Producing synthetic and confidential data using R-Package simPop

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### Why synthetic populations?

- methods comparison (e.g design-based simulation studies)
- policy modelling on individual level (e.g health planning, climate change, demographic change, economic change, ...)
- teaching (e.g. teaching of survey methods)
- Creation of public-/scientific-use files with low disclosure risk
- data availability is often a problem (legal issues, costs,...)

### Close-to-reality data

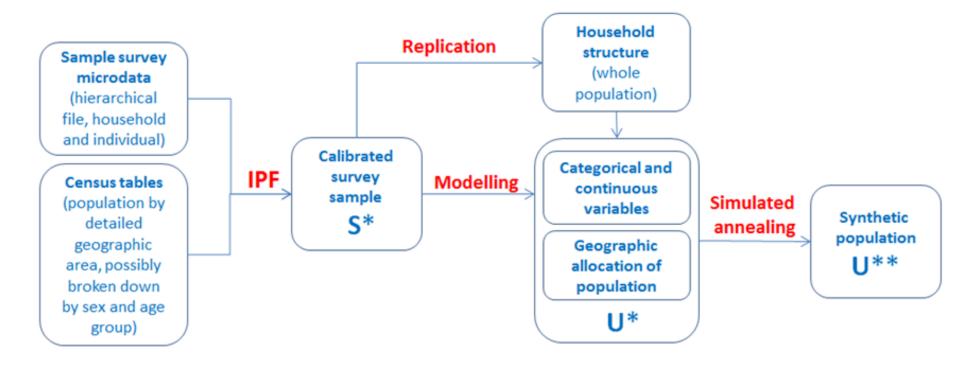
- actual sizes of regions and strata need to be reflected
- marginal distributions and interactions between variables should be represented correctly
- hierarchical and cluster structures have to be preserved
- Data confidentiality must be ensured
- Pure replication of units from the underlying sample should be avoided
- Sometimes some marginal distributions must exactly match known values

# **Approaches**

- choice of methods highly depends on available information
  - survey samples
  - aggregated information from samples
  - known marginal distributions from population
- in simPop: model-based approach is forced

# Model-based approach (1)

Possible flow, in reality often (much) more complex



# Model-based approach (2)

- In general, the procedure consists of four steps:
  - setup of the household structure
  - simulation of categorical variables
  - simulation of continuous variables
  - the splitting continuous variables into components
- Stratification: allows to account for heterogenities (e.g. regional differences)
- Sampling weights are considered in each step

### Model-based approach (3)

- Household structure (core-variables): simulated separately for each combination of strata and household size.
- # of households: estimated using the HT-estimator
- As few variables as possible (due to confidentiality reasons) are simulated using a sampling approach
- This builds up a realistic structure of the core variables
- Finally, additional variables are simulated using a regression-based approach using all existing variables.

## R package simPop

- Developed with support of the
  - International Household Survey Network
  - Department for International Development (DFID)
- latest version on CRAN
- highly object-oriented approach, similar to sdcMicro
- lets you produce synthetic confidential data
- efficiently programmed to work for (very) large data sets
- parallel computing is automatically be applied

### Example: EU-SILC

```
library(simPop)
str(origData)
```

```
'data.frame':
              11725 obs. of 18 variables:
$ db030
           : int 1123444555...
           : int 2 2 1 1 3 3 3 5 5 5 ...
$ hsize
$ db040
           : Factor w/ 9 levels "Burgenland", "Carinthia", ...: 4 4 7 5 7 7 7 4 4 4 ...
$ age
           : int 72 66 56 67 70 46 37 41 35 9 ...
           : Factor w/ 2 levels "male", "female": 1 2 2 2 2 1 1 1 2 2 ...
$ rb090
$ pl030
           : Factor w/ 7 levels "1", "2", "3", "4", ...: 5 5 2 5 5 3 1 1 3 NA ...
$ pb220a
           : Factor w/ 3 levels "AT", "EU", "Other": 1 1 1 1 1 1 3 1 1 NA ...
$ netIncome: num 22675 16999 19274 13319 14366 ...
$ py010n
           : num 0 0 19274 0 0 ...
$ py050n
           : num 00000 ...
$ py090n
           : num 00000 ...
$ pv100n
           : num 22675 0 0 13319 14366 ...
$ py110n
           : num 00000000 NA ...
$ py120n
           : num 000000000 NA ...
$ py130n
           : num 0 16999 0 0 0 ...
$ py140n
           : num 00000000 NA ...
$ db090
           : num 7.82 7.82 8.79 8.11 7.51 ...
$ rb050
           : num 7.82 7.82 8.79 8.11 7.51 ...
```

### Structure your data (once to be defined)

Create an object of class dataObj with function specifyInput().

```
inp <- specifyInput(data=origData, hhid="db030", hhsize="hsize", strata="db040", weight="rb050"); inp</pre>
```

#### Additional information

- (external) Population charateristics on EU-SILC variables can be specified as
  - data frame or
  - n-dimensional table

```
data(totalsRGtab); totalsRGtab # here: 2-dimensional table
        db040
         Burgenland Carinthia Lower Austria Salzburg Styria Tyrol
rb090
  female
             146980
                       285797
                                     828087
                                              722883 274675 619404
 male
             140436
                       270084
                                      797398
                                               702539 259595 595842
        db040
rb090
         Upper Austria Vienna Vorarlberg
  female
                368128 916150
                                  190343
  male
                353910 850596
                                  184939
```

• Calibration to this given known totals:

```
addWeights(inp) <- calibSample(inp, totalsRGtab)</pre>
```

# Simulating the basic structure

```
synthP <- simStructure(data=inp, method="direct", basicHHvars=c("age", "rb090", "db040"))
class(synthP)</pre>
```

```
[1] "simPopObj"
attr(,"package")
[1] "simPop"
```

- Output object ("synthP") is of class simPopObj
- various functions can be directly applied such objects

# Simulation of categorical variables

```
synth P <- sim Categorical(synth P, additional = c("pl030", "pb220a"), method = "multinom") \\ synth P
```

```
synthetic population of size
85057 x 10

build from a sample of size
11725 x 19
------
variables in the population:
db030,hsize,db040,age,rb090,db040.1,pid,weight,pl030,pb220a
```

# Simulating continuous variables

```
# multinomial model with random draws
synthP <- simContinuous(synthP, additional = "netIncome", upper = 2e+05, equidist = FALSE)</pre>
```

```
synthetic population of size
85057 x 12

build from a sample of size
11725 x 19
------
variables in the population:
db030,hsize,db040,age,rb090,db040.1,pid,weight,pl030,pb220a,netIncomeCat,netIncome
```

- Simulation of components with simComponents()
- Simulation of finer regional variables (like districts) with simInitSpatial()

#### Census information to calibrate

- Assumption: external information (n-dimensional table) is available, e.g marginals on  $region \times gender \times exonomic status$ .
- We add these marginals to the object and calibrate afterwards

```
synthP <- addKnownMargins(synthP, margins) # add margins

# calibration using simulated annealing
synthPadj <- calibPop(synthP, split="db040", temp=1, eps.factor=0.00005, maxiter=200, temp.cooldown=0.975, factor.cooldown=0.85, min.temp=0.001, verbose=FALSE)</pre>
```

# Continued, we end up with

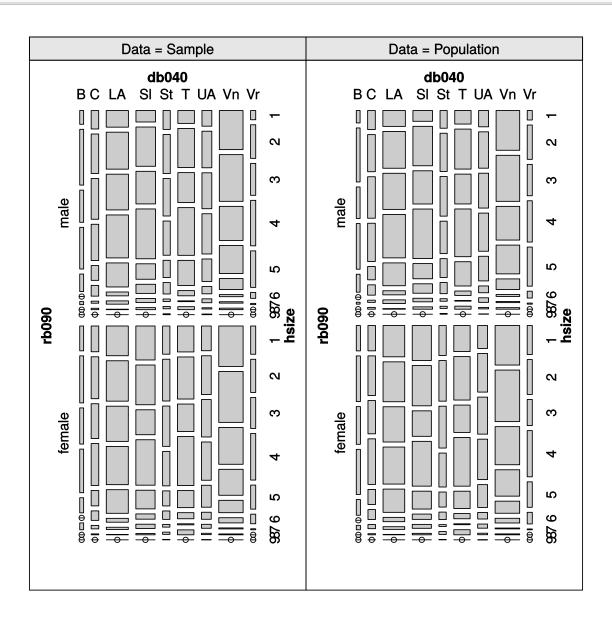
```
synthPadj
```

```
synthetic population of size
84975 x 12

build from a sample of size
11725 x 19
----------
variables in the population:
db030,hsize,db040,age,rb090,db040.1,pid,weight,pl030,pb220a,netIncomeCat,netIncome
```

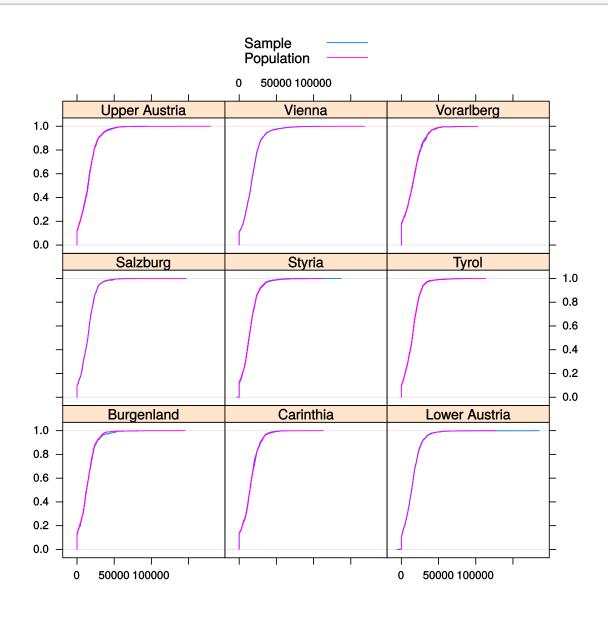
### Results

```
tab <- spTable(synthP, select=c("rb090","db040","hsize"))
spMosaic(tab, labeling=labeling_border(abbreviate=c(db040=TRUE)))</pre>
```



# Results

```
spCdfplot(synthPadj, "netIncome", cond="db040", layout=c(3, 3))
```



#### Conclusions

- margins of synthetic populations are calibrated
- all statistics can be very precisely estimated
- the synthetic populations are confidential
- code of **simPop** is quite efficient
- many other methods (IPU, IPF) are also included