

Variance Estimation of Indicators on Social Exclusion and Poverty using the R Package **laeken**

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Abstract This vignette illustrates the application of variance estimation procedures to indicators on social exclusion and poverty using the R package **laeken**. To be more precise, it describes a general framework for estimating variance and confidence intervals of indicators under complex sampling designs. Currently, the package is focused on bootstrap approaches. While the naive bootstrap does not modify the weights of the bootstrap samples, a calibrated version allows to calibrate each bootstrap sample on auxiliary information before deriving the bootstrap replicate estimate.

1 Introduction

When point estimates of indicators are computed from samples, it is important to also obtain variance estimates and confidence intervals in order to account for variability due to sampling. Other sources of variability such as data editing or imputation may need to be considered as well, but this is not further discussed in this paper. While this vignette targets the topic of variance and confidence interval estimation for the indicators on social exclusion and poverty according to Eurostat (2004, 2009), the aim is not to describe and evaluate the different approaches that have been proposed to date. Instead, the aim is to present the functionality for the statistical environment R (R Development Core Team 2012) implemented in the add-on package **laeken** (Alfons et al. 2012).

It should be noted that the basic design of the package, as well as standard point estimation of the indicators on social exclusion and poverty, is discussed in detail in vignette **laeken-standard** (Templ and Alfons 2011). In addition, vignette **laeken-pareto** (Alfons et al. 2011b) presents more sophisticated methods for point estimation of the indicators, which are less influenced by outliers. Those documents can be viewed from within R with the following commands:

```
R> vignette("laeken-standard")
R> vignette("laeken-pareto")
```

The data basis for the estimation of the indicators on social exclusion and poverty is the *European Union Statistics on Income and Living Conditions* (EU-SILC), which is an annual panel survey conducted in EU member states and other European countries. Package **laeken** provides the synthetic example data **eusilc** consisting of 14 827 observations from 6 000 households. Furthermore, the data were generated from Austrian EU-SILC survey data from 2006 using the data simulation methodology proposed by Alfons et al. (2011a) and implemented in the R package **simPopulation** (Alfons and Kraft 2012). The data set **eusilc** is used in the code examples throughout the paper.

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```
R> library("laeken")
R> data("eusilc")
```

The rest of the paper is organized as follows. Section 2 presents the general wrapper function for estimating variance and confidence intervals of indicators in package **laeken**. The naive and calibrated bootstrap approaches are discussed in Sections 3 and 4, respectively. Section 5 concludes.

2 General wrapper function for variance estimation

The function `variance()` provides a flexible framework for estimating the variance and confidence intervals of indicators such as the *at-risk-of-poverty rate*, the *Gini coefficient*, the *quintile share ratio* and the *relative median at-risk-of-poverty gap*. For a mathematical description and details on the implementation of these indicators in the R package **laeken**, the reader is referred to vignette `laeken-standard` (Templ and Alfons 2011). In any case, `variance()` acts as a general wrapper function for computing variance and confidence interval estimates of indicators on social exclusion and poverty with package **laeken**. The arguments of function `variance()` are shown in the following:

```
R> args(variance)

function (inc, weights = NULL, years = NULL, breakdown = NULL,
  design = NULL, data = NULL, indicator, alpha = 0.05, na.rm = FALSE,
  type = "bootstrap", gender = NULL, method = "mean", ...)
NULL
```

All these arguments are fully described in the R help page of function `variance()`. The most important arguments are:

inc: the income vector.

weights: an optional vector of sample weights.

breakdown: an optional vector giving different domains in which variances and confidence intervals should be computed.

design: an optional vector or factor giving different strata for stratified sampling designs.

data: an optional `data.frame`. If supplied, each of the above arguments should be specified as a character string or an integer or logical vector specifying the corresponding column.

indicator: an object inheriting from the class "indicator" that contains the point estimates of the indicator, such as "arpr" for the at-risk-of-poverty rate, "qsr" for the quintile share ratio, "rmpg" for the relative median at-risk-of-poverty gap, or "gini" for the Gini coefficient.

type: a character string specifying the type of variance estimation to be used. Currently, only "bootstrap" is implemented for variance estimation based on bootstrap resampling.

In the following sections, two bootstrap methods for estimating the variance and confidence intervals of point estimates for complex survey data are described. Furthermore, their application using the function `variance()` from package **laeken** is demonstrated.

3 Naive bootstrap

Let $\mathbf{X} := (\mathbf{x}_1, \dots, \mathbf{x}_n)'$ denote a survey sample with n observations and p variables. Then the *naive bootstrap algorithm* for estimating the variance and confidence interval of an indicator can be summarized as follows:

1. Draw R independent bootstrap samples $\mathbf{X}_1^*, \dots, \mathbf{X}_R^*$ from \mathbf{X} .

2. Compute the bootstrap replicate estimates $\hat{\theta}_r^* := \hat{\theta}(\mathbf{X}_r^*)$ for each bootstrap sample \mathbf{X}_r^* , $r = 1, \dots, R$, where $\hat{\theta}$ denotes an estimator for a certain indicator of interest. Of course the sample weights always need to be considered for the computation of the bootstrap replicate estimates.
3. Estimate the variance $V(\hat{\theta})$ by the variance of the R bootstrap replicate estimates:

$$\hat{V}(\hat{\theta}) := \frac{1}{R-1} \sum_{r=1}^R \left(\hat{\theta}_r^* - \frac{1}{R} \sum_{s=1}^R \hat{\theta}_s^* \right)^2. \quad (1)$$

4. Estimate the confidence interval at confidence level $1 - \alpha$ by one of the following methods (for details, see [Davison and Hinkley 1997](#)):

Percentile method: $\left[\hat{\theta}_{((R+1)\frac{\alpha}{2})}^*, \hat{\theta}_{((R+1)(1-\frac{\alpha}{2}))}^* \right]$, as suggested by [Efron and Tibshirani \(1993\)](#).

Normal approximation: $\hat{\theta} \pm z_{1-\frac{\alpha}{2}} \cdot \hat{V}(\hat{\theta})^{1/2}$ with $z_{1-\frac{\alpha}{2}} = \Phi^{-1}(1 - \frac{\alpha}{2})$.

Basic bootstrap method: $\left[2\hat{\theta} - \hat{\theta}_{((R+1)(1-\frac{\alpha}{2}))}^*, 2\hat{\theta} - \hat{\theta}_{((R+1)\frac{\alpha}{2})}^* \right]$.

For the percentile and the basic bootstrap method, $\hat{\theta}_{(1)}^* \leq \dots \leq \hat{\theta}_{(R)}^*$ denote the order statistics of the bootstrap replicate estimates.

In the following example, the variance and confidence interval of the at-risk-of-poverty rate are estimated with the naive bootstrap procedure. The output of function `variance()` is an object of the same class as the point estimate supplied as the `indicator` argument, but with additional components for the variance and confidence interval. In addition to the point estimate, the income and the sample weights need to be supplied. Furthermore, a stratified sampling design can be considered by specifying the `design` argument, in which case observations are resampled separately within the strata. To ensure reproducibility of the results, the seed of the random number generator is set.

```
R> a <- arpr("eqIncome", weights = "rb050", data = eusilc)
R> variance("eqIncome", weights = "rb050", design = "db040",
+         data = eusilc, indicator = a, bootType = "naive", seed = 123)
```

```
Value:
[1] 14.44422
```

```
Variance:
[1] 0.0920564
```

```
Confidence interval:
      lower      upper
13.87663 15.19417
```

```
Threshold:
[1] 10859.24
```

One of the most convenient features of package `laeken` is that indicators can be evaluated for different subdomains using a single command. This also holds for variance estimation. Using the `breakdown` argument, the example below produces variance and confidence interval estimates for each NUTS2 region in addition to the overall values.

```
R> b <- arpr("eqIncome", weights = "rb050", breakdown = "db040", data = eusilc)
R> variance("eqIncome", weights = "rb050", breakdown = "db040", design = "db040",
+         data = eusilc, indicator = b, bootType = "naive", seed = 123)
```

Value:
[1] 14.44422

Variance:
[1] 0.0920564

Confidence interval:
lower upper
13.87663 15.19417

Value by domain:

	stratum	value
1	Burgenland	19.53984
2	Carinthia	13.08627
3	Lower Austria	13.84362
4	Salzburg	13.78734
5	Styria	14.37464
6	Tyrol	15.30819
7	Upper Austria	10.88977
8	Vienna	17.23468
9	Vorarlberg	16.53731

Variance by domain:

	stratum	var
1	Burgenland	3.5105237
2	Carinthia	1.4133369
3	Lower Austria	0.4456053
4	Salzburg	1.2937926
5	Styria	0.4615967
6	Tyrol	1.0299617
7	Upper Austria	0.3785766
8	Vienna	0.6384621
9	Vorarlberg	1.7601223

Confidence interval by domain:

	stratum	lower	upper
1	Burgenland	16.072806	23.63099
2	Carinthia	10.640776	15.23716
3	Lower Austria	12.196265	15.17182
4	Salzburg	11.913708	16.13315
5	Styria	13.020339	15.89730
6	Tyrol	13.084487	17.51124
7	Upper Austria	9.960467	12.54200
8	Vienna	15.712609	18.96003
9	Vorarlberg	13.604720	19.23431

Threshold:
[1] 10859.24

It should be noted that the workhorse function `bootVar()` is called internally by `variance()` for bootstrap variance and confidence interval estimation. The function `bootVar()` could also be called directly by the user in exactly the same manner. Moreover, variance and confidence interval estimation for any other indicator implemented in package **laeken** is straightforward—the application using function `variance()` or `bootVar()` remains the same.

4 Calibrated bootstrap

Rao and Wu (1988) showed that the naive bootstrap is biased when used in the complex survey context. They propose to increase the variance estimate in the h -th stratum by a factor of $\frac{n_h-1}{n_h}$ (if the bootstrap sample is of the same size). In addition, they describe extensions to sampling without replacement, unequal probability sampling, and two-stage cluster sampling with equal probabilities and without replacement.

Deville and Särndal (1992) and Deville et al. (1993) provide a general description on how to calibrate sample weights to account for known population totals. The naive bootstrap does not include the recalibration of bootstrap samples in order to fit known population totals and therefore is, strictly formulated, not suitable for many practical applications. However, even though a bias might be introduced, the naive bootstrap works well in many situations and is faster to compute than the calibrated version. Hence it is a popular method often used in practice.

In real-world data, the inclusion probabilities for observations in the population are in general not all equal, resulting in different *design weights* for the observations in the sample. Furthermore, the initial design weights are in practice often adjusted by calibration, e.g., to account for non-response or so that certain known population totals can be precisely estimated from the survey sample. To give a simplified example, if the population sizes in different regions are known, the sample weights may be calibrated so that the Horvitz-Thompson estimates (Horvitz and Thompson 1952) of the population sizes equal the known true values. However, when bootstrap samples are drawn from survey data, resampling observations has the effect that such known population totals can no longer be precisely estimated. As a remedy, the sample weights of each bootstrap sample should be calibrated.

The calibrated version of the bootstrap thus results in more precise variance and confidence interval estimation, but comes with higher computational costs than the naive approach. In any case, the *calibrated bootstrap algorithm* is obtained by adding the following step between Steps 1 and 2 of the naive bootstrap algorithm from Section 3:

- 1b. Calibrate the sample weights for each bootstrap sample X_r^* , $r = 1, \dots, R$. Generalized raking procedures are thereby used for calibration: either a multiplicative method known as *raking*, an additive method or a logit method (see Deville and Särndal 1992, Deville et al. 1993).

The function call to `variance()` for the calibrated bootstrap is very similar to its counterpart for the naive bootstrap. A matrix of auxiliary calibration variables needs to be supplied via the argument `X`. In addition, the argument `totals` can be used to supply the corresponding population totals. If the `totals` argument is omitted, as in the following example, the population totals are computed from the sample weights of the original sample. This follows the assumption that those weights are already calibrated on the supplied auxiliary variables.

```
R> variance("eqIncome", weights = "rb050", design = "db040",
+          data = eusilc, indicator = a, X = calibVars(eusilc$db040),
+          seed = 123)
```

Value:

```
[1] 14.44422
```

Variance:

```
[1] 0.09165169
```

Confidence interval:

```
      lower      upper
13.87817 15.19303
```

Threshold:

```
[1] 10859.24
```

Note that the function `calibVars()` transforms a factor into a matrix of binary variables, as required by the calibration function `calibWeights()`, which is called internally. While the default is to use raking for calibration, other methods can be specified via the `method` argument.

5 Conclusions

Both bootstrap procedures for variance and confidence interval estimation of indicators on social exclusion and poverty currently implemented in the R package **laeken** have their strengths. While the naive bootstrap is faster to compute, the calibrated bootstrap in general leads to more precise results. The implementation of other procedures such as linearization techniques (Kovačević and Binder 1997, Deville 1999, Hulliger and Münnich 2006, Osier 2009) or the delete-a-group jackknife (Kott 2001) is future work.

Furthermore, Alfons et al. (2009) demonstrated how the variance of indicators computed from data with imputed values may be underestimated in bootstrap procedures, depending on the indicator itself and the imputation procedure used. They proposed to use the method described in Little and Rubin (2002), which consists of drawing bootstrap samples from the original data with missing values, and to impute the missing data for each bootstrap sample before computing the corresponding bootstrap replicate estimate. Of course, this results in an additional increase of the computation time. The implementation of this procedure in package **laeken** is future work. It should also be noted that multiple imputation is a further possibility to consider the additional uncertainty from imputation when estimating the variance of an indicator (see Little and Rubin 2002).

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