

CALIBRATING ON PRINCIPAL COMPONENTS IN THE PRESENCE OF MULTIPLE AUXILIARY VARIABLES FOR NONRESPONSE ADJUSTMENT

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Abstract: Nonresponse is a major impediment to valid inference in sample surveys. In the nonresponse scenario, the driver of successful estimation is the efficient use of available auxiliary information. As electronic devices provide considerable data storage capacities, at the estimation stage it is natural for survey statisticians to face large datasets of auxiliary variables. It is unwise to use all available data as doing so may lead to poor estimators, especially if some variables are strongly correlated. Furthermore, selecting a subset of available auxiliary variables may not be the best alternative given the issues related to selection criteria. In this paper, we propose reducing the dimensions of the original set of auxiliary variables by using principal components. The use of principal components in place of the original auxiliary variables is evaluated via two calibration approaches, linear calibration using no explicit response model and propensity calibration of a known response model. For the latter, we propose selecting components based on their canonical correlation with the model variables. The results of two simulation studies suggest that using principal components is appropriate, as it offers the great advantage of reducing the computational burden.

1. Introduction

When adjusting for nonresponse in sample surveys, auxiliary information plays a prominent role in successful estimation. Rizzo, Kalton and Brick (1996) note that, providing it is carefully chosen, the particular adjustment scheme used at the estimation stage is not that important. The relation with the study variable or response pattern is usually taken as a benchmark in the choice of auxiliary variables (see Kreuter and Olson, 2011; Särndal and Lundström, 2005, p. 110).

Calibration estimation (Deville and Särndal, 1992), initially designed to reduce sampling error in surveys with complete response, was eventually extended to surveys affected by nonresponse, (see

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e.g. Lundström and Särndal, 1999; Kott, 2006). The method relies on an efficient choice of auxiliary variables.

When many auxiliary variables are available, calibrating on all of them may lead to ‘over-calibration’, the term used by Guggemos and Tillé (2010). According to Särndal and Lundström (2005), a problem may arise when the candidate auxiliary vector contains variables likely to cause multicollinearity or variables with highly skewed distributions. These problems may result in a very inefficient estimator almost less efficient than, for example, the Horvitz-Thompson estimator (Cardot, Goga and Shehzad, 2015).

Large sets of auxiliary variables have also been considered by many authors in various estimation settings, as in the following examples, Bardsley and Chambers (1984) propose a ridge-type estimator in the context of model-based estimation, an approach that relaxes the principle that the calibration weights ‘exactly’ reproduce the totals of known characteristics by holding only ‘approximately’. Guggemos and Tillé (2010) introduce a penalized calibration estimator. Bilen, Khan and Yadav (2004) suggest a principal component approach for reducing the multicollinearity and dimensions of the auxiliary variables in a regression context. Cardot et al. (2015) propose calibration on reduced data via principal components (PCs) in surveys with complete response.

Variable selection criteria are also suggested in the literature as an alternative way to deal with large sets of auxiliary variables and related problems. McHenry (1978) suggests an algorithm to select the best subset of auxiliary variables in the context of multiple regression or multivariate analysis. Silva and Skinner (1997) suggest a selection criterion based on the variability of the regression estimator. Särndal and Lundström (2007) propose a selection device based on the variability of estimated inverse propensities determined under the assumption that the auxiliary variables satisfy some pre-specified condition. The variable selection is conditioned on an increase in the variability of the inverse propensities. A potential auxiliary variable must predict the key survey variables and the propensities to respond. Geuzinge, Rooijen and Bakker (2000) propose a selection indicator based on the product of (a) the correlation between the auxiliary vector and the study variables and (b) the correlation between the auxiliary vector and the response propensity. When adjusting for nonresponse through regression estimation, Bethlehem and Schouten (2004) and Schouten (2007) propose a selection based on minimizing the maximal absolute bias of the estimator; the method relies on computing an interval for the maximal absolute bias and selecting those variables that minimize its width.

The common practice of using a subset of the full set of potential auxiliary variables and discarding others may result in the loss of important information. For example, in a regression context, it is known that the R^2 tends to decrease with the removal of regressors from the regression equation. This phenomenon can be interpreted in many ways, but in some cases is due to the loss of valuable information. Furthermore, most of the suggested selection algorithms are computationally intensive and, impractical for large sets of candidate auxiliary variables.

In this paper, we calibrate on reduced data via principal components. Thus, we account for the exponential growth in computing time due to dimensionality in the auxiliary data and most importantly, the problem of large weights due to outliers is also accounted turning the estimator more efficient. The idea was initially suggested by Cardot et al. (2015) in surveys with complete response, and we extend it to estimation in surveys affected by nonresponse. Furthermore, the ideas in Cardot et al. (2015) are centred on the Greg-type-calibration (the complete response linear

calibration), while we study this and the propensity score calibration estimators in the nonresponse context. Note that the use of principal components in weighting does not stand for data interpretation, but is a tool for alleviating the problem of managing high-dimensional auxiliary data. Specifically, the PCs approach assists in the construction of new auxiliary variables from the original variables by taking into account all available candidate variables through linear combinations. Furthermore, we implement a rejection of PCs based on their canonical correlation (Hotelling, 1939) with the model variables.

Two calibration estimators are considered in the paper:

1. Linear calibration (LC) using no explicit form of response model Särndal and Lundström (2005).
2. Instrumental variable or propensity score calibration (PSC) with an explicit form of response model (Chang and Kott, 2008).

This suggests two sources of auxiliary information for estimation: an $\mathbf{X}_{(N \times P)}$ data matrix carrying information on the N population elements of a P -dimensional vector of auxiliary variables and an $\mathbf{H}_{(m \times L)}$ data matrix carrying information on the m respondent elements of an L -dimensional vector of instrumental variables. The LC estimator uses only the first source of auxiliary information, while the PSC combines the two sources.

The rest of the article is organized as follows: Section 2 provides background information on calibration estimators for nonresponse adjustment; Section 3 provides a summary theoretical framework on principal components; Section 4 provides a theoretical combination of calibration estimators and principal components; Section 5 provides numerical support for Section 4; and the final section discusses the results.

2. Calibration Estimators

Define a finite population, U , of distinguishable units indexed by integers $1, 2, \dots, k, \dots, N$. A probability sample, s , of distinguishable elements indexed by integers $1, 2, \dots, k, \dots, n$ is drawn from U according to a probability sampling design, $p(s)$, yielding the first- and second-order inclusion probabilities, $\pi_k = P(k \in s) > 0$ and $\pi_{kl} = P(k \& l \in s) > 0$, respectively for all $k, l \in \{1, 2, \dots, N\}$, where $\pi_{kk} = \pi_k$. Suppose that data are observed for subset $r \subset s$ with $|r| = m$. The elements of r are assumed to be generated by a random process, $q(r)$, on s . Thus, each element $k \in r$ is associated with probability $\theta_k = P(k \in r | k \in s) > 0$. The random process $q(r)$ on a given s is usually termed a response mechanism, while θ_k is the response probability for the individual k . Here, it is assumed that events $k \in r$ and $l \in r$ for a given s are independent of one another given that $k \neq l$.

Calibration estimators were introduced by Deville and Särndal (1992) in the context of surveys with complete response; the approach was then extended to surveys affected by nonresponse. In this context, Särndal and Lundström (2005) define the calibration estimator for total $t_y = \sum_U y_k$ as,

$$\hat{t}_{ycal} = \mathbf{w}_{(r)}' \mathbf{y}_{(r)},$$

where $\mathbf{w}_{(r)} = \text{vec}\{w_k\}^m$ and $\mathbf{y}_{(r)} = \text{vec}\{y_k\}^m$ are m -dimensional column vectors of calibrated weights w_k and study variable values y_k respectively. The term ‘calibrated weights’ means that the weights

satisfy the calibration property $\mathbf{X}_{(r)}^t \mathbf{w}_{(r)} = \mathbf{T}_x$, where $\mathbf{T}_x = \sum_U \mathbf{X}_k$ and \mathbf{X}_k being the transpose of the k^{th} line of $\mathbf{X}_{(N \times P)}$. Calibrated weights, w_k , are constructed to be as close as possible to the reciprocals of the sample inclusion probabilities, $d_k = 1/\pi_k$, according to a distance metric $\Omega(\mathbf{w}_{(r)}; \mathbf{d}_{(r)})$, while satisfying the above calibration property. Using Lagrange reasoning, calibrated weights can be derived by minimizing $\Omega(\mathbf{w}_{(r)}; \mathbf{d}_{(r)}) + \gamma^t (\mathbf{T}_x - \mathbf{X}_{(r)}^t \mathbf{w}_{(r)})$, where γ is a column vector of Lagrange multipliers, $\mathbf{d}_{(r)} = \text{vec}\{d_k\}^m$. The resulting calibrated weights take the form

$$w_k = d_k h(\gamma^t \mathbf{X}_k), \quad (1)$$

where $d_k h_k = \psi^{-1}(\cdot, d_k)$, $\psi = \partial \Omega / \partial w$, given the assumptions in Deville and Särndal (1992).

A different choice of Ω leads to a different weight system (1). Deville and Särndal (1992) establish conditions under which any choice of distance function leads to estimators that are asymptotically equivalent to the regression estimator obtained through a Chi-square-type distance measure. Thus, the choice of distance measure may be influenced by the computational aspects or other properties of w_k , such as its non-negativity or degree of stability.

Using the Chi-square distance, i.e., $\Omega(\mathbf{w}_{(r)}; \mathbf{d}_{(r)}) = (\mathbf{w}_{(r)} - \mathbf{d}_{(r)})^t (2\mathbf{D})^{-1} (\mathbf{w}_{(r)} - \mathbf{d}_{(r)})$, with $\mathbf{D} = \text{diag}\{d_1, d_2, \dots, d_k, \dots, d_m\}$, leads to the linear calibrated weights of the form

$$w_k = d_k + d_k \gamma^t \mathbf{X}_k, \quad (2)$$

where $\gamma = \left(\mathbf{X}_{(r)}^t \mathbf{D} \mathbf{X}_{(r)} \right)^{-1} \left(\mathbf{T}_x - \mathbf{X}_{(r)}^t \mathbf{d}_{(r)} \right)$.

The linear calibration estimator for t_y is:

$$\hat{t}_{ycal} = \mathbf{w}_{(r)}^t \mathbf{y}_{(r)} = \mathbf{d}_{(r)}^t \mathbf{e}_{(r)} + \mathbf{T}_x^t \left(\mathbf{X}_{(r)}^t \mathbf{D} \mathbf{X}_{(r)} \right)^{-1} \mathbf{X}_{(r)}^t \mathbf{D} \mathbf{y}_{(r)}, \quad (3)$$

where $\mathbf{e}_{(r)} = \text{vec}\{e_k\}^m$ and $\mathbf{y}_{(r)} = \text{vec}\{y_k\}^m$ are m -dimensional column vectors of residuals $e_k = y_k - \hat{y}_k$ and study variable values y_k respectively, and $\hat{y}_k = \mathbf{X}_k^t \left(\mathbf{X}_{(r)}^t \mathbf{D} \mathbf{X}_{(r)} \right)^{-1} \mathbf{X}_{(r)}^t \mathbf{D} \mathbf{y}_{(r)}$.

In the complete response context, estimator (3) is equivalent to the GREG estimator (Särndal, Swensson and Wretman, 1992) derived under superpopulation model ξ , which assumes a linear relationship between the survey variable, y_k , and the auxiliary vector, \mathbf{X}_k , given by $\xi : y_k = \beta^t \mathbf{X}_k + \varepsilon_k$. Since, $\mathbf{X}_{(s)}^t \mathbf{d}_{(s)}$ is unbiased for \mathbf{T}_x , the weights (2) are on average equal to d_k which leads to zero average differences $y_k - \hat{y}_k$.

3. A Brief Summary of Principal Components

Suppose that \mathbf{X} is defined as in Section 1 except that each $\mathbf{X}_j, j = 1, \dots, P$ is rescaled to zero mean and unit variance, then, $\mathbf{X}^t \mathbf{X}$ is the covariance matrix of \mathbf{X} . Let $(\lambda_j, \mathbf{b}_j; j = 1, \dots, P)$ be eigenvalue-eigenvector pairs of $\mathbf{X}^t \mathbf{X}$. The j^{th} principal component is given by $\mathbf{Z}_j = \mathbf{b}_j^t \mathbf{X} = \sum_{l=1}^P \mathbf{b}_{lj} \mathbf{X}_l$ with the properties

$$\text{cov}(\mathbf{Z}_j, \mathbf{Z}_i) = \begin{cases} 0, & j \neq i \\ \lambda_i, & j = i \end{cases},$$

where \mathbf{b}_j is a P -dimensional column vector and the $\lambda_i, i = 1, \dots, P$ satisfy $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_P \geq 0$. The proportion of total variance accounted for by the first $R < P$ principal components is given by $(\sum_{i=1}^R \lambda_i / \sum_{i=1}^P \lambda_i) \times 100\%$.

Suppose now that $\mathbf{X} = \mathbf{X}_{(s)}$, that is, auxiliary data observed only at sample level. The covariance matrix of $\mathbf{X}_{(s)}$ is estimated without bias by $\mathbf{X}^t \mathbf{D} \mathbf{X}$, where $\mathbf{D} = \text{diag}\{d_1, d_2, \dots, d_k, \dots, d_n\}$. The estimated principal components are given by $\hat{\mathbf{Z}}_j = \hat{\mathbf{b}}_j^t \mathbf{X}_{(s)} = \sum_{l=1}^P \hat{\mathbf{b}}_{lj} \mathbf{X}_{l(s)}$. The pair $(\hat{\lambda}_j, \hat{\mathbf{b}}_j; j = 1, \dots, P)$ comprise the eigenvalue and eigenvector of $\mathbf{X}^t \mathbf{D} \mathbf{X}$.

4. Calibrating on Principal Components

The calibration estimator in the principal components setting can be derived by solving the following problem:

$$\min \Omega(\mathbf{w}_{(r)}^{pc}; \mathbf{d}_{(r)}) \quad \text{s.t.} : \mathbf{Z}_{(r)}^t \mathbf{w}_{(r)}^{pc} = \mathbf{T}_z.$$

4.1. The linear calibration estimator based on principal components

If we follow the same reasoning that led to weights (2), we will then arrive at principal components calibrated weights given by

$$\mathbf{w}_k^{pc} = d_k + d_k \gamma_{(pc)}^t \mathbf{Z}_k,$$

where $\gamma_{(pc)} = \left(\mathbf{Z}_{(r)}^t \mathbf{D} \mathbf{Z}_{(r)} \right)^{-1} \left(\mathbf{T}_z^t - \mathbf{Z}_{(r)}^t \mathbf{d}_{(r)} \right)$ and $\mathbf{Z}_k = \{Z_{k1}, Z_{k2}, \dots, Z_{kR} | R < P\}$ is the vector whose elements are the retained components. The nonresponse principal-components-based calibration estimator for t_y is given by

$$\hat{t}_{ycal(pc)} = \mathbf{d}_{(r)}^t \mathbf{e}_{(r)}^{pc} + \mathbf{T}_z^t \left(\mathbf{Z}_{(r)}^t \mathbf{D} \mathbf{Z}_{(r)} \right)^{-1} \mathbf{Z}_{(r)}^t \mathbf{D} \mathbf{y}_{(r)},$$

where $\mathbf{e}_{(r)}^{pc} = \text{vec} \left\{ y_k - \mathbf{Z}_k^t \left(\mathbf{Z}_{(r)}^t \mathbf{D} \mathbf{Z}_{(r)} \right)^{-1} \mathbf{Z}_{(r)}^t \mathbf{D} \mathbf{y}_{(r)} \right\}^r$.

4.2. The propensity score calibration based on principal components

Consider a framework of unit response resulting according to a known parametric model, $\phi^{-1}(\cdot; \mathbf{H}_k)$. Observe that this model is known only up to an unknown L -dimensional vector of parameters, $\delta = \delta^*$, where $\delta \in \Upsilon$, $\dim(\mathbf{H}_k) = L \leq R$ and R is the number of selected PCs. Then, the model parameters can be estimated from the calibration constraint below (see Kott, 2012)

$$\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z = \mathbf{0}, \quad (4)$$

where $\dim(\mathbf{Z}_{(r)}) = m \times R$ and $\Phi(\delta) = \text{diag}\{\phi(\delta; \mathbf{H}_1), \phi(\delta; \mathbf{H}_2), \dots, \phi(\delta; \mathbf{H}_k), \dots, \phi(\delta; \mathbf{H}_m)\}$. This is a principle suggested by Chang and Kott (2008). The solution to (4) is the minimizer of the objective function

$$\left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right)^t \mathbf{W}_n \left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right). \quad (5)$$

When $L = R$, the form of weighting matrix \mathbf{W}_n of dimension $R \times R$ is irrelevant, as system (4) is just identified, otherwise \mathbf{W}_n is a suitably chosen nonnegative definite matrix. Note that \mathbf{Z}_k is an R -dimensional column vector of retained PCs of P originals. Under this setting, to make the system of equations (4) feasible, the minimal requirement is that the number of PCs in \mathbf{Z}_k be at least L retained components.

Having estimated the response model parameter, δ^* , the calibration estimator for t_y (the propensity score calibration) is

$$\hat{t}_{PSC(pc)} = \sum_r d_k \phi(\hat{\delta}_{(pc)}^t Z_k) y_k,$$

where $\hat{\delta}_{(pc)}$ is the estimated value of δ . To obtain $\hat{\delta}_{(pc)}$, we can follow the ideas by Beaumont (2006), who propose an iterative procedure based on the Taylor approximation of (4). This is similar to the procedure suggested by Binder (1983). We apply a slightly different perspective in the estimation of δ in (4).

Assume the following conditions to hold:

1. Function $\phi(\delta)$ is continuous and twice differentiable with respect to δ .
2. $E_{pq}(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z) = \mathbf{0}$ if and only if $\delta = \delta^*$ for all $\delta \in \Upsilon$
3. Set Υ is a compact set .
4. $E_{pq} \left[\left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right) \left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right)^t \right]$ is finite
5. $\mathbf{Z}_{(r)}^t \Psi(\delta) \mathbf{H} = \frac{\partial}{\partial \delta} \left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right) = \sum_r d_k \phi_1(\mathbf{H}_k; \delta) \mathbf{Z}_k \mathbf{H}_k^t$ exists and is continuous in Υ , where $\phi_1(\mathbf{H}_k; \delta) = \partial \phi(\mathbf{H}_k; \delta) / \partial \delta$ and the $m \times m$ diagonal matrix $\Psi(\delta)$ has its k^{th} diagonal element given by $d_k \phi_1(\mathbf{H}_k; \delta)$
6. $\mathbf{Z}_{(r)}^t \Psi(\delta) \mathbf{H}$ is a full-column rank matrix.

Define the quadratic distance as follows:

$$\left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right)^t \frac{\mathbf{W}_n}{2} \left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right). \quad (6)$$

The solution to (4) is defined as the minimizer of objective function (6). In the generalized method of moments setting, minimizing (6) is equivalent to solving the set of estimating equations defined by

$$\left(\mathbf{Z}_{(r)}^t \Psi(\delta) \mathbf{H} \right)^t \mathbf{W}_n \left(\mathbf{Z}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \mathbf{T}_z \right) = \mathbf{0}. \quad (7)$$

We use the following approximation:

$$\left(\mathbf{Z}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \mathbf{T}_z \right) \approx \left(\mathbf{Z}_{(r)}^t \Phi(\hat{\delta}_{(pc)}) \mathbf{d}_{(r)} - \mathbf{T}_z \right) + \left(\mathbf{Z}_{(r)}^t \Psi(\hat{\delta}_{(pc)}) \mathbf{H} \right) (\delta^* - \hat{\delta}_{(pc)}). \quad (8)$$

Introducing equation (8) into (7) yields the following updating equation:

$$\hat{\delta}_{(pc)}^1 \approx \hat{\delta}_{(pc)}^0 + \left[\left(\mathbf{Z}_{(r)}^t \Psi^0 \mathbf{H} \right)^t \mathbf{W}_n \left(\mathbf{Z}_{(r)}^t \Psi^0 \mathbf{H} \right) \right]^{-1} \left(\mathbf{Z}_{(r)}^t \Psi^0 \mathbf{H} \right)^t \mathbf{W}_n \left(\mathbf{Z}_{(r)}^t \Phi(\delta^0) \mathbf{d}_{(r)} - \mathbf{T}_z \right), \quad (9)$$

where $\Psi^0 = \Psi(\hat{\delta}_{(pc)}^0)$. In (5), $\hat{\delta}_{(pc)}$ is the value of $\hat{\delta}_{(pc)}^1$ obtained upon convergence of (9).

In Appendix A we provide the derivation of the asymptotic variances of the estimated coefficients of the propensity functions when population- or sample-level auxiliary information is used. A comparison of these variances shows that sample-level auxiliary information provides more accurate estimated coefficients than the population-level does.

4.3. Suggested retention criterion (a canonical correlation-based criterion)

Many authors have discussed PCs retention criteria, for example, Jollifé (1972), Cadima and Jollifé (1995), Jollifé, Trendafilov and Uddin (2003), and McCabe (1984), though there is no unified recommendation on this matter (Johnson and Wichern, 2007). Common practice is based on one or combinations of the following three criteria: the eigenvalue-one, scree plot, and proportion of total variance explained criteria. Mansfield, Webster and Gunst (1977) noted that it is common in PCs analysis for significant data variation to be accounted for by the first few components. According to these criteria, the components with small variability are excluded. Note, however that we are not concerned with interpreting PCs, instead using them as a tool for constructing new auxiliary variables that take into account all original candidate auxiliary variables.

In a canonical correlation setting, the goal is to determine sets of linearly independent vectors for two groups of variables that result in the maximum correlation between the projections of these variables onto the space spanned by these linearly independent vectors. According to Borga (2001), the correlation between two sets of multidimensional variables, if it exists, may be blurred if an inappropriate coordinate system is used to represent the variables. However, in canonical correlation, each of the two sets is linearly transformed, so that the corresponding pairs of coordinates of these transformed variables have the maximum correlation.

Recall that \mathbf{H} is an $m \times L$ data matrix where H_1, H_2, \dots, H_L are the model variables and let $\tilde{\mathbf{Z}}$ be an $m \times D$ data matrix, where $1 \leq D \leq P$ is the number of principal component variables in $\tilde{\mathbf{Z}}$. Let $\mathbf{P}_\mathbf{H}$ be the projection of \mathbf{H} onto the space spanned by linear combinations of its elements and suppose that $\mathbf{P}_{\tilde{\mathbf{Z}}}$ is the analogous projection of elements in $\tilde{\mathbf{Z}}$. We want to approximate the correlation ($\tilde{\rho}_{\mathbf{H}, \tilde{\mathbf{Z}}}$) of sets \mathbf{H} and $\tilde{\mathbf{Z}}$ by the canonical correlation defined by $\max_{\mathbf{P}_\mathbf{H}, \mathbf{P}_{\tilde{\mathbf{Z}}}} \Gamma(\mathbf{P}_\mathbf{H} \mathbf{H}^t, \mathbf{P}_{\tilde{\mathbf{Z}}} \tilde{\mathbf{Z}}^t)$.

$$\tilde{\rho}_{\mathbf{H}, \tilde{\mathbf{Z}}} \equiv \max_{\mathbf{P}_\mathbf{H}, \mathbf{P}_{\tilde{\mathbf{Z}}}} \Gamma(\mathbf{P}_\mathbf{H} \mathbf{H}^t, \mathbf{P}_{\tilde{\mathbf{Z}}} \tilde{\mathbf{Z}}^t) = \max_{\mathbf{P}_\mathbf{H}, \mathbf{P}_{\tilde{\mathbf{Z}}}} \frac{[\mathbf{P}_\mathbf{H} (\mathbf{H}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t]}{[\mathbf{P}_\mathbf{H} (\mathbf{H}^t \mathbf{H}) \mathbf{P}_\mathbf{H}^t]^{1/2} [\mathbf{P}_{\tilde{\mathbf{Z}}} (\tilde{\mathbf{Z}}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t]^{1/2}}. \quad (10)$$

We can equivalently reformulate (10) as

$$\begin{aligned} & \max \mathbf{P}_\mathbf{H} (\mathbf{H}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t, \\ & \text{s.t. } \begin{cases} [\mathbf{P}_\mathbf{H} (\mathbf{H}^t \mathbf{H}) \mathbf{P}_\mathbf{H}^t]^{1/2} = 1 \\ [\mathbf{P}_{\tilde{\mathbf{Z}}} (\tilde{\mathbf{Z}}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t]^{1/2} = 1. \end{cases} \end{aligned} \quad (11)$$

Using the Lagrange multiplier principle, (11) is solved by maximizing the objective function

$$\mathbf{L}(\mu_*, \mathbf{P}_*) = [\mathbf{P}_\mathbf{H} (\mathbf{H}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t] - 2^{-1} (\mu_1 [\mathbf{P}_\mathbf{H} (\mathbf{H}^t \mathbf{H}) \mathbf{P}_\mathbf{H}^t - 1] - \mu_2 [\mathbf{P}_{\tilde{\mathbf{Z}}} (\tilde{\mathbf{Z}}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t - 1])$$

yielding the system of equations

$$\begin{cases} \frac{\partial \mathbf{L}}{\partial \mathbf{P}_\mathbf{H}} = (\mathbf{H}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t - \mu_1 (\mathbf{H}^t \mathbf{H}) \mathbf{P}_\mathbf{H}^t = 0 \\ \frac{\partial \mathbf{L}}{\partial \mathbf{P}_{\tilde{\mathbf{Z}}}} = (\mathbf{H}^t \tilde{\mathbf{Z}})^t \mathbf{P}_\mathbf{H}^t - \mu_2 (\tilde{\mathbf{Z}}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t = 0. \end{cases} \quad (12)$$

Premultiplying the first equation in (12) by $\mathbf{P}_\mathbf{H}$ and subtracting $\mathbf{P}_{\tilde{\mathbf{Z}}}$ times the second equation from the first, results in $\mu_1 \mathbf{P}_\mathbf{H} (\mathbf{H}^t \mathbf{H}) \mathbf{P}_\mathbf{H}^t = \mu_2 \mathbf{P}_{\tilde{\mathbf{Z}}} (\tilde{\mathbf{Z}}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t$, because $\mathbf{P}_\mathbf{H} (\mathbf{H}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t = (\mathbf{P}_\mathbf{H} (\mathbf{H}^t \tilde{\mathbf{Z}}) \mathbf{P}_{\tilde{\mathbf{Z}}}^t)^t$, where by $\mu_1 = \mu_2 = \mu$.

Assuming that $\mathbf{H}'\mathbf{H}$ is invertible, the first equation gives

$$\mu \mathbf{P}'_{\mathbf{H}} = (\mathbf{H}'\mathbf{H})^{-1} (\mathbf{H}'\tilde{\mathbf{Z}}) \mathbf{P}'_{\tilde{\mathbf{Z}}}. \quad (13)$$

After appropriate replacements in the second, we get $(\mathbf{H}'\tilde{\mathbf{Z}})' (\mathbf{H}'\mathbf{H})^{-1} (\mathbf{H}'\tilde{\mathbf{Z}}) \mathbf{P}'_{\tilde{\mathbf{Z}}} - \mu^2 (\tilde{\mathbf{Z}}'\tilde{\mathbf{Z}}) \mathbf{P}'_{\tilde{\mathbf{Z}}} = 0$, which is equivalent to writing this last equation as

$$Qx = \lambda Rx, \quad (14)$$

where $Q = (\mathbf{H}'\tilde{\mathbf{Z}})' (\mathbf{H}'\mathbf{H})^{-1} (\mathbf{H}'\tilde{\mathbf{Z}})$, $x = \mathbf{P}'_{\tilde{\mathbf{Z}}}$, $\lambda = \mu^2$ and $R = (\tilde{\mathbf{Z}}'\tilde{\mathbf{Z}})$.

Equation (14) is in the form of a generalized eigenvalue equation (Parra and Sajda, 2003). Let, $R = MM'$ be a Cholesky decomposition of R ; then (14) becomes

$$(M^{-1}QM^{-1'})M'x = \lambda M'x \Leftrightarrow \tilde{Q}\tilde{x} = \lambda\tilde{x},$$

which is the standard eigenvalue equation. Solving this, we obtain a solution for $\mathbf{P}_{\tilde{\mathbf{Z}}}$, which naturally leads to a solution for $\mathbf{P}_{\mathbf{H}}$ in (13). These solutions represent the optimal projections of the variables in $\{\tilde{\mathbf{Z}}\}$ and $\{\mathbf{H}\}$ onto spaces spanned by their respective linear combinations. The coordinate systems resulting from $\mathbf{P}_{\tilde{\mathbf{Z}}}$, and $\mathbf{P}_{\mathbf{H}}$ are mutually maximally correlated. See, for example, Borga (2001) and Hardoon, Szedmak and Shawe-Taylor (2004), for more insight on canonical correlation analysis.

Our PCs selection criterion is based on the value of the canonical correlation between the PCs and the instrumental variables. The PCs are selected in order of their appearance and the canonical correlations are used to measure the representativeness of the selected components. The canonical correlations are calculated in a forward stepwise manner: the first canonical correlation is the correlation between the instrumental vector and a vector comprising the first PC; the second canonical correlation is the maximal correlation between the instrument vector and the vector comprising the first two PCs, and so on. The values of these canonical correlations are obtained in an increasing order. The stopping rule is based on the amount by which this correlation increases from a previous step to the actual step. If the addition of a further component to the vector of PCs does not significantly change the correlation among these two groups, then that component and the remaining components are discarded from the final auxiliary vector.

Remark 1 Unlike $\mathbf{Z}'\mathbf{Z}$, which is a diagonal matrix with eigenvalues of $\mathbf{X}'\mathbf{X}$ being its diagonal elements, matrix $\tilde{\mathbf{Z}}'\tilde{\mathbf{Z}}$ is no longer a diagonal since $\tilde{\mathbf{Z}}$ is made of elements of \mathbf{Z} falling into response set r .

Remark 2 We maximize the relation $(\mathbf{H}, \tilde{\mathbf{Z}})$ rather than (\mathbf{H}, \mathbf{Z}) as the latter is impossible because information on \mathbf{H} is assumed to be known at response level. The variables' distributions are generally distorted by nonresponse and the resulting correlation is expected to deviate from the true correlation. This is not of concern here, as the main goal is to guarantee at the response level selected auxiliary variables closely linked to the instruments.

5. Simulation Study

This section provides empirical illustrations of the points discussed in the previous sections. It is known that the principal components data reduction approach is effective when the relations among

the variables involved are strong. In this article we present two simulation studies: in the first study, the structure of correlation among the variables is very strong, the first principal component alone explaining more than 90% of the total data variation, as can be observed in Figure 1; in the second study, the structure of correlation among the variables is weak, and several components are needed to meaningfully explain the total variation of the data as illustrated by the scree plot shown in Figure 2. The data source for the first study is ‘Unemployment and median household income for the U.S., States, and counties, 2006–2014’ from the Unemployment – Bureau of Labor Statistics – LAUS data. The data are freely and publicly accessible for use at <http://www.bls.gov/laui/>. According to the source, *‘the concepts and definitions underlying LAUS data come from the Current Population Survey (CPS), the household survey that is the official measure of the labor force for the nation. State monthly model estimates are controlled in real time to sum to national monthly labor force estimates from the CPS. These models combine current and historical data from the CPS, the Current Employment Statistics (CES) program, and State unemployment insurance (UI) systems’*.

The data source for the second study is ‘Small Area Income and Poverty Estimates (SAIPE)’, which is a 1989, 1993, and 1995 – 2013 dataset, also freely and publicly accessible at <https://www.census.gov/did/www/saipe/>. According to the source, *‘Small Area Income and Poverty Estimates (SAIPE) are produced for school districts, counties, and states. The main objective of this program is to provide updated estimates of income and poverty statistics for the administration of federal programs and the allocation of federal funds to local jurisdictions’*.

5.1. Simulation setup

5.1.1. Study 1

From the data of the first study we selected 27 quantitative variables. We applied data transformation to induce the correlation among them to a desired pattern. The transformed variables are named v_1 to v_{27} . For example, from uncorrelated variables x_1 and x_2 we can generate new corresponding correlated variables $v_1 = x_1$ and $v_2 = \sqrt{x_1 * x_2}$, respectively. From these 27 variables, two (v_1, v_5), were chosen to be the model variables, that is, the variables governing the response behaviour, and another one (v_{27}), was chosen to be the study variable y . The correlations between each model variable with the study variable are approximately 0.5. The remaining 24 were assumed to be auxiliary variables. These data correspond to our population of 3 260 observations. The simulation process was the following:

1. From this population we draw a sample of size 300 using simple random sampling without replacement.
2. A response set was generated using a logistic regression model $p_k = 1/(1 + \exp(-\delta' \mathbf{H}_k))$, where $\mathbf{H} = \{1, v_1, v_5\}^t$ is a vector of model variables, whereas δ is a vector of model parameters. The elements $k \in S$ for which a Bernoulli trial was 1 with probability p_k , were selected to the response set.
3. Estimates of interest were calculated using the data in the response set.
4. The process was repeated 1 000 times. Higher replication numbers basically produced similar results.

5. Indicators of the properties of the estimators were calculated. These indicators are the relative bias ($Rel.bias = \frac{bias(\hat{\theta})}{\theta} * 100\%$), the standard error ($S.E. = \sqrt{var(\hat{\theta})}$), and the root mean squared error ($RMSE = \sqrt{bias(\hat{\theta})^2 + var(\hat{\theta})}$), where $bias(\hat{\theta}) = mean(\hat{\theta}) - \theta$, $mean(\hat{\theta}) = \frac{\sum_{i=1}^{1000} \hat{\theta}_i}{1000}$, and $var(\hat{\theta}) = \frac{1}{999} \sum_{i=1}^{1000} (\hat{\theta}_i - mean(\hat{\theta}))^2$.

The points 1 to 5 were repeated for samples of sizes 400, 500, and 600. We chose $\delta = \{1.311, -0.199, -0.083\}^t$, which led to an average response rate of 57% for each sample size.

Recall that we base this article on two calibration approaches, the linear calibration (LC) estimator of Särndal and Lundström (2005) and the propensity score calibration (PSC) of Chang and Kott (2008). For the former estimator, the auxiliary vector was given by $\mathbf{X}_k = \{1, v_{1k}, \dots, v_{26k}\}^t$, whereas the latter used $\mathbf{X}_k = \{1, v_{2k}, \dots, v_{4k}, v_{6k}, \dots, v_{26k}\}^t$ and $\mathbf{H}_k = \{1, v_{1k}, v_{5k}\}^t$.

The principal components auxiliary variables for both the LC and PSC estimators were generated from their corresponding values of \mathbf{X} . The retention criterion for the LC estimator was the proportion of total variance explained by the set of selected components. This led to the selection of three principal components in population LC, while for the PSC estimator, the retention criterion is the one suggested in Section 4.3. The scree plot given in Figure 1 below illustrates the population correlation structure of the variables.

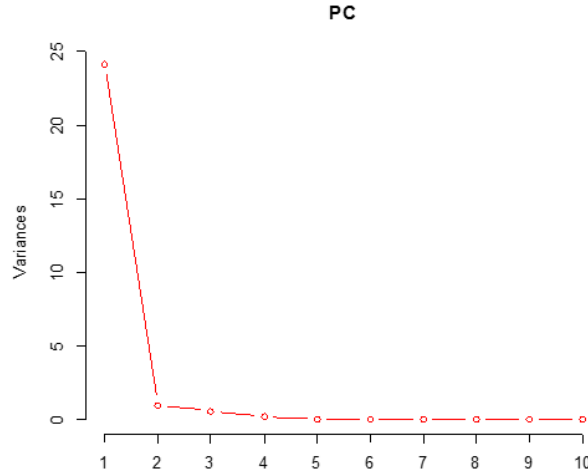


Figure 1: Scree plot of the auxiliary data in the first study.

5.1.2. Study 2

In this study we perform the simulations following the same 5 points of study 1. However, here we use the ‘Small Area Income and Poverty Estimates’ dataset (size 3 173). We selected some variables from the 2006 and others from the 2013 data, for a total of 19 variables. The original data

were square root transformed and the new variables were named v_1 to v_{19} . Again we chose three variables, two of which were used as model variables, say, v_1 and v_5 , and the other was the study variable, say, v_{19} . The correlations of v_1 and v_5 with the study variable v_{19} were $cor(v_1, v_{19}) = 0.52$ and $cor(v_5, v_{19}) = 0.45$, respectively. This resembles the correlation structure of the corresponding variables in study 1. As in study 1, we used $\delta = \{1.311, -0.199, -0.083\}^t$. The LC estimator uses $\mathbf{X}_k = \{1, v_{1k}, \dots, v_{18k}\}^t$, whereas the PSC estimator uses $\mathbf{X}_k = \{1, v_{2k}, \dots, v_{4k}, v_{6k}, \dots, v_{18k}\}^t$ and $\mathbf{H}_k = \{1, v_{1k}, v_{5k}\}^t$. The proportion of total variance explained by the selected PCs is again the retention criterion used for the LC estimator based on PCs. This criterion led to a selection of eight components. The retention criterion for the PSC estimator based on PCs is again the one described in Section 4.3. The following is the scree plot of the principal components of the population auxiliary data used in the second simulation study.

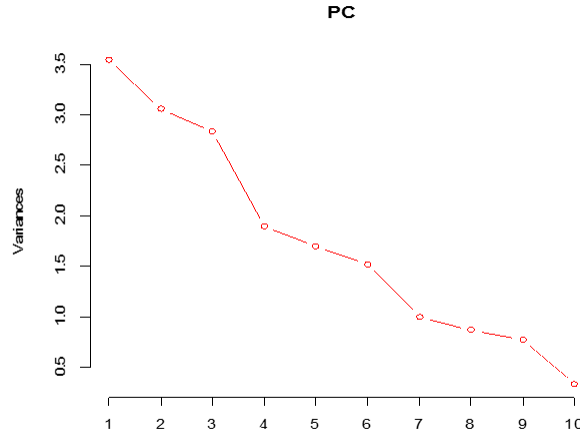


Figure 2: Scree plot of the auxiliary data in the second study.

5.2. Simulation results

5.2.1. Results of Study 1

The results are presented in two versions, a tabular version in Tables 1–4 and a graphic version in Figures 3–6 (the figures are in Appendix B). These representations show the behaviour of each considered estimator when the sample size increases. For each table or graph the performance of the estimator is evaluated from two perspectives, i.e., when the estimator is based on the complete original auxiliary variables (X) and when it is based on the PCs of the auxiliary variables.

Tables 1 and 2 show the LC estimator results when auxiliary information is observed at the population and sample levels, respectively. Both tables show that the relative bias, standard error, and the root mean squared error values of the PCs-based linear calibration are smaller than their

counterparts computed based on the original auxiliary variables.

Table 1: LC on original population auxiliary variables vs. LC on population PCs – Study 1.

Sample size	Properties	Estimators	
		L. Calibration on X	L. Calibration on PCs
300	Rel.bias(%)	5.474	1.296
	S.E.	3 519	935
	RMSE	8 661	2 094
400	Rel.bias(%)	4.771	1.224
	S.E.	3 231	872
	RMSE	7 616	1 973
500	Rel.bias(%)	4.462	1.222
	S.E.	3 083	804
	RMSE	7 150	1 941
600	Rel.Bias(%)	3.974	1.149
	S.E.	3 135	846
	RMSE	6 544	1 864

Table 2: LC on original sample auxiliary variables vs. LC on sample PCs – Study 1.

Sample size	Properties	Estimator	
		L. Calibration on X	L. Calibration on PCs
300	Rel.bias(%)	3.930	0.192
	S.E.	21 936	11 202
	RMSE	22 660	11 206
400	Rel.bias(%)	3.341	0.037
	S.E.	17 089	9 621
	RMSE	17 758	9 621
500	Rel.bias(%)	3.551	0.328
	S.E.	14 332	8 250
	RMSE	15 224	8 263
600	Rel.bias(%)	2.951	0.369
	S.E.	12 422	7 608
	RMSE	13 134	7 626

Tables 3 and 4 show results obtained under conditions similar to those used to obtain the results in Tables 1 and 2, except that PSC replaces LC. The results obtained by using PCs of the auxiliary variables are comparable to those obtained using original auxiliary variables; this is true in both levels of auxiliary information.

Table 3: PSC on original population auxiliary variables vs. PSC on population PCs – Study 1.

Sample size	Properties	Estimator			
		PS on X	Time (in hr)	PS on PCs	Time (in hr)
300	Rel.bias(%)	0.280		0.153	
	S.E.	16 182	7	15 912	0.35
	RMSE	16 188		15 914	
400	Rel.bias(%)	0.105		0.209	
	S.E.	13 815	13	13 660	0.57
	RMSE	13 816		13 663	
500	Rel.bias(%)	0.338		0.434	
	S.E.	11 953	22	11 837	0.83
	RMSE	11 963		11 854	
600	Rel.bias(%)	0.169		0.264	
	S.E.	10 899	36	10 757	1.30
	RMSE	10 902		10 764	

Table 4: PSC on original sample auxiliary variables vs. PSC on sample PCs – Study 1.

Sample size	Properties	Estimator			
		PS on X	Time (in hr)	PS on PCs	Time (in hr)
300	Rel.bias(%)	0.255		0.125	
	S.E.	16 162	0.25	16 010	0.18
	RMSE	16 166		16 011	
400	Rel.bias(%)	0.120		0.189	
	S.E.	13 820	0.32	13 711	0.22
	RMSE	13 821		13 713	
500	Rel.bias(%)	0.353		0.421	
	S.E.	11 952	0.45	11 834	0.23
	RMSE	11 963		11 850	
600	Rel.bias(%)	0.191		0.263	
	S.E.	10 880	0.50	10 795	0.25
	RMSE	10 884		10 801	

5.2.2. Results of Study 2

Tables 5–10 below present the results of this study. The process of evaluating the estimators is similar to that used in Study 1. The results of the LC, presented in Tables 5 and 6, display consistency when comparing X-and PCs- based estimators and when comparing population- and sample-based estimators.

The results of the PSC estimators for the second study are displayed in Tables 7 and 8. As with the LC estimator, the PSC results are also consistent in terms of the type (X or PCs) and level (population or sample) of the auxiliary information used.

Table 5: LC on original population auxiliary variables vs LC on population PCs – Study 2.

Sample size	Properties	Estimator	
		L. Calibration on X	L. Calibration on PCs
300	Rel.bias(%)	0.735	0.899
	S.E.	2 262	2 136
	MSE	2 282	2 168
400	Rel.bias(%)	0.810	1.077
	S.E.	1 871	1 798
	MSE	1 901	1 852
500	Rel.bias(%)	0.829	1.029
	S.E.	1 558	1 529
	MSE	1 596	1 588
600	Rel.bias(%)	0.672	0.836
	S.E.	1 402	1 382
	MSE	1 429	1 425

Table 6: LC on original sample auxiliary variables vs. LC on sample PCs – Study 2.

Sample size	Properties	Estimator	
		L. Calibration on X	L. Calibration on PCs
300	Rel.bias(%)	0.725	0.949
	S.E.	2 489	2 494
	MSE	2 507	2 525
400	Rel.bias(%)	0.882	1.205
	S.E.	2 068	2 068
	MSE	2 100	2 129
500	Rel.bias(%)	0.841	1.104
	S.E.	1 792	1 814
	MSE	1 825	1 871
600	Rel.bias(%)	0.711	0.937
	S.E.	1 639	1 658
	MSE	1 666	1 703

Table 7: PSC on original population auxiliary variables vs. PSC on population PCs – Study 2.

Sample size	Properties	Estimator	
		PSC on X	PSC on PCs
300	Rel.bias(%)	1.345	1.566
	S.E.	3 748	3 791
	MSE	3 789	3 846
400	Rel.bias(%)	1.627	1.925
	S.E.	3 293	3 385
	MSE	3 362	3 478
500	Rel.bias(%)	1.487	1.848
	S.E.	2 921	2 994
	MSE	2 985	3 091
600	Rel.bias(%)	1.757	2.072
	S.E.	2 708	2 846
	MSE	2 804	2 973

Table 8: PSC on original sample auxiliary variables vs. PSC on sample PCs – Study 2.

Sample size	Properties	Estimator	
		PSC on X	PSC on PCs
300	Rel.bias(%)	1.347	1.480
	S.E.	3 634	3 643
	MSE	3 676	3 695
400	Rel.bias(%)	1.482	1.701
	S.E.	3 166	3 219
	MSE	3 226	3 296
500	Rel.bias(%)	1.559	1.648
	S.E.	2 775	2 815
	MSE	2 849	2 897
600	Rel.bias(%)	1.778	1.908
	S.E.	2 567	2 651
	MSE	2 672	2 767

The results shown in Tables 9 and 10 comprise estimated model parameters (with associated standard errors in parentheses) in the PSC estimation using the data of the second study.

Table 9: Estimated model coefficients (population auxiliary information – Study 2).

Coefficient estimates						
$(\delta_0, \delta_1, \delta_2)$						
True coefficients (1.311, -0.199, -0.083)						
Sample size	PSC on X			PSC on PCs		
	$\hat{\delta}_0$	$\hat{\delta}_1$	$\hat{\delta}_2$	$\hat{\delta}_0$	$\hat{\delta}_1$	$\hat{\delta}_2$
300	1.129	-0.182	-0.044	1.097	-0.188	-0.026
	(0.197)	(0.008)	(0.011)	(0.247)	(0.012)	(0.014)
400	1.125	-0.174	-0.052	1.079	-0.176	-0.035
	0.149	0.006	0.008	(0.213)	(0.010)	(0.010)
500	1.147	-0.178	-0.056	1.096	-0.179	-0.038
	(0.133)	(0.005)	(0.006)	(0.182)	(0.008)	(0.009)
600	1.140	-0.178	-0.054	1.092	-0.179	-0.037
	(0.117)	(0.005)	(0.005)	(0.190)	(0.007)	(0.008)

Table 10: Estimated model coefficients (sample auxiliary information – Study 2).

Coefficient estimates						
$(\delta_0, \delta_1, \delta_2)$						
True coefficients (1.311, -0.199, -0.083)						
Sample size	PSC on X			PSC on PCs		
	$\hat{\delta}_0$	$\hat{\delta}_1$	$\hat{\delta}_2$	$\hat{\delta}_0$	$\hat{\delta}_1$	$\hat{\delta}_2$
300	1.139	-0.177	-0.054	1.119	-0.179	-0.046
	(0.092)	(0.003)	(0.005)	(0.122)	(0.005)	(0.005)
400	1.149	-0.175	-0.061	1.118	-0.174	-0.052
	(0.068)	(0.003)	(0.003)	(0.108)	(0.004)	(0.005)
500	1.148	-0.175	-0.061	1.132	-0.175	-0.054
	(0.063)	(0.0002)	(0.003)	(0.003)	(0.094)	(0.004)
600	1.145	-0.175	-0.060	1.123	-0.174	-0.054
	(0.06)	(0.002)	(0.002)	(0.099)	(0.003)	(0.004)

6. Discussion

The results of two simulation studies are presented in the previous section and for each study we assess two calibration approaches, namely, the LC estimator using no explicit form of response function and the PSC estimator with explicit functional form. Both estimators are evaluated using the original large set of auxiliary variables (X) and using the principal components (PCs) of the original auxiliary variables. The results of the first study are given in two versions, tabular and graphic, while the results of the second study are given in tabular form only. The graphic form enables the convenient visual inspection of the estimator behaviour, while the tabular form gives a quantitative illustration. Study 1 demonstrates that the LC estimator based on principal components auxiliary variables is always superior in terms of relative bias, standard error, and root mean squared error (RMSE), to its counterpart using the original auxiliary information. This is true regardless of the level of the auxiliary information, that is, the population or sample levels, as demonstrated in Tables 1 and 2, respectively.

There is a large discrepancy between the standard errors and RMSEs between population- and sample-based LC estimators. The RMSE of the population-based LC estimator based on the original auxiliary information ranges from 6 544 to 8 661 while the range for its counterpart based on sample-level auxiliary information is 13 134 to 22 660. The RMSE of the LC based on PCs auxiliary information ranges from 1 864 to 2 094 and from 7 626 to 11 206 for population- and sample-based auxiliary information, respectively. Thus, the results differ greatly when comparing estimators of population- and sample-based auxiliary information. Considerable differences are also observed in the standard errors and RMSEs when comparing the estimators in terms of the type of auxiliary information used, that is, original X- auxiliaries and PCs- auxiliaries. This is not a surprising behaviour of the LC calibration estimator as this is a regression-type estimator.

In the response propensity calibration approach, auxiliary information is used in estimating response propensities; the estimation of population characteristics then proceeds by adjusting the design weights through multiplication by the corresponding reciprocals of the estimated propensities, which is usually called ‘double weighting’. Here it is observed that these results are more consistent. As Tables 3 and 4 illustrate, the principal-components-based estimator provides results similar to those obtained using the original auxiliary information. This is true regardless of the level of information on which the estimator is based. Furthermore, the results display consistency when comparing the properties of the corresponding estimators when population- and sample-based auxiliary data are used. The corresponding interval ranges of the RMSEs when using population-level auxiliary information are close to those when sample-level auxiliary information is used. As the sample size increases, the RMSEs tend to converge to the same level, irrespective of the type (X or PCs) or level (population or sample) of the auxiliary information. One of the major advantages of using PCs in place of the original auxiliary variables is the computational effort measured in terms of computational time; as reported in Tables 3–4, due to dimensionality reduction, the principal-components-based estimates are computed much more quickly than are the estimates based on the original auxiliary information.

Tables 5–10 report the results of the second simulation study. In contrast to the previous study, here the LC calibration (Tables 5–6) results are consistent regardless of the type of auxiliary information used for estimation as well as when comparing the properties of the estimator across levels

of information. The RMSEs of the estimators lie in virtually the same interval, regardless of the level of auxiliary information (population or sample levels) or type (original X or PCs). A similar observation can be made with respect to the PSC estimator in Tables 7–8. We can still compare the performances of the LC and PSC estimators as we are using the same set of auxiliary variables, however, the estimators are conceptually different in terms of how auxiliary information is used.

The levels of bias are approximately the same, that is, they are less than 0.1% in Study 2 while in Study 1 some differences are observed, especially in the LC estimator where the bias level attains 5.5%, as Tables 1 and 2 demonstrate. An interesting property of the auxiliary information in the PSC scheme, is the ability to appropriately estimate the response model. Tables 9–10 provide the population- and sample-based model-estimated coefficients, and the results suggest equally good model coefficients estimates when PCs are used compared with estimates resulting from the use of the original X variables. As the results of model estimates are good, we can further improve the target estimates by performing a two-step estimation in which the products of design weights and the reciprocal of the estimated response probabilities are used as initial weights in the linear calibration estimator.

Both Study 1 and Study 2 illustrate how the use of principal components in place of original auxiliary data when adjusting for nonresponse does not lead to distorted results and has the great advantage of reducing the computational effort.

The reported PSC results based on principal components are very similar to those obtained using a fixed number of components via the eigenvalue-one rule. However, the eigenvalue-one results are worse than those of our approach based on canonical correlation for very small samples. When the sample size increases, the number of selected components converges to the number of components based on the eigenvalue-one rule. Figure 7 in Appendix B illustrates the behaviour of our components selection method using the data of Study 1.

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Appendix A

Asymptotic variance of the estimated coefficients of the propensity functions

Let

$$E_{pq} \left[\left(\mathbf{Z}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \mathbf{T}_z \right) \left(\mathbf{Z}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \mathbf{T}_z \right)^t \right] = \Pi_1 + \Pi_2,$$

where $\Pi_1 = \sum_{k \in U} \sum_{l \in U} (\pi_{kl} - \pi_k \pi_l) d_k d_l \mathbf{Z}_k \mathbf{Z}_l^t$ and $\Pi_2 = \sum_U d_k (h(\mathbf{H}_k^t \delta^*) - 1) \mathbf{Z}_k \mathbf{Z}_k^t$ (see Chang and Kott, 2008).

Then

$$Avar \sqrt{n} (\hat{\delta}_{(pc)} - \delta^*) = [\mathbf{F}^t \mathbf{W} \mathbf{F}]^{-1} \mathbf{F}^t \mathbf{W} \Theta \mathbf{W} \mathbf{F} [\mathbf{F}^t \mathbf{W} \mathbf{F}]^{-1},$$

with $\mathbf{F} = \mathbf{Z}^t \Psi \mathbf{H}$ and $\Theta = Avar \left[n^{-1/2} \left(\mathbf{Z}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \mathbf{T}_z \right) \right]$.

We choose

$$\mathbf{W}^{-1} = \Theta$$

and obtain

$$Avar \sqrt{n} (\hat{\delta}_{(pc)} - \delta^*) = \left[(\mathbf{Z}^t \Psi \mathbf{H})^t \Theta^{-1} (\mathbf{Z}^t \Psi \mathbf{H}) \right]^{-1}, \quad (15)$$

where $\mathbf{W} = \text{plim}_{n \rightarrow \infty} \mathbf{W}_n$ is a positive definite matrix, $(\mathbf{Z}^t \Psi \mathbf{H}) = \text{plim}_{n \rightarrow \infty} \frac{1}{n} E_{pq} \left(\mathbf{Z}_{(r)}^t \Psi^0 \mathbf{H} \right)$ and

$$\Theta = \text{plim}_{n \rightarrow \infty} \frac{1}{n} E_{pq} \left[\left(\mathbf{Z}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \mathbf{T}_z \right) \left(\mathbf{Z}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \mathbf{T}_z \right)^t \right].$$

Alternatively, the calibration (4) is on estimated principal components, that is,

$$\hat{\mathbf{Z}}_{(r)}^t \Phi(\delta) \mathbf{d}_{(r)} - \hat{\mathbf{T}}_z = \mathbf{0},$$

where $\hat{\mathbf{T}} = \sum_s d_k \hat{\mathbf{Z}}_k$.

Observe that

$$var_{pq}(\hat{\mathbf{Z}}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \hat{\mathbf{T}}_z) = V_1 + V_2,$$

where $V_1 = var_p E_q \left(\hat{\mathbf{Z}}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \hat{\mathbf{T}}_z | s \right)$ and $V_2 = E_p var_q \left(\hat{\mathbf{Z}}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \hat{\mathbf{T}}_z | s \right)$. The first variance component is zero, implying that

$$E_{pq} \left[\left(\hat{\mathbf{Z}}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \hat{\mathbf{T}}_z \right) \left(\hat{\mathbf{Z}}_{(r)}^t \Phi(\delta^*) \mathbf{d}_{(r)} - \hat{\mathbf{T}}_z \right)^t \right] = \sum_U d_k (h(\mathbf{H}_k^t \delta^*) - 1) \mathbf{Z}_k \mathbf{Z}_k^t,$$

therefore the sample version analogous to \mathbf{W} in (15) is $\tilde{\mathbf{W}} = \text{plim}_{n \rightarrow \infty} \frac{1}{n} \sum_U d_k (h(\mathbf{H}_k^t \delta^*) - 1) \mathbf{Z}_k \mathbf{Z}_k^t$.

Appendix B

Figures

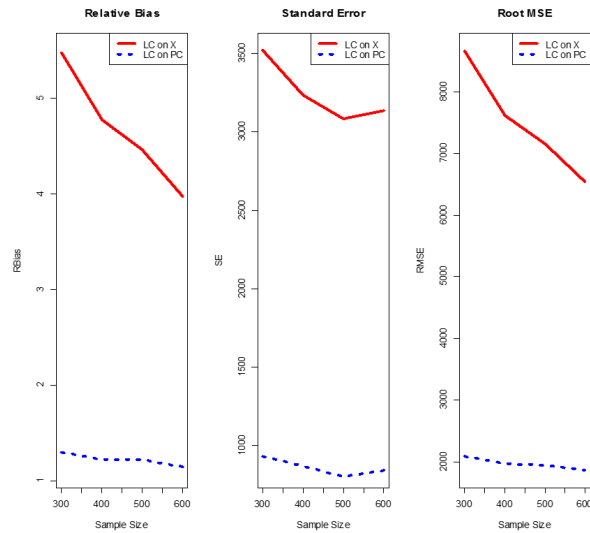


Figure 3: LC on original population auxiliary variables vs. LC on population PCs – Study 1.

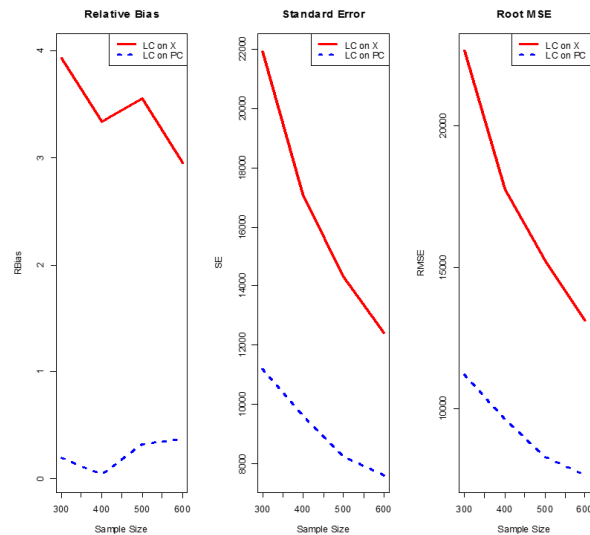


Figure 4: LC on original sample auxiliary variables vs. LC on sample PCs – Study 1.

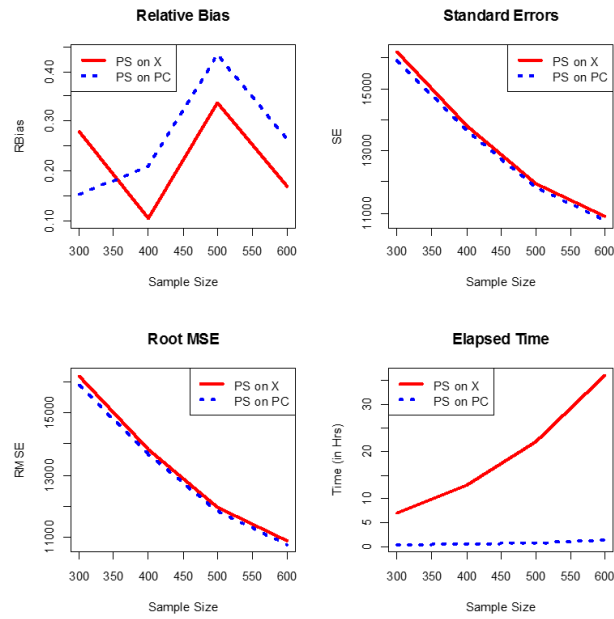


Figure 5: PSC on original population auxiliary variables vs. PSC on population PCs – Study 1.

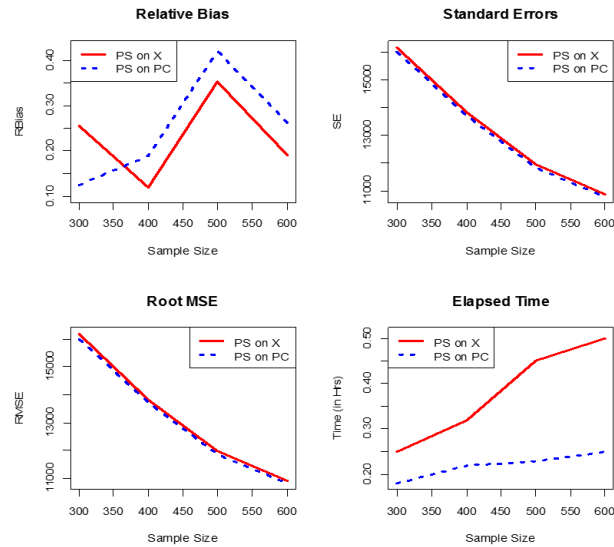


Figure 6: PSC on original sample auxiliary variables vs. PSC on sample PCs – Study 1.

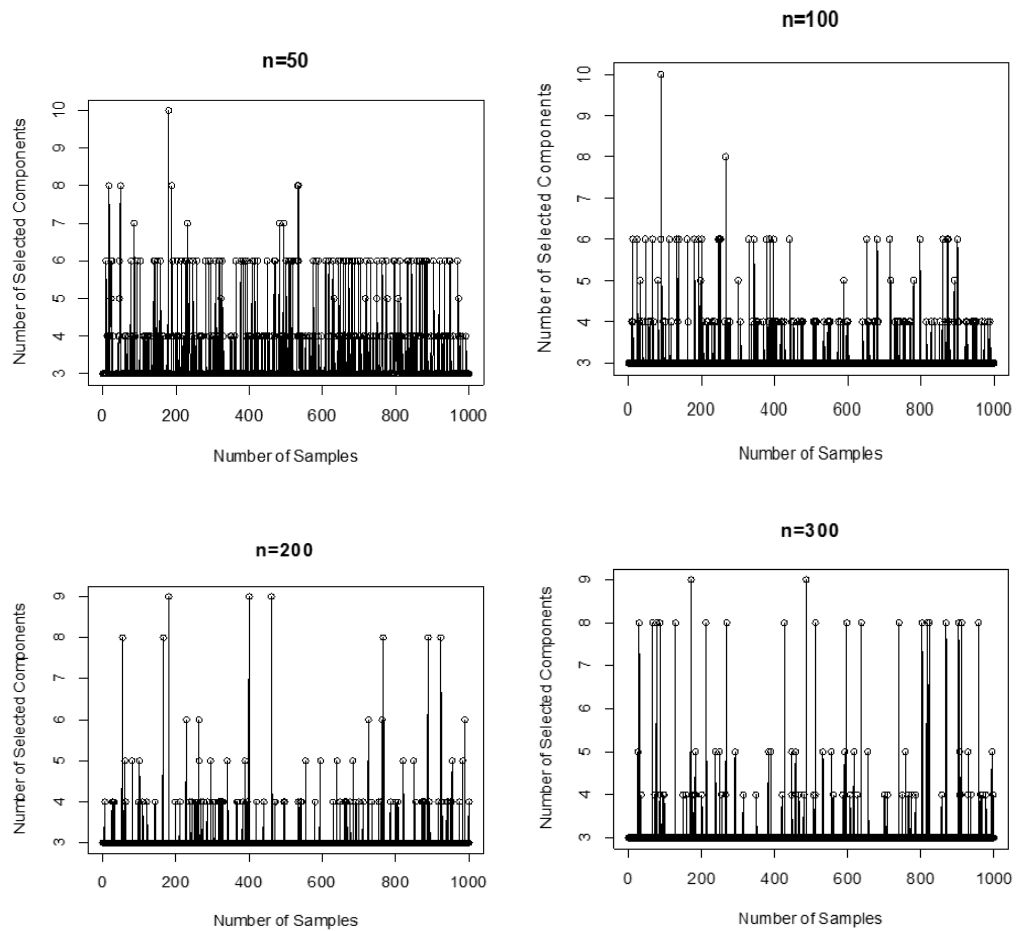


Figure 7: Behaviour of the number of selected components when sample sizes increase – Study 1.

