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Variance estimation methods in samples from household surveys

ABSTRACT

OBJECTIVE: Knowledge of sampling errors is essential for correctly interpreting the results from household surveys and evaluating their sampling designs. The composition of household samples used in surveys gives rise to situations of complex estimation. In this light, the study was conducted with the aim of evaluating the performance of the variance estimators in surveys carried out among urban populations in Brazil.

METHODS: The reference population was the sample drawn by the Fundação Sistema Estadual de Análise de Dados Estatísticos (SEADE – State Statistical Data Analysis System Foundation) for carrying out an employment and unemployment survey in the metropolitan region of São Paulo. Three techniques were used for estimating variance: Taylor linearization and Jackknife and BRR replication. Repeated samples were selected from the reference population, using stratified cluster sampling in two stages (census tracts and households). Three different designs were used and 2,000 samples were drawn within each design. To obtain an estimator ratio, the accuracy of the variance estimators was evaluated by means of the mean square error and the confidence interval coverage.

RESULTS: According to the mean square error, the three techniques provided similar accuracy. The bias ratios were approximately 0.10, for the smaller samples. The confidence interval coverage indicated that the confidence levels observed were lower than what was set (95%), and were around 90% for the smaller samples.

CONCLUSIONS: The variance estimators showed similar performance with regard to accuracy and confidence interval coverage. The bias was irrelevant in relation to the magnitude of the standard error. Although the real confidence levels were lower than the nominal levels for normal distribution, the changes did not prevent construction of interval estimates with reasonable confidence.

KEY WORDS: Estimation techniques. Data collection. Stratified sampling. Cluster sampling. Taylor series linearization. Complex sample design.

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INTRODUCTION

Knowledge of sampling errors is essential for correctly interpreting the results from household surveys and evaluating their sampling designs. However, the composition of household samples with multi-stage cluster draws makes estimation complex. In such cases, the ordinary approach of using variance estimation methods based on simple random sampling (SRS) can be regarded as inappropriate.

In SRS with replacement in which the units have equal probabilities of being selected and are independent, the estimated variance for the mean of a variable

"y" is expressed by
$$\frac{s^2}{n} = \frac{\sum_{i=1}^{n} (y_i - \overline{y})^2}{n(n-1)}$$
, where n is the number of elements

in the sample, y_i is the value of these elements and

$$\overline{y} = \frac{\sum_{i=1}^{n} y_i}{n}$$
 is its mean.⁸

In household surveys, because the population elements are scattered over wide geographic areas, it becomes necessary to use cluster sampling in order to reduce the cost of data collection. By using clusters with differing sizes and self-weighted sampling procedures, the number of sample members becomes a random variable, which transforms the mean into a ratio of variables, with consequences for estimating the sampling error. In this context, the majority of first-order estimators are nonlinear, and because there are no exact expressions for calculating their variances, it becomes necessary to use approximations adjusted to the complex nature of the sampling design and the estimation procedure.

Nowadays, the most frequently used methods for estimating variance in complex sampling designs are Taylor-series linearization and replication. The first provides a linear approximation for the nonlinear estimator of interest, by means of the Taylor series, to which the usual linear estimator formulas are applied. In order to obtain an adequate approximation for the variance of the ratio estimator by means of a Taylor series, the sample size (the denominator of the ratio) should not be subject to great variation, which would however be the case when the clusters have widely different sizes. The difficulty in keeping the denominator variability under control increases if the estimation is directed towards subclasses, since it is then impossible to control the number of units belonging to each subclass. This is especially true for phenomena that are rare and/or unevenly distributed throughout the clusters.

For the ratio estimator, $r = \frac{y}{x}$, the variance estimator expression obtained by using the Taylor linearization method has been widely used in sampling, and it is as follows: $var(r) = \frac{1}{x^2} [var(y) + r^2 var(x) - 2r cov(xy)].^8$ In spite of being frequently presented in the literature on statistics, this expression cannot be regarded as simple from a computational standpoint, since it requires the numerator variance, var(y), the denominator variance, var(x) and their covariance, cov(x,y).

Replication techniques were developed as a simplifying alternative approach towards the variance estimation procedure. They consist of obtaining subsets from the full sample (replicate samples), repeating the estimation procedure for each subsample and calculating the variance from these estimates. Therefore if θ is the parameter of interest and its estimator, K replicates are formed and estimates are $\hat{\theta}$ obtained for each replicate

using $\hat{\theta}_k$ (k = 1,...,K). The variance estimator for $\hat{\theta}$ is given by: $var(\hat{\theta}) = \frac{1}{c} \sum_{k=1}^{K} (\hat{\theta}_k - \hat{\theta})^2$, where c is a constant

associated with the replication method adopted.17

This construction follows the simplicity of variance estimation found in SRS, which uses the deviations of each observation from the mean. The basic concept was developed by Mahalanobis in studies in 1944 and 1946, in which the use of replicate samples called "interpenetrating samples" was proposed to make it easier to estimate sampling errors and to investigate non-sampling errors such as bias related to interviewers.

The most common replication techniques are "Balanced Repeated Replication" (BRR) and "Jackknife". BRR was developed as an estimation alternative for designs that, for efficiency reasons, use a large number of strata and consequently the least possible number of primary sampling units for each stratum, i.e. two units. It originated from the pseudo-replication scheme known as "half-sample replication", which was proposed by the United States Bureau of the Census and later adapted and modified by experts from the National Center of Health Statistics. In 1966, McCarthy¹² introduced the balancing referred to in the name of the technique. The replicates are composed of one sampling units from each stratum.

In the Jackknife technique, each replicate is obtained by successively omitting one sampling unit from each stratum. Jackknife estimation procedures were originally developed by Quenouille, in 1949, with the purpose of reducing the bias of the correlation coefficient estimator in time series. Tukey15 suggested that the individual estimators from the subsamples created by this technique could be seen as independent and identically distributed random variables, thereby yielding a very simple variance estimator. He named it Jackknife in reference to the multipurpose tool of the same name. Its use in the context of finite populations seems to have been considered for the first time by Durbin, in 1959, in connection with the ratio estimator. Is

Besides providing simpler variance expressions than those obtained by the Taylor linearization method, one attractive feature of replication techniques is that, for a given design, the same analysis procedure is used for almost all statistics, regardless of their complexity. Another property of replication techniques is that they enable users of secondary survey data to estimate sampling errors without knowing the detailed sample design. The replicates created by the investigators involved in the survey can simply be used and included in the data file. This is especially useful when there are confidentiality issues involving the sample units and it is necessary to avoid dissemination of any information identifying the sample units. 14.*

^{*} Brick JM, Morganstein D, Valliant R. Analysis of complex samples using replication. Rockville: Westat; 1998 [Accessed on month day, year]. Available at: http://www.westat.com/wesvar/techpapers/ACS-Replication.pdf

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Present-day electronic processing capabilities make it possible to apply any of these techniques for estimating sampling errors in surveys designed with cluster sampling. The comparative studies available are based on empirical results obtained in various countries under different designs. 1,3,4,9,10,13

The present study had the objective of showing the performance of the Jackknife, BRR and Taylor linearization techniques for variance estimation, in samples that reflect the sociodemographic structure of the metropolitan region of São Paulo. With this aim, census tract samples (which form the geographic units most used in household surveys in Brazil) were used to compare the accuracy of variance estimators. Through this, the intention was to contribute towards the knowledge and dissemination of the alternatives for sampling error estimation that exist, and thus to stimulate the use of adequate techniques for making statistical inferences from household surveys.

METHODS

The reference population for developing the study was a sample drawn by the *Fundação Sistema Estadual de Análise de Dados Estatísticos* (SEADE - State Statistical Data Analysis System Foundation) for carrying out an employment and unemployment survey in the metropolitan region of São Paulo.⁶

Repeated samples were taken from this population, using stratified cluster sampling in two stages. The census tracts constituted the primary sampling units (PSUs) and the households were the secondary units. From each stratum, two census tracts were drawn with probability proportional to the number of households. From each of these census tracts, five households were drawn, thus totaling ten households per stratum. The sampling fraction for stratum h was given by:

$$f_h = \frac{2 \cdot M_{h\alpha}}{M_h} \cdot \frac{5}{M_{h\alpha}} = \frac{10}{M_h} \text{ , where } M_h \text{ is the number}$$

of households in stratum h and $M_{h\alpha}$ is the number of households in census tract α of stratum h.

In order to evaluate the accuracy of the variance estimators in relation to increases in the number of primary sampling units, three designs were defined, without changing the general model presented previously. For the first, second and third designs, the population was organized into 8, 16 and 32 strata respectively. After drawing two census tracts per stratum, the numbers of tracts included in the samples were 16, 32 and 64. Also taking into account that five households were drawn in each tract, the final sample sizes in the three designs were 80, 160 and 240 households.

A total of 2,000 total of samples were drawn under each design.

The estimated mean family income was for each sample,

by using the following expression:
$$r = \frac{\sum_{h} w_{h} \sum_{\alpha} \sum_{\beta} Y_{h\alpha\beta}}{\sum_{h} w_{h} \sum_{\alpha} \sum_{\beta} x_{h\alpha\beta}},$$

where $y_{ha\beta}$ is the family income of the household β , belonging to census tract α , in stratum h; $x_{ha\beta}=I$, for households with income and $x_{ha\beta}=0$ for those with no response; $w_h=1/f_h$, the weight of each household, is given by the inverse of the sampling fraction of the stratum that the household belongs to, $f_h=10/\mathrm{M_b}$.

From the frequency distribution, the population variance

was calculated for each design:
$$S^2 = \frac{\sum_{i=1}^{2000} [r_i - E(r)]^2}{2000}$$

where E(r) is the expected value of r corresponding to

the mean of its sampling distribution,
$$E(r) = \frac{\sum_{i=1}^{2000} r_i}{2000}$$
, and

 r_i is the mean income calculated for the i-th sample. Variance estimates using the Jackknife and BRR techniques were calculated using the WesVar software, version $4.0,^{17}$ according to the expression:

$$var(r) = \frac{\sum_{g=1}^{G} (r_{(g)} - r)^{2}}{C}$$
, where *r* is an estimate based

on all the primary units; $r_{(g)}$ is an estimate based on the g-th replicate; G is the number of replicates and c is a constant that depends on the replication technique, (c=G for BRR; c=1 for Jackknife).

Variance estimates using the Taylor linearization method were obtained as follows:⁸

$$\operatorname{var}(r) = \frac{1}{x^2} \left[\sum_{h} \operatorname{var}(y_h) + r^2 \sum_{h} \operatorname{var}(x_h) - 2r \sum_{h} \operatorname{cov}(y_h, x_h) \right]$$

$$\operatorname{var}(y_h) = \frac{1}{a_h - 1} \left[a_h \sum_{\alpha} (w_{h\alpha} y_{h\alpha})^2 - \left(\sum_{\alpha} w_{h\alpha} y_{h\alpha} \right)^2 \right]$$

$$\operatorname{var}(x_h) = \frac{1}{a_h - 1} \left[a_h \sum_{\alpha} (w_{h\alpha} x_{h\alpha})^2 - \left(\sum_{\alpha} w_{h\alpha} x_{h\alpha} \right)^2 \right]$$

$$\operatorname{cov}(y_h, x_h) = \frac{1}{a_h - 1} \left[a_h \sum_{\alpha} (w_{h\alpha} y_{h\alpha}) (w_{h\alpha} x_{h\alpha}) - \left(\sum_{\alpha} w_{h\alpha} y_{h\alpha} \right) \left(\sum_{\alpha} w_{h\alpha} x_{h\alpha} \right) \right]$$

where $a_h = 2$, i.e. the number of primary sampling units; w_{ha} is the weight of household α in stratum h; y_{ha} is the value of the study variable in household α in stratum h; $x_{ha}=1$ if there is information on the household and $x_{ha}=0$ if not.

Considering that the accuracy refers to small total errors, including bias and sampling variability, the accuracy of the estimators was evaluated by means of the mean square error: $MSE[var(r)] = Var[var(r) + Bias^2[var(r)]]$ and by means of its relative measure: $RMSE = \frac{MSE[var(r)]}{E[var(r)]^2}$.

The sampling distribution variance was expressed by:

$$Var[var(r)] = \frac{\sum_{i=1}^{2000} \left\{ var(r_i) - E[var(r)] \right\}^2}{2000}, \text{ and } E[var(r)],$$

i.e. the expected value for the variance of r, was calculated as follows:
$$E[var(r)] = \frac{\sum_{i=1}^{2000} var(r_i)}{2000}$$
.

The bias of the estimators, measured as the distance between the means of the sampling distribution and of the population, was expressed by: $Bias[var(r)] = E[var(r)] - S^2$, for which the terms were presented earlier. The effect of the bias on the accuracy of the estimators was evaluated by the bias ratio, which measures the bias in standard deviation units:

$$BR[var(r)] = \frac{Bias[var(r)]}{\sqrt{Var[var(r)]}}.$$

$$\sqrt{\text{Var}[\text{Var}(r)]}$$
The distribution of the standardized ratio $\frac{r - E(r)}{\sqrt{\text{Var}(r)}}$ was

studied with a view to checking the real confidence level of the constructed intervals. Using Student's t distribution values for 8, 16 and 24 degrees of freedom, the following intervals were respectively constructed: [-2.306;+2.306], [-2.120;+2.120] and [-2.064;+2.064]. Next, the proportion of times that the standardized ratio was within the above intervals was calculated and checked for its closeness to 0.95. Because the exact distribution of the standardized ratio is unknown for complex estimators, calculation of the coverage and checking its pertinence to predetermined intervals makes it possible to evaluate the applicability of the confidence intervals.1

RESULTS

The sampling distributions for the estimator r were constructed for each of the three designs that were defined, assuming that the shape of the distribution of 2,000 estimates was stable enough to be regarded as a sampling distribution. The variance of these distributions was taken to be the real variance of r for the fixed design, as done by Bean and Kish & Frankel in their studies.1,9

Likewise, the sampling distributions of the variance estimators for the BRR, Jackknife and Taylor techniques were constructed and are shown in Figures 1 to 3. Each figure refers to one of the sampling designs. The means and standard errors of the distributions are indicated on each figure.

It can be noted by overlaying the curves that the results regarding the precision and bias of the estimators were very similar. These results are shown in Table 1.

According to the relative mean square error, the BRR, Jackknife and Taylor techniques provide similar accuracy. The differences between the measurements are only in the third decimal places, except for Jackknife in the design with 16 primary sampling units.

In the first design, with 16 census tracts, the bias ratios were approximately 0.10, which means that the bias corresponded to 10% of the standard error of the variance. The ratios decreased to approximately 0.04 and 0.06 in the designs with 32 and 48 census tracts, respectively.

Table 2 shows the results regarding the inclusion of the population parameters within the calculated confidence intervals. The observed confidence levels were lower than the predetermined values (95%), ranging from around 90% for the samples with 16 primary sampling units to around 94% for the larger samples. The results for the three techniques were very similar.

DISCUSSION

From the sampling distributions of the variance estimators, a pattern was seen in relation to the standard error and bias, which were indicators of the precision, reliability and validity of the results obtained. There were small differences among the estimators, with very similar results for BRR and Taylor and slightly inferior performance by the Jackknife technique. The differences were more accentuated for the results with smaller degrees of freedom, while they ceased to exist or became much smaller as the number of primary sampling units increased.

With regard to accuracy, the differences were so small that they would not have been detected had the mean square error been calculated to only two decimal places. Thus, it is hard to speak of greater accuracy for one estimator or another. Data from other studies corroborate the observed similarity of the results relating to the different estimators.

Bean¹ carried out an empirical investigation on the behavior of the Taylor linearization and replication (two BRR estimators) variance estimation methods by using several measurements from the U.S. Health Interview Survey. This author observed that the best results were obtained alternately by BRR and Taylor, and concluded Rev Saúde Pública 2007;41(6)

Table 1. Expected	values stan	dard errors	hips and m	nean callare	errors of var(r)
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Estimate	Estimator	Sampling design			
		8 strata 16 tracts	16 strata 32 tracts	24 strata 48 tracts	
Expected value	Taylor	7.242 x 10 ⁴	3.198 x 10 ⁴	2.100 x 10 ⁴	
E[var(r)]	BRR	7.293 x 10 ⁴	3.198 x 10 ⁴	2.105 x 10 ⁴	
	Jackknife	7.345 x 10 ⁴	3.207 x 10 ⁴	2.107 x 10 ⁴	
Standard error	Taylor	7.260 x 10 ⁴	2.222 x 10 ⁴	1.211 x 10 ⁴	
SE[var(r)]	BRR	7.303 x 10 ⁴	2.219 x 10 ⁴	1.214 x 10 ⁴	
	Jackknife	7.398 x 10 ⁴	2.233 x 10 ⁴	1.217 x 10 ⁴	
Bias	Taylor	6.493×10^3	2.533×10^3	2.079×10^3	
B[var(r)]	BRR	7.003×10^3	2.525×10^3	2.129×10^3	
	Jackknife	7.522×10^3	2.622×10^3	2.147×10^3	
Relative bias	Taylor	0.089	0.036	0.054	
BR[var(r)]	BRR	0.096	0.036	0.055	
	Jackknife	0.102	0.037	0.056	
Mean square	Taylor	5.313 x 10 ⁹	4.942 x 10 ⁸	1.471 x 10 ⁸	
error	BRR	5.383 x 10 ⁹	4.929 x 10 ⁸	1.478 x 10 ⁸	
MSE[var(r)]	Jackknife	5.530 x 10 ⁹	4.992 x 10 ⁸	1.486 x 10 ⁸	
Relative mean	Taylor	1.013	0.483	0.333	
square error	BRR	1.012	0.482	0.333	
RMSE[var(r)]	Jackknife	1.025	0.485	0.335	

Table 2. Coverage of the confidence intervals constructed over 2,000 samples with Student's t values, according to estimation method and sampling design.

Estimator	Sampling design					
	8 strata	16 strata	24 strata			
	16 tracts	32 tracts	48 tracts			
Taylor	90.20	93.90	94.10			
BRR	90.10	93.85	93.95			
Jackknife	90.15	93.85	94.10			

that the latter showed slightly lower mean square errors. Kish & Frankell9 studied BRR, Jackknife and Taylor estimators, using the data from the Current Population Survey carried out by the U. S. Bureau of the Census. The authors concluded that the variability of the Taylor method was the lowest, although the differences in relation to the other estimators were quite small, especially when ratio estimators were used. In terms of precision, Taylor was followed by Jackknife. Kovar et al10 compared the Taylor, BRR, Jackknife and bootstrap estimators in a simulation study based on hypothetical populations that were constructed to resemble the population in the National Assessment of Educational Progress. With regard to the precision of the variance estimates of the ratio estimator, the authors concluded that the best performances were obtained by Taylor and Jackknife, which had equal results.

By means of the bias ratio, it was noted that, for all the techniques, the bias is irrelevant when compared to the magnitude of the standard error of the variance estimates. This leads to the conclusion that, under the conditions prevailing in this study, the accuracy problems of the estimators are more intensely associated with precision problems than with bias.

These results coincide with those obtained by Kish & Frankel⁹ and Bean,¹ who concluded that there is no consistent pattern of lower bias for any particular estimator. Rather, the bias is small and acceptable for all estimators under evaluation. Kovar et al. 10 came to the same conclusion when evaluating the bias in different situations. For designs in which two primary sampling units per stratum were drawn, they took coefficients of variation for the mean denominator of the r ratio that were either less than or equal to, or greater than 10%. They found that, for low coefficients of variation, the bias of the variance estimators was irrelevant. However, when the coefficient of variation for the mean denominator became greater than 10%, BRR showed considerable positive bias, whereas Jackknife and Taylor tended to slightly underestimate the real variance.

With regard to the coverage of the confidence intervals, the results indicated similar performances by the three techniques, as had previously been concluded in

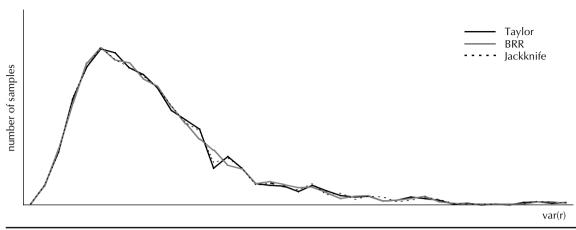


Figure 1. Sampling distribution of the variance of the ratio estimator (mean 1353.955 and standard error 256.750), for the sampling design with 8 strata and 16 primary sampling units, according to variance estimation technique.

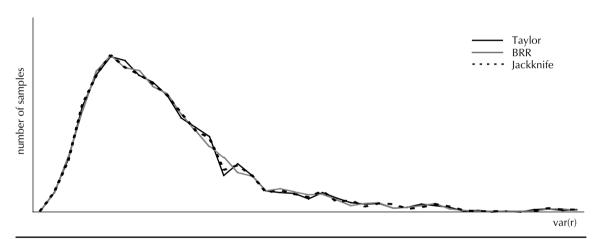


Figure 2. Sampling distribution of the variance of the ratio estimator (mean 1353.767 and standard error 171.616), for the sampling design with 16 strata and 32 primary sampling units, according to variance estimation technique.

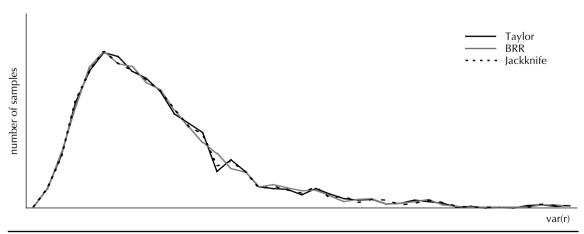


Figure 3. Sampling distribution of the variance of the ratio estimator (mean 1353.107 and standard error 137.564), for the sampling design with 24 strata and 48 primary sampling units, according to variance estimation technique.

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other studies. Although the observed confidence levels were lower than the predetermined ones, they were always higher than 90%, even for the design that only included 16 census tracts. These changes in the confidence level can be regarded as tolerable. However, for inference purposes, researchers should be aware of their existence.

Other empirical studies that evaluated the applicability of confidence intervals to different populations, using different estimators and designs, have also found that the real confidence levels were lower than the nominal levels of normal distribution. The authors of those studies considered that the changes did not prevent the interval estimates from being made with reasonable confidence. In his study, Bean¹ noted coverage greater than 90% for the BRR and Taylor estimators, with results closer to 95% for the first of these. For the ratio estimator, Kish & Frankel9 also found that the coverage was best with BRR (ranging from 90.4 to 94.4%), followed by Jackknife (from 89.4 to 94.3%) and lastly by Taylor (from 88.8 to 94.0%). Kovar et al¹0 considered BRR and Jackknife to be equivalent.

The results from the present study showed that the inferences were valid even for the design in which only eight census tracts were drawn. Even in surveys carried out with many primary sampling units, it may be of interest to study population subgroups that are restricted to some of these clusters. This often occurs in health surveys and it gives rise to the need to obtain interval estimates with a much lower number of primary units than in the overall design. In a study on the performance of the Jackknife estimator for systematic sampling with two to 30 primary sampling units, Burke & Rust³ showed that valid inferences could be made with samples from at least six primary units.

Taking into consideration that the methods evaluated showed equivalent results with regard to precision and bias, decisions on which estimation methods to use in health surveys will depend heavily on operational issues. Furthermore, the availability of software to calculate variance estimates under complex designs is a relevant criterion in this choice.

Various software possibilities have been developed over recent decades. Specific software for variance estimates using one or both of these methods has been created.^{5,11} Moreover, widely used data analysis software such as SAS, STATA and SPSS, has become capable of handling variance estimation in complex study designs, thus expanding the range of alternatives available for analyzing the data coming from household surveys.²

Efforts need to be made by the researchers responsible for conducting these surveys such that information relating to the sampling designs is always included in the data files. The basic information needed is the primary sampling units and the strata to which the study units belong, and their weights, if any.

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