$ and $\gamma_i = \sigma_u^2 / [\sigma_u^2 + \sigma_i^2]$.

Thus, for the *sampled areas* $E_c(y_{il} | D_s, I_i = 1)$ is obtained by computing the expectation of the right hand side of (4.4) with respect to the normal distribution of $\theta_i | D_s, I_i = 1$. We find that,

$$E_c(y_{il} | D_s, I_i = 1) = \frac{n_i}{N_i - n_i} \left\{ c_i [\hat{\theta}_i + b(v_i + \sigma_e^2)] \exp[\hat{\theta}_i b + \frac{b^2}{2} (\sigma_e^2 + v_i)] - \hat{\theta}_i \right\} \quad (4.6)$$

Notice that if $b=0$ (noninformative sampling within the areas with equal inclusion probabilities) $c_i = N_i / n_i$ and $E_c(y_{il} | D_s, I_i = 1) = \hat{\theta}_i$.

Comment: The optimal predictor obtained for the case of noninformative sampling, $E_p(\bar{Y}_i | D_s, I_i = 1) = [\sum_{j \in s_i} y_{ij} + (N_i - n_i) \hat{\theta}_i] / N_i$ (Eq. 2.2) is different from the common predictor, $\hat{\theta}_i$. This is so because the target parameter is defined to be the finite area mean \bar{Y}_i rather than θ_i . See also Prasad and Rao (1990).

For the *nonsampled areas* the optimal predictor is defined in (2.3) to be, $E_p(\bar{Y}_i | D_s, I_i = 0) = \sum_{k=1}^N E_c(y_{ik} | D_s, I_i = 0) / N_i$. In order to compute the expectations $E_c(y_{ik} | D_s, I_i = 0)$ we note first that

$$f_p(y_{ij} | \theta_i, I_i = 1) = f_p(y_{ij} | \theta_i, I_i = 0) = f_p(y_{kl} | \theta_k) \quad (4.7)$$

signifying that conditionally on the area means θ_i , the *population pdf* is the same for all the areas irrespective of whether the areas are sampled or not. The *pdf* $f_p(y_{il} | \theta_i)$ is obtained from (1.5), (1.6) and (4.2) similarly to the derivation of $f_c(y_{il} | \theta_i, I_i = 1)$ in (4.3) as,

$$\begin{aligned}
f_p(y_{il} | \theta_i) &= \frac{E_s(w_{li} | y_{il}, \theta_i) f_s(y_{il} | \theta_i)}{E_s(w_{li} | \theta_i)} \\
&= \frac{c_i n_i}{N_i} \exp(\theta_i b + \frac{\sigma_e^2 b^2}{2}) \frac{1}{\sigma_e} \phi\left[\frac{y_{il} - (\theta_i + b\sigma_e^2)}{\sigma_e}\right]
\end{aligned} \tag{4.8}$$

Notice that the population *pdf* is different from the sample *pdf* defined by (4.1) unless the sampling scheme within the areas is noninformative ($b=0$).

By (4.8),

$$E_p(y_{il} | \theta_i) = \frac{c_i n_i}{N_i} \exp(\theta_i b + \frac{\sigma_e^2 b^2}{2}) (\theta_i + b\sigma_e^2) \tag{4.9}$$

Now,

$$\begin{aligned}
E_c(y_{ik} | D_s, I_i = 0) &\stackrel{Def}{=} E_p(y_{ik} | D_s, I_i = 0) \\
&= E_p[E_p(y_{ik} | \theta_i, D_s, I_i = 0) | D_s, I_i = 0] \\
&= E_p[E_p(y_{ik} | \theta_i) | D_s, I_i = 0] \stackrel{Def}{=} E_c[E_p(y_{ik} | \theta_i) | D_s]
\end{aligned} \tag{4.10}$$

where the exterior expectations in the last row are with respect to the conditional distribution $f_p(\theta_i | D_s, I_i = 0) = f_c(\theta_i | D_s)$.

Finally, by (1.8) and (4.10),

$$E_c(y_{ik} | D_s, I_i = 0) = E_c[E_p(y_{ik} | \theta_i) | D_s] = E_s\left[\frac{(w_i - 1)E_p(y_{ik} | \theta_i)}{E_s(w_i | D_s) - 1} | D_s\right] \tag{4.11}$$

Denoting $\theta_{i,p} = E_p(y_{ik} | \theta_i)$, an estimator of the expectation $E_c(y_{ik} | D_s, I_i = 0)$ is obtained from (4.11) as,

$$\hat{E}_c(y_{ik} | D_s, I_i = 0) = \frac{1}{m} \sum_{i \in s} \frac{(w_i - 1)\hat{\theta}_{i,p}}{\hat{E}_s(w_i | D_s) - 1} = \sum_{i \in s} \frac{(w_i - 1)\hat{\theta}_{i,p}}{\sum_{i \in s} (w_i - 1)} \tag{4.12}$$

where $\hat{E}_s(w_i | D_s) = \frac{1}{m} \sum_{i \in s} w_i$ and $\hat{\theta}_{i,p}$ is obtained by substituting θ_i in (4.9) by $\hat{\theta}_i = \gamma_i \bar{y}_i + (1 - \gamma_i) \bar{y} = E_s(\theta_i | D_s, I_i = 1)$ or by use direct Hajek estimator of $\theta_{i,p} = E_p(y_{ik} | \theta_i)$. Notice that the right hand side of (4.12) defines the predictor of the mean θ_i 's in the *nonsampled* areas.

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