Poverty and Inequality with Complex Survey Data

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2020-05-21

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Chapter 1

Introduction

The R convey library estimates measures of poverty, inequality, and wellbeing. There are two other R libraries covering this subject, vardpoor and laeken, however, only convey integrates seamlessly with the R survey package.

convey is free and open-source software that runs inside the R environment for statistical computing. Anyone can review and propose changes to the source code for this software. Readers are welcome to propose changes to this book as well.

1.1 Installation

In order to work with the **convey** library, you will need to have R running on your machine. If you have never used R before, you will need to install that software before **convey** can be accessed. Check out FlowingData for a concise list of resources for new R users. Once you have R loaded on your machine, you can install..

• the latest released version from CRAN with

```
install.packages("convey")
```

• the latest development version from github with

```
devtools::install_github("djalmapessoa/convey")
```

1.2 Complex surveys and statistical inference

In this book, we demonstrate how to measure poverty and income concentration in a population based on microdata collected from a complex survey sample. Most surveys administered by government agencies or larger research organizations utilize a sampling design that violates the assumption of simple random sampling (SRS), including:

- 1. Different units selection probabilities;
- 2. Clustering of units;
- 3. Stratification of clusters;
- 4. Reweighting to compensate for missing values and other adjustments.

Therefore, basic unweighted R commands such as mean() or glm() will not properly account for the weighting nor the measures of uncertainty (such as the confidence intervals) present in the dataset. For some examples of publicly-available complex survey data sets, see http://asdfree.com.

Unlike other software, the R convey package does not require that the user specify these parameters throughout the analysis. So long as the svydesign object or svrepdesign object has been constructed properly at the outset of the analysis, the convey package will incorporate the survey design automatically and produce statistics and variances that take the complex sample into account.

1.3 Usage Examples

In the following example, we've loaded the data set eusilc from the R library laeken.

```
library(laeken)
data(eusilc)
```

Next, we create an object of class survey.design using the function svydesign of the library survey:

```
library(survey)
des_eusilc <- svydesign(ids = ~rb030, strata =~db040, weights = ~rb050, data = eusilc</pre>
```

Right after the creation of the design object des_eusilc, we should use the function convey_prep that adds an attribute to the survey design which saves information on the design object based upon the whole sample, needed to work with subset designs.

```
library(convey)
des_eusilc <- convey_prep( des_eusilc )</pre>
```

To estimate the at-risk-of-poverty rate, we use the function svyarpt:

```
svyarpr(~eqIncome, design=des_eusilc)
                     SE
            arpr
eqIncome 0.14444 0.0028
To estimate the at-risk-of-poverty rate across domains defined by the variable
db040 we use:
svyby(~eqIncome, by = ~db040, design = des_eusilc, FUN = svyarpr, deff = FALSE)
                      db040 eqIncome
Burgenland
                 Burgenland 0.1953984 0.017202243
Carinthia
                  Carinthia 0.1308627 0.010610622
Lower Austria Lower Austria 0.1384362 0.006517660
                   Salzburg 0.1378734 0.011579280
Salzburg
Styria
                     Styria 0.1437464 0.007452360
Tyrol
                      Tyrol 0.1530819 0.009880430
Upper Austria Upper Austria 0.1088977 0.005928336
Vienna
                     Vienna 0.1723468 0.007682826
Vorarlberg
                 Vorarlberg 0.1653731 0.013754670
Using the same data set, we estimate the quintile share ratio:
# for the whole population
svyqsr(~eqIncome, design=des_eusilc, alpha1= .20)
          qsr
                  SE
eqIncome 3.97 0.0426
# for domains
svyby(~eqIncome, by = ~db040, design = des_eusilc,
 FUN = svyqsr, alpha1= .20, deff = FALSE)
                      db040 eqIncome
Burgenland
                 Burgenland 5.008486 0.32755685
Carinthia
                  Carinthia 3.562404 0.10909726
Lower Austria Lower Austria 3.824539 0.08783599
Salzburg
                   Salzburg 3.768393 0.17015086
Styria
                     Styria 3.464305 0.09364800
Tyrol
                      Tyrol 3.586046 0.13629739
Upper Austria Upper Austria 3.668289 0.09310624
Vienna
                     Vienna 4.654743 0.13135731
                 Vorarlberg 4.366511 0.20532075
Vorarlberg
These functions can be used as S3 methods for the classes survey.design and
svyrep.design.
```

Let's create a design object of class svyrep.design and run the function

convey_prep on it:

```
des_eusilc_rep <- as.svrepdesign(des_eusilc, type = "bootstrap")</pre>
des_eusilc_rep <- convey_prep(des_eusilc_rep)</pre>
and then use the function svyarpr:
svyarpr(~eqIncome, design=des_eusilc_rep)
            arpr
                      SE
eqIncome 0.14444 0.0025
svyby(~eqIncome, by = ~db040, design = des_eusilc_rep, FUN = svyarpr, deff = FALSE)
                       db040 eqIncome se.eqIncome
Burgenland
                  Burgenland 0.1953984 0.016713791
Carinthia
                  Carinthia 0.1308627 0.012061625
Lower Austria Lower Austria 0.1384362 0.007294696
                    Salzburg 0.1378734 0.010050357
Salzburg
Styria
                      Styria 0.1437464 0.008558783
Tyrol
                       Tyrol 0.1530819 0.010328225
Upper Austria Upper Austria 0.1088977 0.006212301
                      Vienna 0.1723468 0.007259732
Vienna
Vorarlberg
                 Vorarlberg 0.1653731 0.012792618
The functions of the library convey are called in a similar way to the functions
in library survey.
```

It is also possible to deal with missing values by using the argument na.rm.

```
# survey.design using a variable with missings
svygini( ~ py010n , design = des_eusilc )
       gini SE
py010n
        NA NA
svygini( ~ py010n , design = des_eusilc , na.rm = TRUE )
          gini
                   SE
py010n 0.64606 0.0036
# svyrep.design using a variable with missings
svygini( ~ py010n , design = des_eusilc_rep )
       gini SE
py010n
        NA NA
svygini( ~ py010n , design = des_eusilc_rep , na.rm = TRUE )
          gini
py010n 0.64606 0.0043
```

1.4 Underlying Calculations

In what follows, we often use the linearization method as a tool to produce an approximation for the variance of an estimator. From the linearized variable z of an estimator T, we get from the expression (1.1) an estimate of the variance of T

If T can be expressed as a function of the population totals $T=g(Y_1,Y_2,\ldots,Y_n)$, and if g is linear, the estimation of the variance of $T=g(Y_1,Y_2,\ldots,Y_n)$ is straightforward. If g is not linear but is a 'smooth' function, then it is possible to approximate the variance of $g(Y_1,Y_2,\ldots,Y_n)$ by the variance of its first order Taylor expansion. For example, we can use Taylor expansion to linearize the ratio of two totals. However, there are situations where Taylor linearization cannot be immediately possible, either because T cannot be expressed as functions of the population totals, or because g is not a smooth function. An example is the case where T is a quantile.

In these cases, it might work an alternative form of linearization of T, by Influence Function, as defined in (1.2), proposed in (Deville, 1999). Also, it could be used replication methods such as bootstrap and jackknife.

In the convey library, there are some basic functions that produce the linearized variables needed to measure income concentration and poverty. For example, looking at the income variable in some complex survey dataset, the quantile of that income variable can be linearized by the function convey::svyiqalpha and the sum total below any quantile of the variable is linearized by the function convey::svyisq.

From the linearized variables of these basic estimates, it is possible by using rules of composition, valid for influence functions, to derive the influence function of more complex estimates. By definition the influence function is a Gateaux derivative and the rules rules of composition valid for Gateaux derivatives also hold for Influence Functions.

The following property of Gateaux derivatives was often used in the library convey. Let g be a differentiable function of m variables. Suppose we want to compute the influence function of the estimator $g(T_1, T_2, \dots, T_m)$, knowing the Influence function of the estimators T_i , $i=1,\dots,m$. Then the following holds:

$$I(g(T_1,T_2,\dots,T_m)) = \sum_{i=1}^m \frac{\partial g}{\partial T_i} I(T_i)$$

In the library convey this rule is implemented by the function contrastinf which uses the R function deriv to compute the formal partial derivatives $\frac{\partial g}{\partial T}$.

For example, suppose we want to linearize the Relative median poverty gap(rmpg), defined as the difference between the at-risk-of-poverty threshold (arpt) and the median of incomes less than the arpt relative to the arprt:

$$rmpg = \frac{arpt - medpoor}{arpt}$$

where medpoor is the median of incomes less than arpt.

Suppose we know how to linearize arpt and medpoor, then by applying the function contrastinf with

$$g(T_1, T_2) = \frac{(T_1 - T_2)}{T_1}$$

we linearize the rmpg.

1.5 The Variance Estimator

Using the notation in (Osier, 2009), the variance of the estimator $T(\hat{M})$ can approximated by:

$$Var\left[T(\hat{M})\right] \cong var\left[\sum_{s} w_{i} z_{i}\right] \tag{1.1}$$

The linearized variable z is given by the derivative of the functional:

$$z_k = lim_{t\rightarrow 0} \frac{T(M+t\delta_k) - T(M)}{t} = IT_k(M) \eqno(1.2)$$

where, δ_k is the Dirac measure in k: $\delta_k(i) = 1$ if and only if i = k.

This **derivative** is called **Influence Function** and was introduced in the area of **Robust Statistics**.

1.6 Influence Functions

Some measures of poverty and income concentration are defined by non-differentiable functions so that it is not possible to use Taylor linearization to estimate their variances. An alternative is to use **Influence functions** as described in (Deville, 1999) and (Osier, 2009). The convey library implements this methodology to work with survey.design objects and also with svyrep.design objects.

Some examples of these measures are:

• At-risk-of-poverty threshold: $arpt = .60q_{.50}$ where $q_{.50}$ is the income median;

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- At-risk-of-poverty rate $arpr = \frac{\sum_{U} 1(y_i \leq arpt)}{N}.100$
- Quintile share ratio

$$qsr = \frac{\sum_{U} 1(y_i > q_{.80})}{\sum_{U} 1(y_i \leq q_{.20})}$$

• Gini coefficient $1+G=\frac{2\sum_U(r_i-1)y_i}{N\sum_Uy_i}$ where r_i is the rank of y_i .

Note that it is not possible to use Taylor linearization for these measures because they depend on quantiles and the Gini is defined as a function of ranks. This could be done using the approach proposed by Deville (1999) based upon influence functions.

Let U be a population of size N and M be a measure that allocates mass one to the set composed by one unit, that is $M(i)=M_i=1$ if $i\in U$ and M(i)=0 if $i\notin U$

Now, a population parameter θ can be expressed as a functional of M $\theta = T(M)$

Examples of such parameters are:

- Total: $Y = \sum_{U} y_i = \sum_{U} y_i M_i = \int y dM = T(M)$
- Ratio of two totals: $R = \frac{Y}{X} = \frac{\int y dM}{\int x dM} = T(M)$
- Cumulative distribution function: $F(x) = \frac{\sum_U 1(y_i \le x)}{N} = \frac{\int 1(y \le x) dM}{\int dM} = T(M)$

To estimate these parameters from the sample, we replace the measure M by the estimated measure \hat{M} defined by: $\hat{M}(i) = \hat{M}_i = w_i$ if $i \in s$ and $\hat{M}(i) = 0$ if $i \notin s$.

The estimators of the population parameters can then be expressed as functional of the measure \hat{M} .

- Total: $\hat{Y} = T(\hat{M}) = \int y d\hat{M} = \sum_s w_i y_i$
- Ratio of totals: $\hat{R} = T(\hat{M}) = \frac{\int y d\hat{M}}{\int x d\hat{M}} = \frac{\sum_s w_i y_i}{\sum_s w_i x_i}$
- Cumulative distribution function: $\hat{F}(x) = T(\hat{M}) = \frac{\int 1(y \le x)d\hat{M}}{\int d\hat{M}} = \frac{\sum_s w_i 1(y_i \le x)}{\sum_s w_i}$

1.7 Influence Function Examples

• Total:

$$\begin{split} IT_k(M) &= lim_{t \rightarrow 0} \frac{T(M + t\delta_k) - T(M)}{t} \\ &= lim_{t \rightarrow 0} \frac{\int y.d(M + t\delta_k) - \int y.dM}{t} \\ &= lim_{t \rightarrow 0} \frac{\int yd(t\delta_k)}{t} = y_k \end{split}$$

• Ratio of two totals:

$$\begin{split} IR_k(M) &= I\left(\frac{U}{V}\right)_k(M) = \frac{V(M) \times IU_k(M) - U(M) \times IV_k(M)}{V(M)^2} \\ &= \frac{Xy_k - Yx_k}{X^2} = \frac{1}{X}(y_k - Rx_k) \end{split}$$

1.8 Examples of Linearization Using the Influence Function

• At-risk-of-poverty threshold:

$$arpt = 0.6 \times m$$

where m is the median income.

$$z_k = -\frac{0.6}{f(m)} \times \frac{1}{N} \times \left[I(y_k \leq m - 0.5)\right]$$

• At-risk-of-poverty rate:

$$\begin{split} arpr &= \frac{\sum_{U} I(y_i \leq t)}{\sum_{U} w_i}.100\\ z_k &= \frac{1}{N} \left[I(y_k \leq t) - t \right] - \frac{0.6}{N} \times \frac{f(t)}{f(m)} \left[I(y_k \leq m) - 0.5 \right] \end{split}$$

where:

N - population size;

t - at-risk-of-poverty threshold;

 y_k - income of person k;

m - median income;

f - income density function;

1.9 Replication Designs

All major functions in the library convey have S3 methods for the classes: survey.design, svyrep.design and DBIdesign. When the argument design is

a survey design with replicate weights created by the library survey, convey uses the method svrepdesign.

Considering the remarks in (Wolter, 1985), p. 163, concerning the deficiency of the Jackknife method in estimating the variance of quantiles, we adopted the type bootstrap instead.

The function bootVar from the library laeken, (Alfons et al., 2014), also uses the bootstrap method to estimate variances.

1.10 Decomposition

Some inequality and multidimensional poverty measures can be decomposed. As of December 2016, the decomposition methods in convey are limited to group decomposition.

For instance, the generalized entropy index can be decomposed into between and within group components. This sheds light on a very simple question: of the overall inequality, how much can be explained by inequalities between groups and within groups? Since this measure is additive decomposable, one can get estimates of the coefficients, SEs and covariance between components. For a more practical approach, see (Lima, 2013).

The Alkire-Foster class of multidimensional poverty indices can be decomposed by dimension and groups. This shows how much each group (or dimension) contribute to the overall poverty.

This technique can help understand where and who is more affected by inequality and poverty, contributing to more specific policy and economic analysis.

Chapter 2

Poverty Indices

Poverty is an issue discussed since long time ago. As Ravallion (2016) points out, Aristotle and Confucius discussed ideas about poverty. In fact, Aristotle's ideas influenced Thomas Aquinas, one of the pillars of Western philosophy. Since then, societies changed, modifying the theories of justice underlying the idea of poverty.

As the concept and the ethics towards poverty change, so does its measurement. From basic measures like the headcount rate to more complex metrics, such as the FGT index, poverty measurement evolved. Nowadays, poverty measures estimates are calculated using household surveys and censuses (Deaton, 1997). Yet, only recently the aspects of statistical inference combining such measures and survey designs were explored¹. These advances become even more important given the recent efforts in poverty mapping, an analytical method that combined poverty analysis and small area estimation, like Elbers et al. (2003) and Bedi et al. (2007).

The following subsections shows how poverty estimates and their sampling errors can be estimated using simple commands from the **convey** package.

2.1 At Risk of Poverty Threshold (svyarpt)

The at-risk-of-poverty threshold (ARPT) is a measure used to define the people whose incomes imply a low standard of living in comparison to the general living standards. I.e., even though some people are not below the effective poverty line, those below the ARPT can be considered "almost deprived".

This measure is defined as 0.6 times the median income for the entire population:

¹For instance, see Deville (1999), Berger and Skinner (2003), Bhattacharya (2007), and Osier (2009).

```
arpt = 0.6 \times median(y),
```

where, y is the income variable and median is estimated for the whole population. The details of the linearization of the arpt are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a arpt coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the laeken library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the arpt coefficient
# using the R vardpoor library
varpoord_arpt_calculation <-</pre>
    varpoord(
        # analysis variable
        Y = "eqincome",
        # weights variable
        w_final = "rb050",
```

```
# row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # arpt coefficient function
        type = "linarpt",
      # poverty threshold range
      order_quant = 50L ,
        # qet linearized variable
      outp_lin = TRUE
# construct a survey.design
# using our recommended setup
des_eusilc <-
   svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc
   )
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_arpt_calculation$all_result$value
```

```
coef( svyarpt( ~ eqincome , des_eusilc ) )
## eqincome
## 10859.24
# linearized variables do match
# vardpoor
lin_arpt_varpoord<- varpoord_arpt_calculation$lin_out$lin_arpt</pre>
# convey
lin_arpt_convey <- attr(svyarpt( ~ eqincome , des_eusilc ),"lin")</pre>
# check equality
all.equal(lin_arpt_varpoord, lin_arpt_convey )
## [1] TRUE
# variances do not match exactly
attr( svyarpt( ~ eqincome , des_eusilc ) , 'var' )
##
            eqincome
## eqincome 2564.027
varpoord_arpt_calculation$all_result$var
## [1] 2559.442
# standard errors do not match exactly
varpoord_arpt_calculation$all_result$se
## [1] 50.59093
SE( svyarpt( ~ eqincome , des_eusilc ) )
            eqincome
## eqincome 50.63622
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions convey::svyarpt and vardpoor::linarpt produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
```

```
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = ~rb030 ,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svyarpt( ~ eqincome , des_eusilc_ultimate_cluster ) , 'var' )
            eqincome
## eqincome 2559.442
varpoord_arpt_calculation$all_result$var
## [1] 2559.442
varpoord_arpt_calculation$all_result$se
## [1] 50.59093
SE( svyarpt( ~ eqincome , des_eusilc_ultimate_cluster ) )
##
            eqincome
## eqincome 50.59093
```

For additional usage examples of svyarpt, type ?convey::svyarpt in the R console.

2.2 At Risk of Poverty Ratio (svyarpr)

The at-risk-of-poverty rate (ARPR) is the share of persons with an income below the at-risk-of-poverty threshold (arpt). The logic behind this measure is that although most people below the ARPT cannot be considered "poor", they are the ones most vulnerable to becoming poor in the event of a negative economic phenomenon.

The ARPR is a composite estimate, taking into account both the sampling error in the proportion itself and that in the ARPT estimate. The details of the linearization of the arpr and are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a ARPR coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the vardpoor library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the arpr coefficient
# using the R vardpoor library
varpoord_arpr_calculation <-</pre>
    varpoord(
        # analysis variable
        Y = "eqincome",
        # weights variable
        w final = "rb050",
        # row number variable
```

```
ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # arpr coefficient function
        type = "linarpr",
      # poverty threshold range
      order_quant = 50L ,
      # get linearized variable
      outp_lin = TRUE
   )
# construct a survey.design
# using our recommended setup
des_eusilc <-
   svydesign(
        ids = ~rb030,
        strata = ~db040,
       weights = \sim rb050,
        data = eusilc
   )
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_arpr_calculation$all_result$value
```

```
coef( svyarpr( ~ eqincome , des_eusilc ) ) * 100
## eqincome
## 14.44422
# linearized variables do match
# vardpoor
lin_arpr_varpoord<- varpoord_arpr_calculation$lin_out$lin_arpr</pre>
# convey
lin_arpr_convey <- attr(svyarpr( ~ eqincome , des_eusilc ),"lin")</pre>
# check equality
all.equal(lin_arpr_varpoord,100*lin_arpr_convey )
## [1] "Mean relative difference: 0.2264738"
# variances do not match exactly
attr( svyarpr( ~ eqincome , des_eusilc ) , 'var' ) * 10000
##
              eqincome
## eqincome 0.07599778
varpoord_arpr_calculation$all_result$var
## [1] 0.08718569
# standard errors do not match exactly
varpoord_arpr_calculation$all_result$se
## [1] 0.2952722
SE( svyarpr( ~ eqincome , des_eusilc ) ) * 100
             eqincome
## eqincome 0.2756769
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions $\verb"convey":svyarp"$ and $\verb"vardpoo"::linarp"$ produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
```

```
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svyarpr( ~ eqincome , des_eusilc_ultimate_cluster ) , 'var' ) * 10000
              eqincome
## eqincome 0.07586194
varpoord_arpr_calculation$all_result$var
## [1] 0.08718569
# matches
varpoord_arpr_calculation$all_result$se
## [1] 0.2952722
SE( svyarpr( ~ eqincome , des_eusilc_ultimate_cluster ) ) * 100
             eqincome
## eqincome 0.2754305
```

For additional usage examples of svyarpr, type ?convey::svyarpr in the R console.

2.3 Relative Median Income Ratio (svyrmir)

The relative median income ratio (rmir) is the ratio of the median income of people aged above a value (65) to the median of people aged below the same value. In mathematical terms,

$$rmir = \frac{median\{y_i; age_i > 65\}}{median\{y_i; age_i \leq 65\}}.$$

The details of the linearization of the rmir and are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a rmir coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the vardpoor library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the rmir coefficient
# using the R vardpoor library
varpoord_rmir_calculation <-</pre>
    varpoord(
        # analysis variable
        Y = "eqincome",
        # weights variable
        w_final = "rb050",
```

```
# row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
      # age variable
      age = "age",
        # rmir coefficient function
        type = "linrmir",
      # poverty threshold range
      order_quant = 50L ,
      # get linearized variable
     outp_lin = TRUE
   )
# construct a survey.design
# using our recommended setup
des_eusilc <-
   svydesign(
        ids = ~rb030,
        strata = ~db040,
       weights = \sim rb050 ,
       data = eusilc
   )
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
```

```
# coefficients do match
varpoord_rmir_calculation$all_result$value
## [1] 0.9330361
coef( svyrmir( ~ eqincome , des_eusilc, age = ~age ) )
## eqincome
## 0.9330361
# linearized variables do match
# vardpoor
lin_rmir_varpoord<- varpoord_rmir_calculation$lin_out$lin_rmir</pre>
# convey
lin_rmir_convey <- attr(svyrmir( ~ eqincome , des_eusilc, age = ~age ),"lin")</pre>
# check equality
all.equal(lin_rmir_varpoord, lin_rmir_convey[,1] )
## [1] TRUE
# variances do not match exactly
attr( svyrmir( ~ eqincome , des_eusilc, age = ~age ) , 'var' )
##
               eqincome
## eqincome 0.000127444
varpoord_rmir_calculation$all_result$var
## [1] 0.0001272137
# standard errors do not match exactly
varpoord_rmir_calculation$all_result$se
## [1] 0.0112789
SE( svyrmir( ~ eqincome , des_eusilc , age = ~age) )
##
              eqincome
## eqincome 0.01128911
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions convey::svyrmir and vardpoor::linrmir produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )</pre>
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = ~rb030,
        strata = ~ db040
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svyrmir( ~ eqincome , des_eusilc_ultimate_cluster , age = ~age ) , 'var' )
##
                eqincome
## eqincome 0.0001272137
varpoord_rmir_calculation$all_result$var
## [1] 0.0001272137
# matches
varpoord_rmir_calculation$all_result$se
## [1] 0.0112789
SE( svyrmir( ~ eqincome , des_eusilc_ultimate_cluster, age = ~age ) )
##
             eqincome
## eqincome 0.0112789
```

For additional usage examples of svyrmir, type ?convey::svyrmir in the R

console.

2.4 Relative Median Poverty Gap (svyrmpg)

The relative median poverty gap (rmpg) is the relative difference between the median income of people having income below the arpt and the arpt itself:

$$rmpg = \frac{median\{y_i, y_i < arpt\} - arpt}{arpt}$$

The details of the linearization of the rmpg are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a rmpg coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the vardpoor library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the rmpg coefficient
# using the R vardpoor library
varpoord_rmpg_calculation <-</pre>
    varpoord(
        # analysis variable
```

```
Y = "eqincome",
        # weights variable
        w_final = "rb050",
        # row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040"
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # rmpg coefficient function
        type = "linrmpg",
      # poverty threshold range
     order_quant = 50L ,
      # get linearized variable
     outp_lin = TRUE
   )
# construct a survey.design
# using our recommended setup
des_eusilc <-
   svydesign(
       ids = - rb030,
        strata = ~db040,
       weights = \sim rb050 ,
       data = eusilc
   )
```

```
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_rmpg_calculation$all_result$value
## [1] 18.9286
coef( svyrmpg( ~ eqincome , des eusilc ) ) * 100
## eqincome
## 18.9286
# linearized variables do match
# vardpoor
lin_rmpg_varpoord<- varpoord_rmpg_calculation$lin_out$lin_rmpg</pre>
# convey
lin_rmpg_convey <- attr(svyrmpg( ~ eqincome , des_eusilc ),"lin")</pre>
# check equality
all.equal(lin_rmpg_varpoord, 100*lin_rmpg_convey[,1] )
## [1] TRUE
# variances do not match exactly
attr( svyrmpg( ~ eqincome , des_eusilc ) , 'var' ) * 10000
##
            eqincome
## eqincome 0.332234
varpoord_rmpg_calculation$all_result$var
## [1] 0.3316454
# standard errors do not match exactly
varpoord_rmpg_calculation$all_result$se
## [1] 0.5758866
SE( svyrmpg( ~ eqincome , des_eusilc ) ) * 100
##
             eqincome
## eqincome 0.5763974
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions $\verb"convey":svyrmpg"$ and $\verb"vardpoor":linrmpg"$ produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative

setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )</pre>
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = - rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svyrmpg( ~ eqincome , des_eusilc_ultimate_cluster ) , 'var' ) * 10000
##
             eqincome
## eqincome 0.3316454
varpoord_rmpg_calculation$all_result$var
## [1] 0.3316454
# matches
varpoord_rmpg_calculation$all_result$se
## [1] 0.5758866
SE( svyrmpg( ~ eqincome , des_eusilc_ultimate_cluster ) ) * 100
##
             eqincome
## eqincome 0.5758866
```

For additional usage examples of svyrmpg, type ?convey::svyrmpg in the R console.

2.5 Median Income Below the At Risk of Poverty Threshold (svypoormed)

Median income below the at-risk-of-poverty- threshold (poormed) is median of incomes of people having the income below the arpt:

```
poormed = median\{y_i; y_i < arpt\}
```

The details of the linearization of the poormed are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a poormed coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the vardpoor library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the poormed coefficient
# using the R vardpoor library
varpoord_poormed_calculation <-</pre>
   varpoord(
```

```
# analysis variable
        Y = "eqincome",
        # weights variable
        w_{final} = "rb050",
        # row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # poormed coefficient function
        type = "linpoormed",
      # poverty threshold range
     order_quant = 50L ,
      # get linearized variable
     outp_lin = TRUE
   )
# construct a survey.design
# using our recommended setup
des_eusilc <-
   svydesign(
       ids = - rb030,
       strata = ~db040,
       weights = \sim rb050,
       data = eusilc
```

```
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_poormed_calculation$all_result$value
## [1] 8803.735
coef( svypoormed( ~ eqincome , des_eusilc ) )
## eqincome
## 8803.735
# linearized variables do match
# vardpoor
lin_poormed_varpoord<- varpoord_poormed_calculation$lin_out$lin_poormed
lin_poormed_convey <- attr(svypoormed( ~ eqincome , des_eusilc ),"lin")</pre>
# check equality
all.equal(lin_poormed_varpoord, lin_poormed_convey )
## [1] TRUE
# variances do not match exactly
attr( svypoormed( ~ eqincome , des_eusilc ) , 'var' )
            eqincome
## eqincome 5311.47
varpoord_poormed_calculation$all_result$var
## [1] 5302.086
# standard errors do not match exactly
varpoord_poormed_calculation$all_result$se
## [1] 72.81542
SE( svypoormed( ~ eqincome , des_eusilc ) )
##
            eqincome
## eqincome 72.87983
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions convey::svypoormed and vardpoor::linpoormed produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor)

defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )</pre>
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svypoormed( ~ eqincome , des_eusilc_ultimate_cluster ) , 'var' )
##
            eqincome
## eqincome 5302.086
varpoord_poormed_calculation$all_result$var
## [1] 5302.086
# matches
varpoord_poormed_calculation$all_result$se
## [1] 72.81542
SE( svypoormed( ~ eqincome , des_eusilc_ultimate_cluster ) )
            eqincome
## eqincome 72.81542
```

For additional usage examples of svypoormed, type ?convey::svypoormed in the R console.

2.6 Foster-Greer-Thorbecke class (svyfgt, svyfgtdec)

Foster et al. (1984) proposed a family of indicators to measure poverty. This class of FGT measures, can be defined as

$$p = \frac{1}{N} \sum_{k \in U} h(y_k, \theta),$$

where

$$h(y_k,\theta) = \left\lceil \frac{(\theta-y_k)}{\theta} \right\rceil^{\gamma} \delta \left\{ y_k \leq \theta \right\},$$

where: θ is the poverty threshold; δ the indicator function that assigns value 1 if the condition $\{y_k \leq \theta\}$ is satisfied and 0 otherwise, and γ is a non-negative constant.

If $\gamma=0$, the FGT(0) equals the poverty headcount ratio, which accounts for the spread of poverty. If $\gamma=1$, FGT(1) is the mean of the normalized income shortfall of the poor. By doing so, the measure takes into account both the spread and the intensity of poverty. When $\gamma=2$, the relative weight of larger shortfalls increases even more, which yields a measure that accounts for poverty severity, i.e., the inequality among the poor. This way, a transfer from a poor person to an even poorer person would reduce the FGT(2).

Although Foster et al. (1984) already presented a decomposition for the FGT(2) index, Aristondo et al. (2010) provided a general formula that decomposes the FGT(γ) for any $\gamma \geq 2$. Put simply, any such FGT(γ) index can be seen as function of the headcount ratio, the average normalized income gap among the poor and a generalized entropy index of the normalized income gaps among poor. In mathematical terms,

$$FGT_{\gamma} = FGT_0 \cdot I^{\gamma} \cdot \left[1 + \left(\gamma^2 - \gamma\right)GEI_{\gamma}^*\right], \ \gamma \geq 2$$

where I is the average normalized income gap among the poor and GEI_{γ}^{*} is a generalized entropy index of such income gaps among the poor.

This result is particularly useful, as one can attribute cross-sectional differences of a FGT index to differences in the spread, depth and inequality of poverty.

The FGT poverty class and its decomposition is implemented in the library convey by the function svyfgt and svyfgtdec, respectively. The argument thresh_type of this function defines the type of poverty threshold adopted. There are three possible choices:

- 1. abs fixed and given by the argument thresh value
- 2. relq a proportion of a quantile fixed by the argument proportion and the quantile is defined by the argument order.
- 3. relm a proportion of the mean fixed the argument proportion

The quantile and the mean involved in the definition of the threshold are estimated for the whole population. When $\gamma = 0$ and $\theta = .6*MED$ the measure is equal to the indicator arpr computed by the function svyarpr. The linearization of the FGT(0) is presented in Berger and Skinner (2003).

Next, we give some examples of the function **svyfgt** to estimate the values of the FGT poverty index.

Consider first the poverty threshold fixed ($\gamma=0$) in the value 10000. The headcount ratio (FGT0) is

```
svyfgt(~eqincome, des_eusilc, g=0, abs_thresh=10000)
```

```
fgt0 SE eqincome 0.11444 0.0027
```

The poverty gap ratio (FGT(1)) ($\gamma = 1$) index for the poverty threshold fixed at the same value is

```
svyfgt(~eqincome, des_eusilc, g=1, abs_thresh=10000)
```

```
fgt1 SE eqincome 0.032085 0.0011
```

To estimate the FGT(0) with the poverty threshold fixed at 0.6*MED we fix the argument type_thresh="relq" and use the default values for percent and order:

```
svyfgt(~eqincome, des_eusilc, g=0, type_thresh= "relq")
```

```
fgt0 SE eqincome 0.14444 0.0028
```

that matches the estimate obtained by

```
svyarpr(~eqincome, design=des_eusilc, .5, .6)
```

```
arpr SE eqincome 0.14444 0.0028
```

To estimate the poverty gap ratio with the poverty threshold equal to 0.6*MEAN, we use:

```
svyfgt(~eqincome, des_eusilc, g=1, type_thresh= "relm")
```

```
fgt1 SE eqincome 0.051187 0.0011
```

A replication example

In July 2006, Jenkins (2008) presented at the North American Stata Users' Group Meetings on the stata Atkinson Index command. The example below reproduces those statistics.

In order to match the presentation's results using the svyfgt function from the convey library, the poverty threshold was considered absolute despite being directly estimated from the survey sample. This effectively treats the variance of the estimated poverty threshold as zero; svyfgt does not account for the uncertainty of the poverty threshold when the level has been stated as absolute with the abs_thresh= parameter. In general, we would instead recommend using either relq or relm in the type_thresh= parameter in order to account for the added uncertainty of the poverty threshold calculation. This example serves only to show that svyfgt behaves properly as compared to other software.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the foreign library
library(foreign)
# create a temporary file on the local disk
tf <- tempfile()</pre>
# store the location of the presentation file
presentation_zip <- "http://repec.org/nasug2006/nasug2006_jenkins.zip"</pre>
# download jenkins' presentation to the temporary file
download.file( presentation_zip , tf , mode = 'wb' )
# unzip the contents of the archive
presentation_files <- unzip( tf , exdir = tempdir() )</pre>
# load the institute for fiscal studies' 1981, 1985, and 1991 data.frame objects
x81 <- read.dta( grep( "ifs81" , presentation_files , value = TRUE ) )</pre>
x85 <- read.dta( grep( "ifs85" , presentation_files , value = TRUE ) )
x91 <- read.dta( grep( "ifs91" , presentation_files , value = TRUE ) )</pre>
# NOTE: we recommend using ?convey::svyarpt rather than this unweighted calculation #
```

Survey: Mean estimation

```
# calculate 60% of the unweighted median income in 1981
unwtd_arpt81 <- quantile( x81$eybhc0 , 0.5 ) * .6</pre>
# calculate 60% of the unweighted median income in 1985
unwtd_arpt85 <- quantile( x85$eybhc0 , 0.5 ) * .6</pre>
# calculate 60% of the unweighted median income in 1991
unwtd_arpt91 <- quantile( x91$eybhc0 , 0.5 ) * .6</pre>
# stack each of these three years of data into a single data.frame
x <- rbind( x81 , x85 , x91 )
Replicate the author's survey design statement from stata code..
. ge poor = (year=1981)*(x < $z_81) + (year=1985)*(x < $z_85) + (year=1991)*(x < $z_91)
. * account for clustering within HHs
. svyset hrn [pweight = wgt]
.. into R code:
# initiate a linearized survey design object
y <- svydesign( ~ hrn , data = x , weights = ~ wgt )
# immediately run the `convey_prep` function on the survey design
z <- convey_prep( y )</pre>
Replicate the author's headcount ratio results with stata..
. svy: mean poor if year == 1981
(running mean on estimation sample)
Survey: Mean estimation
                                  Number of obs = 9772
Number of strata =
                    1
Number of PSUs = 7476
                                  Population size = 5.5e+07
                                  Design df = 7475
                        Linearized
           | Mean Std. Err. [95% Conf. Interval]
______
      poor | .1410125 .0044859 .132219
                                                   .149806
. svy: mean poor if year == 1985
(running mean on estimation sample)
```

```
Number of strata = 1
                         Number of obs = 8991
Number of PSUs = 6972
                         Population size = 5.5e+07
                         Design df
                                 = 6971
_____
              Linearized
        Mean Std. Err. [95% Conf. Interval]
_____
    poor | .137645 .0046531 .1285235 .1467665
. svy: mean poor if year == 1991
(running mean on estimation sample)
Survey: Mean estimation
Number of strata =
               1
                         Number of obs = 6468
Number of PSUs = 5254
                         Population size = 5.6e+07
                         Design df
                                   = 5253
_____
              Linearized
        | Mean Std. Err. [95% Conf. Interval]
_____
    poor | .2021312 .0062077 .1899615 .2143009
..using R code:
headcount_81 <-
  svyfgt(
     ~ eybhc0 ,
     subset( z , year == 1981 ) ,
     g = 0,
     abs_thresh = unwtd_arpt81
  )
headcount 81
##
         fgt0
               SE
## eybhc0 0.14101 0.0045
confint( headcount_81 , df = degf( subset( z , year == 1981 ) ) )
         2.5 % 97.5 %
## eybhc0 0.1322193 0.1498057
```

```
headcount_85 <-
    svyfgt(
        ~ eybhc0 ,
        subset(z, year == 1985),
        g = 0,
        abs_thresh = unwtd_arpt85
   )
headcount_85
             fgt0
## eybhc0 0.13764 0.0047
confint( headcount_85 , df = degf( subset( z , year == 1985 ) ) )
##
              2.5 %
                       97.5 %
## eybhc0 0.1285239 0.1467661
headcount_91 <-
    svyfgt(
        ~ eybhc0 ,
        subset( z , year == 1991 ) ,
        g = 0,
        abs_thresh = unwtd_arpt91
   )
headcount_91
##
             fgt0
                      SE
## eybhc0 0.20213 0.0062
confint( headcount_91 , df = degf( subset( z , year == 1991 ) ) )
              2.5 % 97.5 %
## eybhc0 0.1899624 0.2143
Confirm this replication applies for the normalized poverty gap as well, compar-
ing stata code..
. ge ngap = poor*($z_81- x)/$z_81 if year == 1981
. svy: mean ngap if year == 1981
(running mean on estimation sample)
Survey: Mean estimation
Number of strata =
                         1
                                    Number of obs
                                                           9772
                                    Population size = 5.5e+07
Number of PSUs =
                      7476
                                    Design df
                                                           7475
```

```
| Linearized | Mean Std. Err. [95% Conf. Interval] | ngap | .0271577 .0013502 .0245109 .0298044
```

..to R code:

```
## fgt1 SE
## eybhc0 0.027158 0.0014

confint( norm_pov_81 , df = degf( subset( z , year == 1981 ) ) )
## 2.5 % 97.5 %
## eybhc0 0.02451106 0.02980428
```

For additional usage examples of svyfgt, type ?convey::svyfgt in the R console.

2.7 Watts poverty measure (svywatts, svywatts-dec)

The measure proposed in Watts (1968) satisfies a number of desirable poverty measurement axioms and is known to be one of the first distribution-sensitive poverty measures, as noted by Haughton and Khandker (2009). It is defined as

$$Watts = \frac{1}{N} \sum_{i \in U} \log \left(\frac{y_i}{\theta} \right) \delta(y_i \leqslant \theta).$$

Morduch (1998) points out that the Watts poverty index can provide an estimate of the expected time to exit poverty. Given the expected growth rate of income per capita among the poor, g, the expected time taken to exit poverty T_{θ} would be

$$T_{\theta} = \frac{Watts}{g}.$$

The Watts poverty index also has interesting decomposition properties. Blackburn (1989) proposed a decomposition for the Watts poverty index, rewriting it in terms of the headcount ratio, the Watts poverty gap ratio and the mean log deviation of poor incomes². Mathematically,

$$Watts = FGT_0(I_w + L_*)$$

where $I_w = \log(\theta/\mu_*)$ is the Watts poverty gap ratio³ and L_* is the mean log deviation of incomes among the poor. This can be estimated using the svywattsdec function.

This result can also be interpreted as a decomposition of the time taken to exit poverty, since

$$\begin{split} T_{\theta} &= \frac{Watts}{g} \\ &= \frac{FGT_0}{g} (I_w + L_*) \end{split}$$

As Morduch (1998) points out, if the income among the poor is equally distributed (i.e., $L_*=0$), the time taken to exit poverty is simply FGT_0I_w/g . Therefore, FGT_0L_*/g can be seen as the additional time needed to exit poverty as a result of the inequality among the poor.

2.8 Clark-Hemming-Ulph class of poverty measures (svychu)

Clark et al. (1981) proposes two classes of distribution-sensitive poverty measures. Yet, the poverty measurement literature focuses on the second class⁴, expressed as

$$CHU_{\alpha} = \begin{cases} \frac{1}{\alpha N} \sum_{i \in U} \left[1 - (y_i/\theta)^{\alpha} \right] \cdot \delta(y_i \leqslant \theta), & \alpha \leqslant 1, \alpha \neq 0 \\ 1 - \left(\prod_{i \in U} y_i^{\delta(y_i \leqslant \theta)} \right)^{1/N} \middle/ \theta, & \alpha = 0 \end{cases}$$

²The mean log deviation (also known as Theil-L or Bourguignon-Theil index) is an inequality measure of the generalized entropy class. The family of generalized entropy indices is discussed in the next chapter.

 $^{^3\}mu_*$ stands for the average income among the poor.

⁴See Atkinson (1987) and Verma and Betti (2011), for instance.

As an special case, $CHU_0=1-\exp{(-Watts)}$. The α parameter defines the sensivity towards regressive income transfers among the poor, such that the lower its value, larger is the regressive transfer impact on the index. When $\alpha \to 1$, $CHU_1 = FGT_0 \cdot I$, a measure insensitive to regressive income transfers among the poor.

Chapter 3

Inequality Measurement

Another problem faced by societies is inequality. Economic inequality can have several different meanings: income, education, resources, opportunities, wellbeing, etc. Usually, studies on economic inequality focus on income distribution.

Most inequality data comes from censuses and household surveys. Therefore, in order to produce reliable estimates from this samples, appropriate procedures are necessary.

This chapter presents brief presentations on inequality measures, also providing replication examples if possible. It starts with an initial attempt to measure the inequality between two groups of a population; then, it presents ideas of overall inequality indices, moving from the quintile share ratio to the Lorenz curve and measures derived from it; then, it discusses the concept of entropy and presents inequality measures based on it. Finally, it ends with a discussion regarding which inequality measure should be used.

3.1 The Gender Pay Gap (svygpg)

Although the GPG is not an inequality measure in the usual sense, it can still be an useful instrument to evaluate the discrimination among men and women. Put simply, it expresses the relative difference between the average hourly earnings of men and women, presenting it as a percentage of the average of hourly earnings of men.

In mathematical terms, this index can be described as,

$$GPG = \frac{\bar{y}_{male} - \bar{y}_{female}}{\bar{y}_{male}}$$

,

which is precisely the estimator used in the package. As we can see from the formula, if there is no difference among classes, GPG = 0. Else, if GPG > 0, it means that the average hourly income received by women are GPG percent smaller than men's. For negative GPG, it means that women's hourly earnings are GPG percent larger than men's. In other words, the larger the GPG, larger is the shortfall of women's hourly earnings.

We can also develop a more straightforward idea: for every \$1 raise in men's hourly earnings, women's hourly earnings are expected to increase \$(1-GPG). For instance, assuming GPG=0.8, for every \$1.00 increase in men's average hourly earnings, women's hourly earnings would increase only \$0.20.

The details of the linearization of the GPG are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a gpg coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the laeken library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# coerce the gender variable to numeric 1 or 2
eusilc$one two <- as.numeric( eusilc$rb090 == "female" ) + 1</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
```

```
# calculate the gpg coefficient
# using the R vardpoor library
varpoord_gpg_calculation <-</pre>
   varpoord(
        # analysis variable
        Y = "eqincome",
        # weights variable
        w_{final} = "rb050",
        # row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # gpg coefficient function
        type = "lingpg" ,
        # gender variable
        gender = "one_two",
      # poverty threshold range
      order_quant = 50L ,
      # get linearized variable
      outp_lin = TRUE
# construct a survey.design
# using our recommended setup
```

```
des_eusilc <-
    svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050 ,
        data = eusilc
    )
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_gpg_calculation$all_result$value
## [1] 7.645389
coef( svygpg( ~ eqincome , des_eusilc , sex = ~ rb090 ) ) * 100
## eqincome
## 7.645389
# linearized variables do match
# vardpoor
lin_gpg_varpoord<- varpoord_gpg_calculation$lin_out$lin_gpg</pre>
# convey
lin_gpg_convey <- attr(svygpg( ~ eqincome , des_eusilc, sex = ~ rb090 ),"lin")</pre>
# check equality
all.equal(lin_gpg_varpoord,100*lin_gpg_convey[,1] )
## [1] TRUE
# variances do not match exactly
attr( svygpg( ~ eqincome , des_eusilc , sex = ~ rb090 ) , 'var' ) * 10000
##
             eqincome
## eqincome 0.6493911
varpoord_gpg_calculation$all_result$var
## [1] 0.6482346
# standard errors do not match exactly
varpoord_gpg_calculation$all_result$se
## [1] 0.8051301
SE( svygpg( ~ eqincome , des_eusilc , sex = ~ rb090 ) ) * 100
##
             eqincome
```

```
## eqincome 0.8058481
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions convey::svygpg and vardpoor::lingpg produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )</pre>
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
attr( svygpg( ~ eqincome , des_eusilc_ultimate_cluster , sex = ~ rb090 ) , 'var' ) * 10000
             eqincome
## eqincome 0.6482346
varpoord_gpg_calculation$all_result$var
## [1] 0.6482346
# matches
varpoord_gpg_calculation$all_result$se
## [1] 0.8051301
```

For additional usage examples of svygpg, type ?convey::svygpg in the R console.

3.2 Quintile Share Ratio (svygsr)

Unlike the previous measure, the quintile share ratio is an inequality measure in itself, depending only of the income distribution to evaluate the degree of inequality. By definition, it can be described as the ratio between the income share held by the richest 20% and the poorest 20% of the population.

In plain terms, it expresses how many times the wealthier part of the population are richer than the poorest part. For instance, a QSR=4 implies that the upper class owns 4 times as much of the total income as the poor.

The quintile share ratio can be modified to a more general function of fractile share ratios. For instance, Cobham et al. (2015) presents interesting arguments for using the Palma index, defined as the ratio between the share of the 10% richest over the share held by the poorest 40%.

The details of the linearization of the QSR are discussed by Deville (1999) and Osier (2009).

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a qsr coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)

# load the survey library
library(survey)

# load the vardpoor library
library(vardpoor)

# load the laeken library
library(laeken)
```

```
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the qsr coefficient
# using the R vardpoor library
varpoord_qsr_calculation <-</pre>
    varpoord(
        # analysis variable
        Y = "eqincome",
        # weights variable
        w_{final} = "rb050",
        # row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # qsr coefficient function
        type = "linqsr",
      # poverty threshold range
      order_quant = 50L ,
      # get linearized variable
      outp_lin = TRUE
```

```
)
# construct a survey.design
# using our recommended setup
des_eusilc <-
    svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc
    )
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_qsr_calculation$all_result$value
## [1] 3.970004
coef( svyqsr( ~ eqincome , des_eusilc ) )
## eqincome
## 3.970004
# linearized variables do match
# vardpoor
lin_qsr_varpoord<- varpoord_qsr_calculation$lin_out$lin_qsr</pre>
# convey
lin_qsr_convey <- attr(svyqsr( ~ eqincome , des_eusilc ),"lin")</pre>
# check equality
all.equal(lin_qsr_varpoord, lin_qsr_convey )
## [1] TRUE
# variances do not match exactly
attr( svyqsr( ~ eqincome , des_eusilc ) , 'var' )
##
               eqincome
## eqincome 0.001810537
varpoord_qsr_calculation$all_result$var
## [1] 0.001807323
```

```
# standard errors do not match exactly
varpoord_qsr_calculation$all_result$se

## [1] 0.04251263

SE( svyqsr( ~ eqincome , des_eusilc ) )

## eqincome
## eqincome
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions convey::svygpg and vardpoor::lingpg produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )</pre>
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = ~rb030,
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svyqsr( ~ eqincome , des_eusilc_ultimate_cluster ) , 'var' )
                eqincome
## eqincome 0.001807323
```

```
varpoord_qsr_calculation$all_result$var
```

```
## [1] 0.001807323
```

```
# matches
```

varpoord_qsr_calculation\$all_result\$se

```
## [1] 0.04251263
```

```
SE( svyqsr( ~ eqincome , des_eusilc_ultimate_cluster ) )
```

```
## eqincome
## eqincome 0.04251263
```

For additional usage examples of svyqsr, type ?convey::svyqsr in the R console.

3.3 Lorenz Curve (svylorenz)

Though not an inequality measure in itself, the Lorenz curve is a classic instrument of distribution analysis. Basically, it is a function that associates a cumulative share of the population to the share of the total income it owns. In mathematical terms,

$$L(p) = \frac{\int_{-\infty}^{Q_p} y f(y) dy}{\int_{-\infty}^{+\infty} y f(y) dy}$$

where Q_p is the quantile p of the population.

The two extreme distributive cases are

- Perfect equality:
 - Every individual has the same income;
 - Every share of the population has the same share of the income;
 - Therefore, the reference curve is

$$L(p) = p \ \forall p \in [0, 1].$$

- Perfect inequality:
 - One individual concentrates all of society's income, while the other individuals have zero income;
 - Therefore, the reference curve is

$$L(p) = \begin{cases} 0, & \forall p < 1 \\ 1, & \text{if } p = 1. \end{cases}$$

In order to evaluate the degree of inequality in a society, the analyst looks at the distance between the real curve and those two reference curves.

The estimator of this function was derived by Kovacevic and Binder (1997):

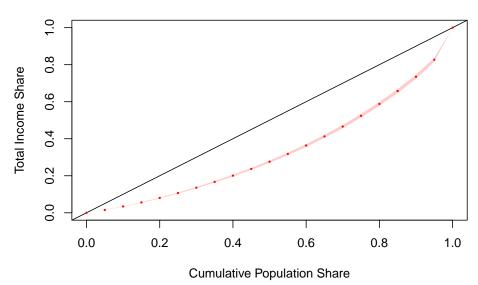
$$L(p) = \frac{\sum_{i \in S} w_i \cdot y_i \cdot \delta\{y_i \leq \widehat{Q}_p\}}{\widehat{Y}}, \ 0 \leq p \leq 1.$$

Yet, this formula is used to calculate specific points of the curve and their respective SEs. The formula to plot an approximation of the continuous empirical curve comes from Lerman and Yitzhaki (1989).

A replication example

In October 2016, (Jann, 2016) released a pre-publication working paper to estimate lorenz and concentration curves using stata. The example below reproduces the statistics presented in his section 4.1.

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the stata-style webuse library
library(webuse)
# load the NLSW 1988 data
webuse("nlsw88")
# coerce that `tbl df` to a standard R `data.frame`
nlsw88 <- data.frame( nlsw88 )</pre>
# initiate a linearized survey design object
des nlsw88 <- svydesign( ids = ~1 , data = nlsw88 )
## Warning in svydesign.default(ids = ~1, data = nlsw88): No weights or
## probabilities supplied, assuming equal probability
# immediately run the `convey_prep` function on the survey design
des_nlsw88 <- convey_prep(des_nlsw88)</pre>
# estimates lorenz curve
result.lin <- svylorenz( ~wage, des_nlsw88, quantiles = seq( 0, 1, .05 ), na.rm = TRUE )
```



```
\# note: most survey commands in R use Inf degrees of freedom by default
# stata generally uses the degrees of freedom of the survey design.
# therefore, while this extended syntax serves to prove a precise replication of stata
# it is generally not necessary.
section_four_one <-
    data.frame(
        estimate = coef( result.lin ) ,
        standard_error = SE( result.lin ) ,
        ci_lower_bound =
            coef( result.lin ) +
            SE( result.lin ) *
            qt( 0.025 , degf( subset( des_nlsw88 , !is.na( wage ) ) ) ) ,
        ci_upper_bound =
            coef( result.lin ) +
            SE( result.lin ) *
            qt( 0.975 , degf( subset( des_nlsw88 , !is.na( wage ) ) ) )
```

	estimate	standard_error	ci_lower_bound	ci_upper_bound
0	0.0000000	0.0000000	0.0000000	0.0000000
0.05	0.0151060	0.0004159	0.0142904	0.0159216
0.1	0.0342651	0.0007021	0.0328882	0.0356420
0.15	0.0558635	0.0010096	0.0538836	0.0578434
0.2	0.0801846	0.0014032	0.0774329	0.0829363
0.25	0.1067687	0.0017315	0.1033732	0.1101642
0.3	0.1356307	0.0021301	0.1314535	0.1398078
0.35	0.1670287	0.0025182	0.1620903	0.1719670
0.4	0.2005501	0.0029161	0.1948315	0.2062687
0.45	0.2369209	0.0033267	0.2303971	0.2434447
0.5	0.2759734	0.0037423	0.2686347	0.2833121
0.55	0.3180215	0.0041626	0.3098585	0.3261844
0.6	0.3633071	0.0045833	0.3543192	0.3722950
0.65	0.4125183	0.0050056	0.4027021	0.4223345
0.7	0.4657641	0.0054137	0.4551478	0.4763804
0.75	0.5241784	0.0058003	0.5128039	0.5355529
0.8	0.5880894	0.0062464	0.5758401	0.6003388
0.85	0.6577051	0.0066148	0.6447333	0.6706769
0.9	0.7346412	0.0068289	0.7212497	0.7480328
0.95	0.8265786	0.0062686	0.8142857	0.8388715
1	1.0000000	0.0000000	1.0000000	1.0000000

For additional usage examples of svylorenz, type ?convey::svylorenz in the R console.

3.4 Gini index (svygini)

The Gini index is an attempt to express the inequality presented in the Lorenz curve as a single number. In essence, it is twice the area between the equality curve and the real Lorenz curve. Put simply:

$$G = 2 \bigg(\int_0^1 p dp - \int_0^1 L(p) dp \bigg)$$

$$\therefore G = 1 - 2 \int_0^1 L(p) dp$$

where G=0 in case of perfect equality and G=1 in the case of perfect inequality.

The estimator proposed by Osier (2009) is defined as:

$$\widehat{G} = \frac{2\sum_{i \in S} w_i r_i y_i - \sum_{i \in S} w_i y_i}{\widehat{Y}}$$

The linearized formula of \widehat{G} is used to calculate the SE.

A replication example

The R vardpoor package (Breidaks et al., 2016), created by researchers at the Central Statistical Bureau of Latvia, includes a gini coefficient calculation using the ultimate cluster method. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the vardpoor library
library(vardpoor)
# load the laeken library
library(laeken)
# load the synthetic EU statistics on income & living conditions
data(eusilc)
# make all column names lowercase
names( eusilc ) <- tolower( names( eusilc ) )</pre>
# add a column with the row number
dati <- data.table::data.table(IDd = 1 : nrow(eusilc), eusilc)</pre>
# calculate the gini coefficient
# using the R vardpoor library
varpoord_gini_calculation <-</pre>
    varpoord(
        # analysis variable
        Y = "eqincome",
        # weights variable
        w_final = "rb050",
```

```
# row number variable
        ID_level1 = "IDd",
        # row number variable
        ID_level2 = "IDd",
        # strata variable
        H = "db040",
        N_h = NULL,
        # clustering variable
        PSU = "rb030",
        # data.table
        dataset = dati,
        # qini coefficient function
        type = "lingini",
      # poverty threshold range
      order_quant = 50L ,
      # get linearized variable
      outp_lin = TRUE
   )
# construct a survey.design
# using our recommended setup
des_eusilc <-
   svydesign(
        ids = ~rb030,
        strata = -db040,
        weights = \sim rb050 ,
        data = eusilc
   )
# immediately run the convey_prep function on it
des_eusilc <- convey_prep( des_eusilc )</pre>
# coefficients do match
varpoord_gini_calculation$all_result$value
```

```
## [1] 26.49652
coef( svygini( ~ eqincome , des_eusilc ) ) * 100
## eqincome
## 26.49652
# linearized variables do match
# varpoord
lin_gini_varpoord<- varpoord_gini_calculation$lin_out$lin_gini</pre>
# convey
lin_gini_convey <- attr(svygini( ~ eqincome , des_eusilc ),"lin")</pre>
# check equality
all.equal(lin_gini_varpoord,100*lin_gini_convey)
## [1] TRUE
# variances do not match exactly
attr( svygini( ~ eqincome , des_eusilc ) , 'var' ) * 10000
              eqincome
## eqincome 0.03790739
varpoord_gini_calculation$all_result$var
## [1] 0.03783931
# standard errors do not match exactly
varpoord_gini_calculation$all_result$se
## [1] 0.1945233
SE( svygini( ~ eqincome , des_eusilc ) ) * 100
##
             eqincome
## eqincome 0.1946982
```

The variance estimate is computed by using the approximation defined in (1.1), where the linearized variable z is defined by (1.2). The functions convey::svygini and vardpoor::lingini produce the same linearized variable z.

However, the measures of uncertainty do not line up, because library(vardpoor) defaults to an ultimate cluster method that can be replicated with an alternative setup of the survey.design object.

```
# within each strata, sum up the weights
cluster_sums <- aggregate( eusilc$rb050 , list( eusilc$db040 ) , sum )
# name the within-strata sums of weights the `cluster_sum`
names( cluster_sums ) <- c( "db040" , "cluster_sum" )</pre>
```

```
# merge this column back onto the data.frame
eusilc <- merge( eusilc , cluster_sums )</pre>
# construct a survey.design
# with the fpc using the cluster sum
des_eusilc_ultimate_cluster <-</pre>
    svydesign(
        ids = "rb030",
        strata = ~db040,
        weights = \sim rb050,
        data = eusilc ,
        fpc = ~ cluster_sum
    )
# again, immediately run the convey_prep function on the `survey.design`
des_eusilc_ultimate_cluster <- convey_prep( des_eusilc_ultimate_cluster )</pre>
# matches
attr( svygini( ~ eqincome , des_eusilc_ultimate_cluster ) , 'var' ) * 10000
              eqincome
## eqincome 0.03783931
varpoord_gini_calculation$all_result$var
## [1] 0.03783931
# matches
varpoord_gini_calculation$all_result$se
## [1] 0.1945233
SE( svygini( ~ eqincome , des_eusilc_ultimate_cluster ) ) * 100
             eqincome
## eqincome 0.1945233
For additional usage examples of svygini, type ?convey::svygini in the R
```

3.5 Amato index (svyamato)

console.

The Amato index is also based on the Lorenz curve, but instead of focusing on the area of the curve, it focuses on its length. Arnold (2012) proposes a formula not directly based in the Lorenz curve, which Barabesi et al. (2016) uses to present the following estimator:

$$\widehat{A} = \sum_{i \in S} w_i \left[\frac{1}{\widehat{N}^2} + \frac{y_i^2}{\widehat{Y}^2} \right]^{\frac{1}{2}},$$

which also generates the linearized formula for SE estimation.

The minimum value A assumes is $\sqrt{2}$ and the maximum is 2. In order to get a measure in the interval [0,1], the standardized Amato index \widetilde{A} can be defined as:

$$\widetilde{A} = \frac{A - \sqrt{2}}{2 - \sqrt{2}} \ .$$

For additional usage examples of svyamato, type ?convey::svyamato in the R console.

3.6 Zenga Index and Curve (svyzenga, svyzengacurve)

The Zenga index and its curve were proposed in Zenga (2007). As Polisicchio and Porro (2011) noticed, this curve derives directly from the Lorenz curve, and can be defined as:

$$Z(p) = 1 - \frac{L(p)}{p} \cdot \frac{1-p}{1-L(p)}.$$

In the **convey** library, an experimental estimator based on the Lorenz curve is used:

$$\widehat{Z(p)} = \frac{p\widehat{Y} - \widehat{\widetilde{Y}}(p)}{p[\widehat{Y} - \widehat{\widetilde{Y}}(p)]}.$$

In turn, the Zenga index derives from this curve and is defined as:

$$Z = \int_0^1 Z(p)dp.$$

However, its estimators were proposed by Langel (2012) and Barabesi et al. (2016). In this library, the latter is used and is defined as:

$$\widehat{Z} = 1 - \sum_{i \in S} w_i \bigg[\frac{(\widehat{N} - \widehat{H}_{y_i})(\widehat{Y} - \widehat{K}_{y_i})}{\widehat{N} \cdot \widehat{H}_{y_i} \cdot \widehat{K}_{y_i}} \bigg]$$

where \widehat{N} is the population total, \widehat{Y} is the total income, \widehat{H}_{y_i} is the sum of incomes below or equal to y_i and \widehat{N}_{y_i} is the sum of incomes greater or equal to y_i .

For additional usage examples of svyzenga or svyzengacurve, type ?convey::svyzenga or ?convey::svyzengacurve in the R console.

3.7 Entropy-based Measures

Entropy is a concept derived from information theory, meaning the expected amount of information given the occurrence of an event. Following (Shannon, 1948), given an event y with probability density function $f(\cdot)$, the information content given the occurrence of y can be defined as $g(f(y)) := -\log f(y)$. Therefore, the expected information or, put simply, the *entropy* is

$$H(f) \colon = -E\big[\log f(y)\big] = -\int_{-\infty}^{\infty} f(y) \log f(y) dy$$

Assuming a discrete distribution, with p_k as the probability of occurring event $k \in K$, the entropy formula takes the form:

$$H = -\sum_{k \in K} p_k \log p_k.$$

The main idea behind it is that the expected amount of information of an event is inversely proportional to the probability of its occurrence. In other words, the information derived from the observation of a rare event is higher than of the information of more probable events.

Using ideas presented in Cowell et al. (2009), substituting the density function by the income share of an individual $s(q) = F^{-1}(q)/\int_0^1 F^{-1}(t)dt = y/\mu$, the entropy function becomes the Theil¹ inequality index

$$I_{Theil} = \int_{0}^{\infty} \frac{y}{\mu} \log \left(\frac{y}{\mu} \right) dF(y) = -H(s)$$

Therefore, the entropy-based inequality measure increases as a person's income y deviates from the mean μ . This is the basic idea behind entropy-based inequality measures.

¹Also known as Theil-T index.

3.8 Generalized Entropy and Decomposition (svygei, svygeidec)

Using a generalization of the information function, now defined as $g(f) = \frac{1}{\alpha-1}[1-f^{\alpha-1}]$, the α -class entropy is

$$H_{\alpha}(f) = \frac{1}{\alpha-1} \bigg[1 - \int_{-\infty}^{\infty} f(y)^{\alpha-1} f(y) dy \bigg].$$

This relates to a class of inequality measures, the Generalized entropy indices, defined as:

$$GE_{\alpha} = \frac{1}{\alpha^2 - \alpha} \int_0^{\infty} \bigg[\bigg(\frac{y}{\mu} \bigg)^{\alpha} - 1 \bigg] dF(x) = -\frac{-H_{\alpha}(s)}{\alpha}.$$

The parameter α also has an economic interpretation: as α increases, the influence of top incomes upon the index increases. In some cases, this measure takes special forms, such as mean log deviation and the aforementioned Theil index.

In order to estimate it, Biewen and Jenkins (2003) proposed the following:

$$GE_{\alpha} = \begin{cases} (\alpha^2 - \alpha)^{-1} \big[U_0^{\alpha - 1} U_1^{-\alpha} U_{\alpha} - 1 \big], & \text{if } \alpha \in \mathbb{R} \quad \{0, 1\} \\ -T_0 U_0^{-1} + \log(U_1/U_0), & \text{if } \alpha \to 0 \\ T_1 U_1^{-1} - \log(U_1/U_0), & \text{if } \alpha \to 1 \end{cases}$$

where $U_{\gamma} = \sum_{i \in S} w_i \cdot y_i^{\gamma}$ and $T_{\gamma} = \sum_{i \in S} w_i \cdot y_i^{\gamma} \cdot \log y_i$. Since those are all functions of totals, the linearization of the indices are easily achieved using the theorems described in Deville (1999).

This class also has several desirable properties, such as additive decomposition. The additive decomposition allows to compare the effects of inequality within and between population groups on the population inequality. Put simply, an additive decomposable index allows for:

$$I_{Total} = I_{Between} + I_{Within}. \label{eq:total_total}$$

A replication example

In July 2006, Jenkins (2008) presented at the North American Stata Users' Group Meetings on the stata Generalized Entropy Index command. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the foreign library
library(foreign)
# create a temporary file on the local disk
tf <- tempfile()</pre>
# store the location of the presentation file
presentation_zip <- "http://repec.org/nasug2006/nasug2006_jenkins.zip"</pre>
# download jenkins' presentation to the temporary file
download.file( presentation_zip , tf , mode = 'wb' )
# unzip the contents of the archive
presentation_files <- unzip( tf , exdir = tempdir() )</pre>
# load the institute for fiscal studies' 1981, 1985, and 1991 data.frame objects
x81 <- read.dta( grep( "ifs81" , presentation_files , value = TRUE ) )</pre>
x85 <- read.dta( grep( "ifs85" , presentation_files , value = TRUE ) )</pre>
x91 <- read.dta( grep( "ifs91" , presentation_files , value = TRUE ) )</pre>
# stack each of these three years of data into a single data.frame
x <- rbind( x81 , x85 , x91 )
Replicate the author's survey design statement from stata code..
. * account for clustering within HHs
. version 8: svyset [pweight = wgt], psu(hrn)
pweight is wgt
psu is hrn
construct an
.. into R code:
# initiate a linearized survey design object
y <- svydesign( ~ hrn , data = x , weights = ~ wgt )
# immediately run the `convey_prep` function on the survey design
z <- convey_prep( y )</pre>
```

Replicate the author's subset statement and each of his svygei results..

```
. svygei x if year == 1981
```

Warning: x has 20 values = 0. Not used in calculations

Complex survey estimates of Generalized Entropy inequality indices

```
pweight: wgt
Strata: <one>
Number of obs = 9752
Number of strata = 1
PSU: hrn
Number of PSUs = 7459
Population size = 54766261
```

Index Estimate Std. Err. z P> z [95% Conf. Inter	
GE(-1) .1902062 .02474921 7.69 0.000 .1416987 .238 MLD .1142851 .00275138 41.54 0.000 .1088925 .119 Theil .1116923 .00226489 49.31 0.000 .1072532 .116 GE(2) .128793 .00330774 38.94 0.000 .1223099 .13	37138 96777 51314 35276 59747

..using R code:

```
z81 \leftarrow subset(z, year == 1981)
svygei( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = -1 )
              gei
## eybhc0 0.19021 0.0247
svygei( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 0 )
##
              gei
## eybhc0 0.11429 0.0028
svygei( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) )
              gei
## eybhc0 0.11169 0.0023
svygei( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 2 )
##
              gei
                      SE
## eybhc0 0.12879 0.0033
svygei( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 3 )
            gei
## eybhc0 0.174 0.0066
```

Confirm this replication applies for subsetted objects as well. Compare stata output..

```
. svygei x if year == 1985 & x >= 1
```

Complex survey estimates of Generalized Entropy inequality indices

GE(-1) .1602358 .00936931 17.10 0.000 .1418723 .1785993 MLD .127616 .00332187 38.42 0.000 .1211052 .1341267 Theil .1337177 .00406302 32.91 0.000 .1257543 .141681 GE(2) .1676393 .00730057 22.96 0.000 .1533304 .1819481	Index		Estimate	Std. Err.	z	P> z	[95% Conf.	Interval]
	GE(-1) MLD Theil GE(2)	 	.1602358 .127616 .1337177 .1676393	.00936931 .00332187 .00406302 .00730057	17.10 38.42 32.91 22.96	0.000 0.000 0.000 0.000	.1418723 .1211052 .1257543 .1533304	.1785993 .1341267 .141681 .1819481 .2972235

..to R code:

```
z85 \leftarrow subset(z, year == 1985)
svygei( ~ eybhc0 , subset( z85 , eybhc0 > 1 ) , epsilon = -1 )
##
              gei
## eybhc0 0.16024 0.0094
svygei( ~ eybhc0 , subset( z85 , eybhc0 > 1 ) , epsilon = 0 )
              gei
## eybhc0 0.12762 0.0033
svygei( ~ eybhc0 , subset( z85 , eybhc0 > 1 ) )
##
                      SE
              gei
## eybhc0 0.13372 0.0041
svygei( ~ eybhc0 , subset( z85 , eybhc0 > 1 ) , epsilon = 2 )
              gei
## eybhc0 0.16764 0.0073
svygei( ~ eybhc0 , subset( z85 , eybhc0 > 1 ) , epsilon = 3 )
##
              gei
## eybhc0 0.26095 0.0185
```

Replicate the author's decomposition by population subgroup (work status) shown on PDF page 57..

```
# define work status (PDF page 22)
z <- update( z , wkstatus = c( 1 , 1 , 1 , 1 , 2 , 3 , 2 , 2 )[ as.numeric( esbu ) ] )</pre>
z <- update( z , factor( wkstatus , labels = c( "1+ ft working" , "no ft working" , "e</pre>
# subset to 1991 and remove records with zero income
z91 <- subset( z , year == 1991 & eybhc0 > 0 )
# population share
svymean( ~wkstatus, z91 )
##
             mean
## wkstatus 1.5594 0.0099
# mean
svyby( ~eybhc0, ~wkstatus, z91, svymean )
##
    wkstatus
              eybhc0
                             se
## 1
           1 278.8040 3.703790
## 2
            2 151.6317 3.153968
## 3
            3 176.6045 4.661740
# subgroup indices: ge_k
svyby( ~ eybhc0 , ~wkstatus , z91 , svygei , epsilon = -1 )
    wkstatus
##
                  eybhc0
## 1
           1 0.2300708 0.02853959
## 2
            2 10.9231761 10.65482557
           3 0.1932164 0.02571991
svyby( ~ eybhc0 , ~wkstatus , z91 , svygei , epsilon = 0 )
##
    wkstatus
                 eybhc0
## 1
           1 0.1536921 0.006955506
## 2
            2 0.1836835 0.014740510
            3 0.1653658 0.016409770
svyby( ~ eybhc0 , ~wkstatus , z91 , svygei , epsilon = 1 )
##
    wkstatus
                 eybhc0
## 1
           1 0.1598558 0.008327994
## 2
            2 0.1889909 0.016766120
            3 0.2023862 0.027787224
svyby( ~ eybhc0 , ~wkstatus , z91 , svygei , epsilon = 2 )
##
    wkstatus
                 eybhc0
## 1
           1 0.2130664 0.01546521
## 2
           2 0.2846345 0.06016394
## 3
           3 0.3465088 0.07362898
```

```
# GE decomposition
svygeidec( ~eybhc0, ~wkstatus, z91, epsilon = -1 )
         total within between
## coef 3.6829 3.6466 0.0363
       3.3999 3.3993 0.0541
svygeidec( ~eybhc0, ~wkstatus, z91, epsilon = 0 )
##
            total
                     within between
## coef 0.1952363 0.1619352 0.0333
       0.0064615 0.0062209 0.0027
svygeidec( ~eybhc0, ~wkstatus, z91, epsilon = 1 )
##
            total
                     within between
## coef 0.2003897 0.1693958 0.0310
       0.0079299 0.0044132 0.0073
svygeidec( ~eybhc0, ~wkstatus, z91, epsilon = 2 )
                   within between
          total
## coef 0.274325 0.245067 0.0293
        0.016694 0.017831 0.0038
```

For additional usage examples of svygei or svygeidec, type ?convey::svygei or ?convey::svygeidec in the R console.

3.9 Rényi Divergence (svyrenyi)

Another measure used in areas like ecology, statistics and information theory is the Rényi divergence measure. Using the formula defined in Langel (2012), the estimator can be defined as:

$$\widehat{R}_{\alpha} = \begin{cases} \frac{1}{\alpha - 1} \log \left[\widehat{N}^{\alpha - 1} \sum_{i \in S} w_i \cdot \left(\frac{y_k}{\widehat{Y}} \right) \right], & \text{if } \alpha \neq 1, \\ \sum_{i \in S} \frac{w_i y_i}{\widehat{Y}} \log \frac{\widehat{N} y_i}{\widehat{Y}}, & \text{if } \alpha = 1, \end{cases}$$

where α is a parameter with a similar economic interpretation to that of the GE_{α} index.

For additional usage examples of svyrenyi, type ?convey::svyrenyi in the R console.

3.10 J-Divergence and Decomposition (svyjdiv, svyjdivdec)

The J-divergence measure (Rohde, 2016) can be seen as the sum of GE_0 and GE_1 , satisfying axioms that, individually, those two indices do not. Using U_{γ} and T_{γ} functions defined in Biewen and Jenkins (2003), the estimator can be defined as:

$$\begin{split} J &= \frac{1}{N} \sum_{i \in S} \left(\frac{y_i - \mu}{\mu} \right) \log \left(\frac{y_i}{\mu} \right) \\ &:: \widehat{J} = \frac{\widehat{T}_1}{\widehat{U}_1} - \frac{\widehat{T}_0}{\widehat{U}_0} \end{split}$$

Since it is a sum of two additive decomposable measures, J itself is decomposable.

For additional usage examples of svyjdiv or svyjdivdec, type ?convey::svyjdiv or ?convey::svyjdivdec in the R console.

3.11 Atkinson index (svyatk)

Although the original formula was proposed in Atkinson (1970), the estimator used here comes from Biewen and Jenkins (2003):

$$\widehat{A}_{\epsilon} = \begin{cases} 1 - \widehat{U}_0^{-\epsilon/(1-\epsilon)} \widehat{U}_1^{-1} \widehat{U}_{1-\epsilon}^{1/(1-\epsilon)}, & \text{if } \epsilon \in \mathbb{R}_+ \\ 1 - \widehat{U}_0 \widehat{U}_0^{-1} exp(\widehat{T}_0 \widehat{U}_0^{-1}), & \text{if } \epsilon \to 1 \end{cases}$$

The ϵ is an inequality aversion parameter: as it approaches infinity, more weight is given to incomes in bottom of the distribution.

A replication example

In July 2006, Jenkins (2008) presented at the North American Stata Users' Group Meetings on the stata Atkinson Index command. The example below reproduces those statistics.

Load and prepare the same data set:

load the convey package
library(convey)

load the survey library

```
library(survey)
# load the foreign library
library(foreign)
# create a temporary file on the local disk
tf <- tempfile()</pre>
# store the location of the presentation file
presentation_zip <- "http://repec.org/nasug2006/nasug2006_jenkins.zip"</pre>
# download jenkins' presentation to the temporary file
download.file( presentation_zip , tf , mode = 'wb' )
# unzip the contents of the archive
presentation_files <- unzip( tf , exdir = tempdir() )</pre>
# load the institute for fiscal studies' 1981, 1985, and 1991 data.frame objects
x81 <- read.dta( grep( "ifs81" , presentation_files , value = TRUE ) )</pre>
x85 <- read.dta( grep( "ifs85" , presentation_files , value = TRUE ) )</pre>
x91 <- read.dta( grep( "ifs91" , presentation_files , value = TRUE ) )</pre>
# stack each of these three years of data into a single data.frame
x <- rbind( x81 , x85 , x91 )
Replicate the author's survey design statement from stata code..
. * account for clustering within HHs
```

```
. version 8: svyset [pweight = wgt], psu(hrn)
pweight is wgt
psu is hrn
construct an
.. into R code:
# initiate a linearized survey design object
y <- svydesign( ~ hrn , data = x , weights = ~ wgt )
# immediately run the `convey_prep` function on the survey design
z <- convey_prep( y )</pre>
```

Replicate the author's subset statement and each of his svyatk results with stata..

```
. svyatk x if year == 1981
```

Warning: x has 20 values = 0. Not used in calculations

Complex survey estimates of Atkinson inequality indices

pweight: wgt
Strata: <one>
Number of obs = 9752
Number of strata = 1
PSU: hrn
Number of PSUs = 7459
Population size = 54766261

Index		Estimate	Std. Err.	z	P> z	[95% Conf.	Interval]
A(0.5)	į	.0543239	.00107583	50.49	0.000	.0522153	.0564324
A(1) A(1.5)	1	.1079964 .1701794	.00245424	44.00 25.42	0.000	.1031862 .1570588	.1128066
A(2)	i	.2755788	.02597608	10.61	0.000	.2246666	.326491
A(2.5)	-	.4992701	.06754311	7.39	0.000	.366888	.6316522

..using R code:

```
z81 \leftarrow subset(z, year == 1981)
svyatk( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 0.5 )
         atkinson
                      SE
## eybhc0 0.054324 0.0011
svyatk( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) )
##
         atkinson
## eybhc0
            0.108 0.0025
svyatk( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 1.5 )
         atkinson
                      SE
## eybhc0 0.17018 0.0067
svyatk( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 2 )
##
         atkinson
## eybhc0 0.27558 0.026
svyatk( ~ eybhc0 , subset( z81 , eybhc0 > 0 ) , epsilon = 2.5 )
         atkinson
## eybhc0 0.49927 0.0675
```

Confirm this replication applies for subsetted objects as well, comparing stata code..

```
. svyatk x if year == 1981 & x >= 1
```

Complex survey estimates of Atkinson inequality indices

```
pweight: wgt
                                       Number of obs
                                                     = 9748
Strata: <one>
                                       Number of strata = 1
PSU: hrn
                                       Number of PSUs = 7457
                                       Population size = 54744234
______
Index | Estimate Std. Err. z > |z|
                                             [95% Conf. Interval]
A(0.5) | .0540059 .00105011 51.43
                                             .0519477 .0560641
                                     0.000
                                             .1022313 .1109852
A(1) | .1066082 .00223318 47.74 0.000
A(1.5) | .1638299 .00483069 33.91 0.000
                                              .154362 .1732979
A(2) | .2443206 .01425258 17.14 0.000
                                              .2163861 .2722552
A(2.5) | .394787
                  .04155221 9.50
                                     0.000
                                              .3133461
                                                       .4762278
..to R code:
z81_two <- subset( z , year == 1981 & eybhc0 > 1 )
svyatk( ~ eybhc0 , z81_two , epsilon = 0.5 )
        atkinson
                   SE
## eybhc0 0.054006 0.0011
svyatk( ~ eybhc0 , z81_two )
##
        atkinson
                   SE
## eybhc0 0.10661 0.0022
svyatk( ~ eybhc0 , z81_two , epsilon = 1.5 )
##
        atkinson
## eybhc0 0.16383 0.0048
svyatk( ~ eybhc0 , z81_two , epsilon = 2 )
        atkinson
## eybhc0 0.24432 0.0143
svyatk( ~ eybhc0 , z81_two , epsilon = 2.5 )
```

For additional usage examples of svyatk, type ?convey::svyatk in the R console.

##

atkinson

eybhc0 0.39479 0.0416

SE

3.12 Which inequality measure should be used?

The variety of inequality measures begs a question: which inequality measure should be used? In fact, this is a very important question. However, the nature of it is not statistical or mathematical, but ethical. This section aims to clarify and, while not proposing a "perfect measure", to provide the reader with an initial guidance about which measure to use.

The most general way to analyze if one distribution is more equally distributed than another is by the Lorenz curve. When $L_A(p) \ge L_B(p), \forall p \in [0,1]$, it is said that A is more equally distributed than B. Technically, we say that A (Lorenz)dominates B^2 . In this case, all inequality measures that satisfy basic properties³ will agree that A is more equally distributed than B.

When this dominance fails, i.e., when Lorenz curves do cross, Lorenz ordering is impossible. Then, under such circumstances, the choice of which inequality measure to use becomes relevant.

Each inequality measure is a result of a subjective understanding of what is a fair distribution. As Dalton (1920, p.348) puts it, "[...] the economist is primarily interested, not in the distribution of income as such, but in the effects of the distribution of income upon the distribution and total amount of economic welfare, which may be derived from income." The importance of how economic welfare is defined is once again expressed by Atkinson (1970), where an inequality measure is directly derived from a class of welfare functions. Even when a welfare function is not explicit, such as in the Gini index, we must agree that an implicit, subjective judgement of the impact of inequality on social welfare is assumed.

The idea of what is a fair distribution is a matter of Ethics, a discipline within the realm of Philosophy. Yet, as Fleurbaey (1996, Ch.1) proposes, the analyst should match socially supported moral values and theories of justice to the set of technical tools for policy evaluation.

Although this can be a useful principle, a more objective answer is needed. By knowing the nature and properties of inequality measures, the analyst can further reduce the set of applicable inequality measures. For instance, choosing from the properties listed in Cowell (2011, p.74), if we require group-decomposability, scale invariance, population invariance, and that the estimate in [0, 1], we must resort to the Atkinson index.

Even though the discussion can go deep in technical and philosophical aspects, this choice also depends on the public. For example, it would not be surprising if a public official doesn't know the Atkinson index; however, he might know the

 $^{^2}$ Krämer (1998) and Mosler (1994) provide helpful insights to how majorization, Lorenz dominance, and inequality measurement are connected. On the topic of majorization, Hardy et al. (1934) is still the main reference, while Marshall et al. (2011) provide a more modern approach.

³Namely, Schur-convexity, population invariance, and scale invariance.

Gini index. The same goes for publications: journalists have been introduced to the Gini index and can find it easier to compare and, therefore, write about it. Also, we must admit that the Gini index is much more straightforward than any other measure.

In the end, the choice is mostly subjective and there is no consensus of which is the "greatest inequality measure". We must remember that this choice is only problematic if Lorenz curves cross and, in that case, it is not difficult to justify the use of this or that inequality measure.

Chapter 4

Multidimensional Indices

Inequality and poverty can be seen as multidimensional concepts, combining several livelihood characteristics. Usual approaches take into account income, housing, sanitation, etc.

In order to transform these different measures from into meaningful numbers, economic theory builds on the idea of utility functions. Utility is a measure of well-being, assigning a "well-being score" to a vector of characteristics. Depending on the utility function, the analyst may allow for substitutions among characteristics: for instance, someone with a slightly lower income, but with access to sanitation, can have a higher wellbeing than someone with a higher income, but without access to sanitation. This depends on the set of weights given to the set of attributes.

Most measures below follow from this kind of two-step procedure: (1) estimating individual scores from an individual's set of characteristics; then (2) aggregating those individual scores into a single measure for the population.

The following section will present measures of multidimensional poverty and inequality, describing the main aspects of the theory and estimation procedures of each.

4.1 Alkire-Foster Class and Decomposition (svyafc, svyafcdec)

This class of measures are defined in Alkire and Foster (2011), using what is called the "dual cutoff" approach. This method applies a cutoffs to define dimensional deprivations and another cutoff for multidimensional deprivation.

To analyze a population of n individuals across d achievement dimensions, the

first step of the method is applying a FGT-like transformation to each dimension, defined as

$$g_{ij}^{\alpha} = \left(\frac{z_j - x_{ij}}{z_j}\right)^{\alpha}$$

where i is an observation index, j is a dimension index and α is an exponent weighting the deprivation intensity. If $\alpha=0$, then g_{ij}^0 becomes a binary variable, assuming value 1 if person i is deprived in dimension j and 0 otherwise. The $n\times d$ matrix G^{α} will be referred to as deprivation matrix.

Each dimension receives a weight w_j , so that the weighted sum of multidimensional deprivation is the matrix multiplication of G^{α} by the $j \times 1$ vector $W = [w_j]$. The $n \times 1$ vector $C^{\alpha} = [c_i^{\alpha}]$ is the weighted sum of dimensional deprivation scores, i.e.,

$$c_i^{\alpha} = \sum_{j \in d} w_j g_{ij}^{\alpha}$$

The second cutoff is defining those considered to be multidimensionally poor. Assuming that $\sum_{j\in d} w_j = 1$, the multidimensional cutoff k belongs to the interval (0,1]. If $c_i^0 \geqslant k$, then this person is considered multidimensionally poor. The censored vector of deprivation sums $C^{\alpha}(k)$ is defined as

$$C^{\alpha}(k) = \bigg[c_{ij}^{\alpha} \cdot \delta \big(c_{ij}^{0} \geqslant k\big)\bigg],$$

where $\delta(A)$ is an indicator function, taking value 1 if condition A is true and 0 otherwise. If $k \geqslant \min w_j$, this is called the "union approach", where a person is considered poor if she is poor in at least one dimension. On the other extreme, the "intersection approach" happens when k=1, meaning that a person is considered poor if she is poor in all dimensions.

The average of vector $C^0(k)$ returns the multidimensional head count ratio. For the multidimensional FGT class, a general measure can be defined as

$$M^{\alpha} = \frac{1}{n} \sum_{i \in n} \sum_{j \in d} w_j g_{ij}^{\alpha}(k), \ \alpha \ge 0,$$

where $g_{ij}^{\alpha}(k) = g_{ij}^{\alpha} \cdot \delta(c_i^0 \geqslant k)$.

For inferential purposes, since this variable is actually the average of scores $\sum_{j \in d} w_j g_{ij}^{\alpha}(k)$, the linearization is straightforward.

The Alkire-Foster index is both dimensional and subgroup decomposable. This way, it is possible to analyze how much each dimension or group contribute to

the general result. The overall poverty measure can be seen as the weighted sum of each group's poverty measure, as in the formula below:

$$M^{\alpha} = \sum_{l \in L} \frac{n_l}{n} M_l^{\alpha}$$

where l is one of L groups.

Also, the overall poverty index can be expressed across dimensions as

$$M^{\alpha} = \sum_{j \in d} w_j \left[\frac{1}{n} \sum_{i \in n} g_{ij}^{\alpha}(k) \right].$$

Since those functions are linear combinations of ratios and totals, it is also possible to calculate standard errors for such measures.

A replication example

In November 2015, Christopher Jindra presented at the Oxford Poverty and Human Development Initiative on the Alkire-Foster multidimensional poverty measure. His presentation can be viewed here. The example below reproduces those statistics.

Load and prepare the same data set:

```
# load the convey package
library(convey)
# load the survey library
library(survey)
# load the stata-style webuse library
library(webuse)
# load the same microdata set used by Jindra in his presentation
webuse("nlsw88")
# coerce that `tbl_df` to a standard R `data.frame`
nlsw88 <- data.frame( nlsw88 )</pre>
# create a `collgrad` column
nlsw88$collgrad <-
    factor(
        as.numeric( nlsw88$collgrad ) ,
        label = c( 'not college grad' , 'college grad' ) ,
        ordered = TRUE
```

```
# coerce `married` column to factor
nlsw88$married <-
    factor(
        nlsw88$married,
        levels = 0:1,
        labels = c( "single" , "married" )
    )
# initiate a linearized survey design object
des_nlsw88 <- svydesign( ids = ~1 , data = nlsw88 )</pre>
# immediately run the `convey_prep` function on the survey design
des_nlsw88 <- convey_prep(des_nlsw88)</pre>
Replicate PDF page 9
page_nine <-
  svyafc(
    ~ wage + collgrad + hours ,
   design = des_nlsw88 ,
   cutoffs = list( 4, 'college grad' , 26 ) ,
   k = 1/3, g = 0,
    na.rm = TRUE
  )
# MO and seMO
print( page_nine )
        alkire-foster
## [1,]
            0.36991 0.0053
# H seH and A seA
print( attr( page_nine , "extra" ) )
          coef
## H 0.8082070 0.008316807
## A 0.4576895 0.004573443
Replicate PDF page 10
page_ten <- NULL</pre>
# loop through every poverty cutoff `k`
for( ks in seq( 0.1 , 1 , .1 ) ){
   this_ks <-
```

4.1. ALKIRE-FOSTER CLASS AND DECOMPOSITION (SVYAFC, SVYAFCDEC)81

Table 4.1: PDF Page 10 Replication

	k	MO	seMO	Н	seH	A	seA
alkire-foster	0.1	0.3699078	0.0053059	0.8082070	0.0083168	0.4576895	0.0045734
alkire-foster1	0.2	0.3699078	0.0053059	0.8082070	0.0083168	0.4576895	0.0045734
alkire-foster2	0.3	0.3699078	0.0053059	0.8082070	0.0083168	0.4576895	0.0045734
alkire-foster3	0.4	0.1865894	0.0068123	0.2582516	0.0092455	0.7225101	0.0051745
alkire-foster 4	0.5	0.1865894	0.0068123	0.2582516	0.0092455	0.7225101	0.0051745
alkire-foster5	0.6	0.1865894	0.0068123	0.2582516	0.0092455	0.7225101	0.0051745
alkire-foster6	0.7	0.0432649	0.0042978	0.0432649	0.0042978	1.0000000	0.0000000
alkire-foster7	0.8	0.0432649	0.0042978	0.0432649	0.0042978	1.0000000	0.0000000
alkire-foster8	0.9	0.0432649	0.0042978	0.0432649	0.0042978	1.0000000	0.0000000
alkire-foster9	1.0	0.0432649	0.0042978	0.0432649	0.0042978	1.0000000	0.0000000

```
svyafc(
            ~ wage + collgrad + hours ,
           design = des_nlsw88 ,
           cutoffs = list( 4 , 'college grad' , 26 ) ,
           k = ks,
           g = 0,
           na.rm = TRUE
   page_ten <-
       rbind(
           page_ten ,
            data.frame(
               k = ks,
               MO = coef( this_ks ) ,
               seMO = SE( this_ks ) ,
               H = attr( this_ks , "extra" )[ 1 , 1 ] ,
               seH = attr( this_ks , "extra" )[ 1 , 2 ] ,
               A = attr(this_ks, "extra")[2, 1],
               seA = attr( this_ks , "extra" )[ 2 , 2 ]
          )
        )
}
```

Replicate PDF page 13

```
page_thirteen <- NULL
```

Table 4.2: PDF Page 13 Replication

	k	MO	seMO	Н	seH	A	seA
alkire-foster	0.50	0.1913470	0.0069137	0.2689563	0.0093668	0.7114428	0.0068474
alkire-foster1	0.75	0.1489741	0.0066918	0.1842105	0.0081889	0.8087167	0.0052160
alkire-foster2	1.00	0.0432649	0.0042978	0.0432649	0.0042978	1.0000000	0.0000000

```
# loop through every poverty cutoff `k`
for( ks in c( 0.5 , 0.75 , 1 ) ){
    this_ks <-
       svyafc(
       ~ wage + collgrad + hours ,
       design = des_nlsw88 ,
       cutoffs = list( 4, 'college grad' , 26 ) ,
       k = ks,
       g = 0,
       dimw = c(0.5, 0.25, 0.25),
       na.rm = TRUE
   page_thirteen <-
       rbind(
           page_thirteen ,
           data.frame(
               k = ks,
               MO = coef( this_ks ) ,
               seMO = SE( this_ks ) ,
               H = attr( this_ks , "extra" )[ 1 , 1 ] ,
               seH = attr( this_ks , "extra" )[ 1 , 2 ] ,
               A = attr(this_ks, "extra")[2, 1],
               seA = attr( this_ks , "extra" )[ 2 , 2 ]
         )
       )
}
```

Replicate PDF page 16

```
page_sixteen <- NULL

# loop through every alpha value `g`
for( gs in 0:3 ){
    this_gs <-</pre>
```

Table 4.3: PDF Page 16 Replication

	g	МО	seMO
alkire-foster	0	0.3699078	0.0053059
alkire-foster1	1	0.2859332	0.0033708
alkire-foster2	2	0.2676266	0.0031164
alkire-foster3	3	0.2616335	0.0030531

```
svyafc(
    ~ wage + collgrad + hours ,
    design = des_nlsw88 ,
    cutoffs = list( 4, 'college grad' , 26 ) ,
    k = 1/3 ,
    g = gs,
    na.rm = TRUE
page_sixteen <-</pre>
    rbind(
        page_sixteen ,
        data.frame(
            g = gs,
            MO = coef( this_gs ) ,
            seMO = SE( this_gs )
      )
    )
```

Replicate k=1/3 rows of PDF page 17 and 19

```
## $overall
## alkire-foster SE
## alkire-foster 0.36991 0.0053
##
## $`raw headcount ratio`
```

```
##
    raw headcount
## wage 0.19492 0.0084
## collgrad 0.76316 0.0090
## hours 0.15165 0.0076
##
## $`censored headcount ratio`
## cens. headcount
## wage 0.19492 0.0084
## collgrad 0.76316 0.0090
## hours 0.15165 0.0076
## $`percentual contribution per dimension`
## dim. % contribution
                       0.17564 0.0061
## wage
## collgrad
                       0.68770 0.0077
## hours
                       0.13666 0.0059
Replicate PDF pages 21 and 22
svyafcdec(
    ~ wage + collgrad + hours ,
    subgroup = ~married ,
    design = des_nlsw88 ,
    cutoffs = list( 4 , 'college grad' , 26 ) ,
    k = 1/3 ,
    g = 0,
    na.rm = TRUE
)
## $overall
##
                 alkire-foster
                                   SE
## alkire-foster 0.36991 0.0053
##
## $`raw headcount ratio`
## raw headcount
             0.19492 0.0084
## wage
## collgrad 0.76316 0.0090
                 0.15165 0.0076
## hours
##
## $`censored headcount ratio`
## cens. headcount
## wage 0.19492 0.0084
## collgrad 0.76316 0.0090
## hours 0.15165 0.0076
## $`percentual contribution per dimension`
       dim. % contribution
##
```

```
## wage
                        0.17564 0.0061
## collgrad
                        0.68770 0.0077
## hours
                        0.13666 0.0059
## $`subgroup alkire-foster estimates`
##
           alkire-foster
## single
                 0.35414 0.0088
## married
                 0.37867 0.0066
##
## $`percentual contribution per subgroup`
           grp. % contribution
                       0.34204 0.012
## single
                       0.65796 0.012
## married
```

For additional usage examples of svyafc or svyafcdec, type ?convey::svyafc or ?convey::svyafcdec in the R console.

(Alkire and Foster, 2011) and (Alkire et al., 2015) and (Pacifico and Poge, 2016)

4.2 Bourguignon-Chakravarty (2003) multidimensional poverty class

A class of poverty measures is proposed in Bourguignon and Chakravarty (2003), using a cross-dimensional function that assigns values to each set of dimensionally normalized poverty gaps. It can be defined as:

$$BCh = \sum_{i \in n} \left[\left(\sum_{j \in d} w_j x_{ij} \right)^{\frac{1}{\theta}} \right]^{\alpha}, \ \theta > 0, \ \alpha > 0$$

where x_{ij} being the normalized poverty gap of dimension j for observation i, w_j is the weight of dimension j, θ and α are parameters of the function.

The parameter θ is the elasticity of substitution between the normalized gaps. In another words, θ defines the order of the weighted generalized mean across achievement dimensions. For instance, when $\theta=1$, the cross-dimensional aggregation becomes the weighted average of all dimensions. As θ increases, the importance of the individual's most deprived dimension increases. As de la Vega et al. (2009) points out, it also weights the inequality among deprivations. In its turn, α works as society's poverty-aversion measure parameter. In another words, as α increases, more weight is given to the most deprived individuals. Similar to θ , when $\alpha=1$, BCh is the average of the weighted deprivation scores.

4.3 Bourguignon (1999) inequality class (svybmi)

Bourguignon (1999) proposes a multidimensional inequality index that possesses interesting properties related to the correlation among the welfare dimensions measured. The estimator used in convey comes from the formula presented in Lugo (2007) and is defined as:

$$B_I = 1 - \frac{1}{\widehat{N}} \frac{\sum_{i \in S} w_i \left[\sum_{j \in d} w_j x_{ij} \right]^{\alpha/\beta}}{\left[\sum_{j \in d} w_j \mu_{ij} \right]^{\alpha/\beta}},$$

where $\alpha \geqslant 0$ is an inequality-aversion parameter and $\beta \leqslant 1$ is a parameter defining the degree of substitution among dimensions.

This measure is strong scale-invariant when $\beta=0$, although Bourguignon (1999) demonstrates that strong scale-dependent measures might be interesting in the context of multidimensional inequality. Also, it can be shown that stronger correlation among dimensions leads to less inequality if $\beta>\alpha$.

For additional usage examples of svybmi, type ?convey::svybmi in the R console.

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