Methodological advancements on the use of administrative data in Official Statistics - User Manual

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Dec 2022

Table of Contents

[Project aims and objectives 2](#_Toc121903161)

[Quality indicators 2](#_Toc121903162)

[R-indicators 3](#_Toc121903163)

[Using sample-based frame information as benchmark 3](#_Toc121903164)

[Distance metrics with Standardization 3](#_Toc121903165)

[Download and inspect the contents 4](#_Toc121903166)

[1 Download 4](#_Toc121903167)

[2 Downloaded contents explained 5](#_Toc121903168)

[Example datasets 5](#_Toc121903169)

[Folders and R scripts 5](#_Toc121903170)

[Launch RStudio and get ready 6](#_Toc121903171)

[1 Open the entire master folder in RStudio 6](#_Toc121903172)

[2 Set custom path 6](#_Toc121903173)

[3 Automatically create output folders 7](#_Toc121903174)

[4 Install packages 7](#_Toc121903175)

[RUNNING 2\_Prep\_Wtsample\_Freq\_Table.R 7](#_Toc121903176)

[PREP PART 1: Generate a weighted sample data 8](#_Toc121903177)

[PREP PART 2: Auxiliary file for R-indicators 10](#_Toc121903178)

[RUNNING 3A\_Distance\_Metrics.R 11](#_Toc121903179)

[RUNNING 3B\_R-indicator.R 15](#_Toc121903180)

[Q & A 19](#_Toc121903181)

[How do I know where to customise the code to suit my needs? 19](#_Toc121903182)

[How to use Starting path in multiple machines? 19](#_Toc121903183)

[What are the commonly used commands? 19](#_Toc121903184)

[How to free up memory space and speed up RStudio? 19](#_Toc121903185)

[I get error messages when a pre-defined function is used. 19](#_Toc121903186)

[How do I modify pre-defined functions? 20](#_Toc121903187)

[Technial notes and programming strategies 21](#_Toc121903188)

[Can I ignore Warning messages? 21](#_Toc121903189)

[Troubleshooting 21](#_Toc121903190)

[Unused argument error 21](#_Toc121903191)

[I get errors when computing… 21](#_Toc121903192)

[I am experiencing slowness in computation. 21](#_Toc121903193)

[Error: cannot allocate vector of size xxxx.x Gb 22](#_Toc121903194)

[What version of R is used? 22](#_Toc121903195)

[References 22](#_Toc121903196)

[Citation 22](#_Toc121903197)

This is the manual accompanying R code publicly available on GitHub for the project, *Methodological advancements on the use of administrative data in Official Statistics,* which is led by Professor Natalie Shlomo. As part of the research team, Sook Kim documented the manual.

# Project aims and objectives

The Office for National Statistics (ONS) have strategic priorities on embedding and advancing the use of administrative data into their official statistics processes. Their immediate priority is the use of administrative data in the quality assurance of the 2021 census and the production of administrative-based population estimates (ABPEs). A more long-term priority is the Population Statistics Transformation Programme which will feed into a recommendation to Government due in 2023 on the future of census and population statistics. In particular, the objective is to create population characteristic estimates from administrative and integrated data sources.

This manual concerns one of the sub-projects related to the quality framework for a single administrative data source. Here. we restrict the scope to distance metrics and R-indicators.

# Quality indicators

Sources of error for representation of administrative data are frame errors, selection errors and missing redundancy. We focus here on errors arising from coverage and representativity of administrative data, particularly when the data is streamed over time such as tax data for the business register or migration statistics for population estimates. It is vital that statistical agencies have good quality indicators to ensure the fit of administrative data to the population and to identify those sub-groups that are missing or over-covered, especially when the administrative data is used to quality assure other data sources such as surveys or a census.

We develop new methodology for calculating a quality indicator to measure representativeness and coverage.

## R-indicators

One such indicator is the R-indicator and its related partial R-indicators that were originally designed to assess the representativeness of responses from a survey and are particularly useful as an objective function in adaptive survey designs where data are collected over time (Schouten, et al. 2009, Schouten and Shlomo, 2009). The R-indicators measure the contrast between those who are missing and not missing in the data and identify those groups that are not represented in the data. We will investigate the usefulness of this framework to assess the representativeness and coverage of administrative data compared to a target population.

## Using sample-based frame information as benchmark

Recent research by Bianchi, et al. 2019 adapts the R-indicator to the case where only population auxiliary information are available instead of sample-based frame information. We draw upon the approach by Bianchi, et al. 2019, and utilise sample-based auxiliary information.

## Distance metrics with Standardization

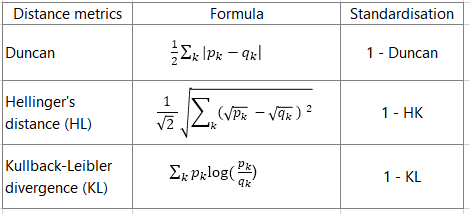
We look at other quality indicators. These are essentially distance metrics, such as the indicator of dissimilarity (Duncan and Duncan, 1955; Agresti, 2013). We also draw upon Hellinger’s distance (HL) and Kull-back-Leibler divergence (KL).

First, we define categories of categorical variable (or cross-classified) categorical variables by 𝑘, 𝑘=1, 2, …𝐾.

Let 𝑝𝑘 be the proportion of individuals in 𝑘 in the census (weighted survey count).

Let 𝑞𝑘 be the proportion of individuals in 𝑘 in the administrative data.

The Entropy is

The formulae for three distance metrics are given below. 

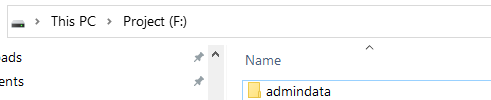
# Download and inspect the contents

## 1 Download

Please visit the github site here: [qualadmin link](https://github.com/sook-tusk/qualadmin). Click on Code at the top-right corner. Then, click on Download ZIP to download to your local machine.

Now, the downloaded folder needs to be placed in the meaningful location. We recommend users decide the appropriate Drive (C, D, E, F, etc) to house the downloaded contents. Then, **create a new folder** called admindata in File Explorer of your PC. Users can customise the new folder name as appropriate. This is your **starting path**.

The screenshot showing **starting path**:



Starting\_path

Under this Starting path, F:/admindata, place the downloaded folder from GitHub. Extract the zip folder as necessary.

As such, F:/admindata/qualadmin becomes the MASTER project folder. We’ll set it as working directory in RStudio later.

Notice that the terms, folder, directory, and path are used interchangeably in the user manual.

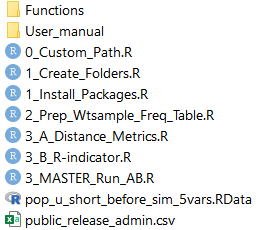
## 2 Downloaded contents explained

### Example datasets

We provide two example data sources.

| Data type | File name |
| --- | --- |
| Administrative | public\_release\_admin.csv |
| Census | pop\_u\_short\_before\_sim\_5vars.Rdata |

### Folders and R scripts

Under F:\admindata\qualadmin folder, you’ll be presented with the following contents. 

The **“User\_manual”** folder contains instructions on using the provided R code files.

The users do not need to do anything with the folder titled **“Functions”**. These pre-defined functions are used to either enclose complex procedures or perform repetitive tasks including cleaning and computing quality indicators. There are two files containing pre-defined functions. There is no need to run function files independently.

The functions will be automatically called in when the three main R script files are run: 2\_Prep\_Wtsample\_Freq\_Table.R 3\_A\_Distance\_Metrics.R 3\_B\_R-indicator.R

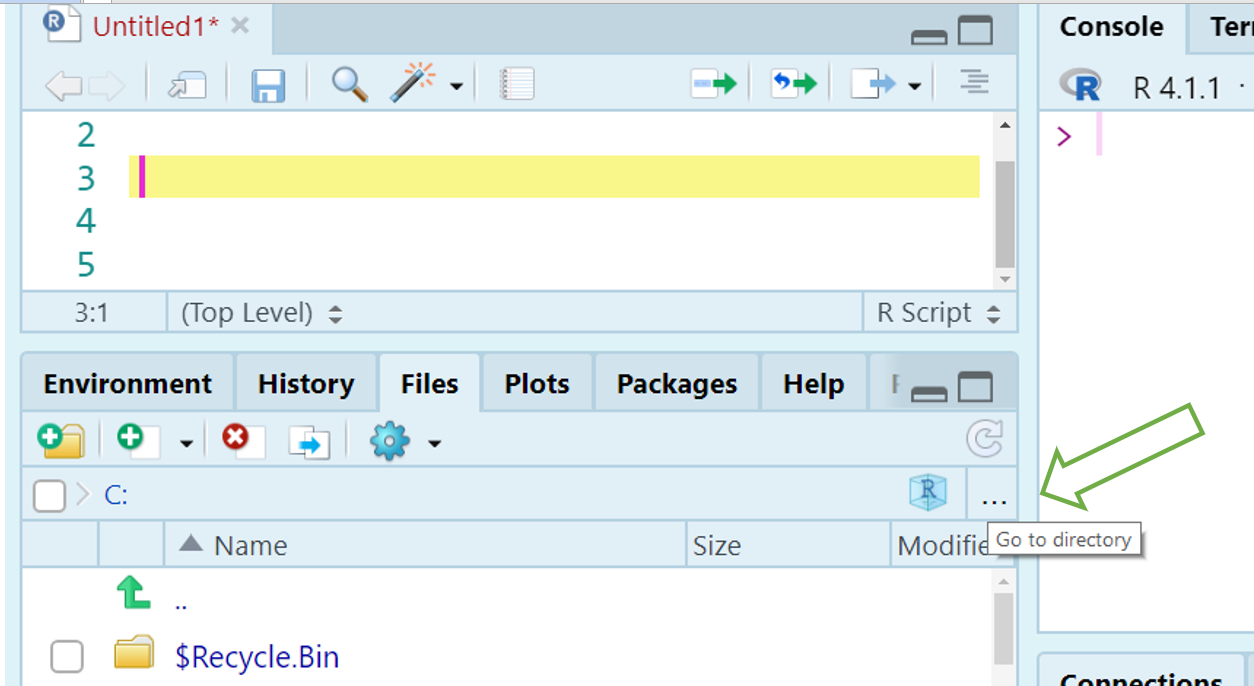
The 2\_Prep\_Wtsample\_Freq\_Table.R file creates necessary data needed to compute distance metrics and R-indicators. The master file, 3\_MASTER\_Run\_AB.R runs the above *two* main R script files, (3\_A\_Distance\_Metrics.R 3\_B\_R-indicator.R) automatically in sequence.

The first three files, 0\_Custom\_Path.R, 1\_Create\_Folders.R and 1\_Install\_Packages.R can be run to get ready to run the above main analysis files, as discussed in the following section.

# Launch RStudio and get ready

## 1 Open the entire master folder in RStudio

First, launch RStudio. Then, we need to **open the entire folder** F:/admindata/qualadmin where downloaded materials are located.

Unfortunately, RStudio has no feature in the menu, but you could do so by accessing **Files** tab. Click on ... as shown below.  Then, locate the master folder. In our example, it is F:/admindata/qualadmin.

## 2 Set custom path

Click open the R script file, 0\_Custom\_Path.R. Customise the starting path as needed, and set the path to indicate the master folder. The example code is:

# Starting path (CUSTOMISE PLEASE)  
 setwd("F:/admindata")  
  
 # Master project folder (USE AS IT IS)  
 setwd("./qualadmin")  
  
 # Check your current directory  
 getwd()

Please ensure to use a single forward slash / as above. R will print an error when backward slash \ is used in path. For instance,

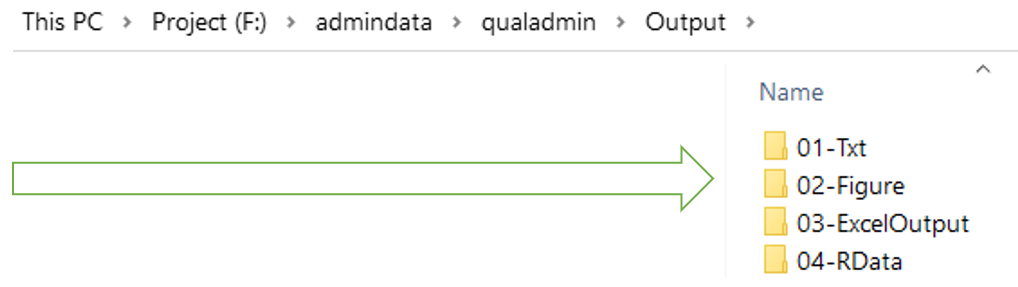
setwd("F:\admindata)`  
 Error: '\a' is an unrecognized escape in character string starting ""F:\a"

Please ensure your working directory is set at the master project path throught the analytical steps.

## 3 Automatically create output folders

The three main R script files 2\_Prep\_Wtsample\_Freq\_Table.R, 3\_A\_Distance\_Metrics.R, 3\_B\_R-indicator.R produce outputs. The outputs may be text, figure or in spreadsheet form. For the existing programmes to work, users need to create dedicated output folders.

To do so, please click on the 1\_Create\_Folders.R file to open. Then run line by line. The resulting folder structure is provided here:



Outputs\_folder

## 4 Install packages

The final preparation step is installing packages. Open 1\_Install\_Packages.R file, and run line by line.

#-----------------------------------  
 # Install packages (Run once)  
 #-----------------------------------  
   
 install.packages("ggplot2")  
 install.packages("tidyverse")  
   
 install.packages("car")

Now, you’re all set to proceed with quality measures indicators!

# RUNNING 2\_Prep\_Wtsample\_Freq\_Table.R

This code file consists of two parts: generating a weighted sample data, and preparing an auxiliary file for R-indicators.

## PREP PART 1: Generate a weighted sample data

* Open the 2\_Prep\_Wtsample\_Freq\_Table.R file.
* Step 1: Load a small percentage of Census data. It is called **pop\_u\_short\_before\_sim\_5vars** in the provided example code.

#H---------------------------------------  
 ##> 1. Load Census data  
 #H--------------------------------------  
  
 load("pop\_u\_short\_before\_sim\_5vars.RData")  
 df <- pop\_u\_short\_before\_sim\_5vars  
 dim(df) # obs = 1163659  
 names(df)

In our example Census data, we have 1,163,659 observations with five categorical variables including geography, sex, age groups, ethnic groups, and economic activity status. One can declare which variable to tabulate. Here, we declare all five variables using var object[[1]](#footnote-1).

var <- c("geog1", "sex", "agecode1",  
 "eth\_code5", "econg")

The description of categories, and distribution is shown below.

| Variable | Category | Description | N | (%) |
| --- | --- | --- | --- | --- |
| Total |  |  | 1163659 |  |
| geog1 | 1 | LA codes | 116128 | (10.0) |
|  | 2 | LA codes | 150139 | (12.9) |
|  | 3 | LA codes | 137520 | (11.8) |
|  | 4 | LA codes | 170624 | (14.7) |
|  | 5 | LA codes | 90873 | (7.8) |
|  | 6 | LA codes | 498375 | (42.8) |
| sex | 1 | male | 564905 | (48.5) |
|  | 2 | female | 598754 | (51.5) |
| agecode1 | 1 | 16-20 | 82426 | (7.1) |
|  | 2 | 21-25 | 94643 | (8.1) |
|  | 3 | 26-30 | 110296 | (9.5) |
|  | 4 | 31-35 | 120398 | (10.3) |
|  | 5 | 36-40 | 119393 | (10.3) |
|  | 6 | 14-45 | 101711 | (8.7) |
|  | 7 | 46-50 | 94209 | (8.1) |
|  | 8 | 51-55 | 100159 | (8.6) |
|  | 9 | 56-60 | 77799 | (6.7) |
|  | 10 | 61-65 | 65833 | (5.7) |
|  | 11 | 66-70 | 57305 | (4.9) |
|  | 12 | 71-75 | 51263 | (4.4) |
|  | 13 | 76-80 | 43678 | (3.8) |
|  | 14 | 81+ | 44546 | (3.8) |
| eth\_code5 | 1 | White | 1081812 | (93.0) |
|  | 2 | Mixed/Multiple ethnic groups | 10487 | (0.9) |
|  | 3 | Asian/Asian British | 46446 | (4.0) |
|  | 3 | Black/African/Caribbean/Black British | 16268 | (1.4) |
|  | 4 | Other ethnic group | 8646 | (0.7) |
| econg | 1 | In employment(FT, PT) | 689140 | (59.2) |
|  | 2 | Unemployed | 27744 | (2.4) |
|  | 3 | Out of workforce | 446775 | (38.4) |

Using this prior information on the population distribution (based on Census), we can mimic the distribution in a random sample. See the next step.

* Step 2: Then, we draw a random sample 1:50.
* Step 3: From the randomly selected sample (1163659/50 = 23273), we then obtain frequency table of categorical variables (count of categories). Users can run the pre-defined function, fn\_maxvar5\_freq\_table() to perform the task. The function[[2]](#footnote-2) automatically obtains counts and structure the output in long form, organised by each variable, and by its discrete category.

fn\_maxvar5\_freq\_table()

This procedure is to *assess distribution* of categories in a random sample (N = 23273). Based on the counts of the randomly selected sample, we multiply the counts by 50. One may wonder why we multiply. As we *reduced* the census sample by drawing a random sample by the 1:50 ratio, we need to *convert* the shrank sample back to the original size (with the priori distribution). That’s why we multiply by 50 (Weighted Sample N = 23273 \* 50 = 1163650). This completes the process of generating weighted sample survey data.

* Step 4: Carry out checks to see if the calculated frequency tables are accurate.

freq\_table[1:8, 1:9]

## seq twdigits raw\_n n p oneway v by1 by2  
## 1 1 101 2265 113250 0.09732308 1 1 geog1 01  
## 2 2 102 2968 148400 0.12752976 1 1 geog1 02  
## 3 3 103 2749 137450 0.11811971 1 1 geog1 03  
## 4 4 104 3502 175100 0.15047480 1 1 geog1 04  
## 5 5 105 1846 92300 0.07931938 1 1 geog1 05  
## 6 6 106 9943 497150 0.42723327 1 1 geog1 06  
## 7 7 201 11307 565350 0.48584196 1 2 sex 01  
## 8 8 202 11966 598300 0.51415804 1 2 sex 02

When we printed the first 8 lines and 10 variables, we can see the count, n, and the corresponding proportion, p by each variable. The following code obtains the total observation size and confirms that the total proportion adds up to 1, for geog1 variable. Here, the total observation size can be viewed as the population size.

# Check whether the total adds up to 1  
 sum(freq\_table[1:6, "n"])

## [1] 1163650

sum(freq\_table[1:6, "p"])

## [1] 1

* Step 5: Rename and save.
* Step 6: Export the output frequency table in Excel with the file name, **Weightedsample\_freq\_table.xlsx**.
* Step 7: Save the R objects as RData. Done.

## PREP PART 2: Auxiliary file for R-indicators

Thus far, we have created a weighted sample data based on the random sampling procedure, and obtained both one-way and two-way frequency tables. We also identified the population size of 1,116,350 (popsize = 1163650).

For R-indicator calculations, we need to compute meanpop by variables which are geography, sex, age groups, ethnic groups, and economic activity status.

# Compute meanpop  
 auxiliary <- freq\_table %>%  
 filter(oneway == 1) %>%  
 group\_by(by1) %>%  
 mutate(meanpop = n / popsize) %>%  
 ungroup() %>%  
 dplyr::select(seq, count = n, by1,  
 v, by2, meanpop, raw\_n)

To do so, we first remove two-way and keep the one-way frequency table only. Then, by variable-level, which is indicated by by1, we compute meanpop. As seen before, the count of each category is stored in n. The meanpop is obtained by dividing n by popsize. For example, the value of **meanpop** for first category of geog1 is calculated as 113250/1163650 = 0.0973, and the second category of geog1 is 148400/1163650 = 0.1275 and so on.

Finally, we add the popsize in the first row, to complete the Auxiliary file.

Let’s look at the Auxiliary file:

print(wtsample\_auxiliary\_econg)

## seq count by1 v by2 meanpop raw\_n type  
## 1 1 1163650 total 0 00 1.000000000 0 wtsample  
## 2 2 113250 geog1 1 01 0.097323078 2265 wtsample  
## 3 3 148400 geog1 1 02 0.127529756 2968 wtsample  
## 4 4 137450 geog1 1 03 0.118119710 2749 wtsample  
## 5 5 175100 geog1 1 04 0.150474799 3502 wtsample  
## 6 6 92300 geog1 1 05 0.079319383 1846 wtsample  
## 7 7 497150 geog1 1 06 0.427233275 9943 wtsample  
## 8 8 565350 sex 2 01 0.485841963 11307 wtsample  
## 9 9 598300 sex 2 02 0.514158037 11966 wtsample  
## 10 10 84600 agecode1 3 01 0.072702273 1692 wtsample  
## 11 11 93300 agecode1 3 02 0.080178748 1866 wtsample  
## 12 12 111200 agecode1 3 03 0.095561380 2224 wtsample  
## 13 13 124250 agecode1 3 04 0.106776092 2485 wtsample  
## 14 14 118200 agecode1 3 05 0.101576935 2364 wtsample  
## 15 15 99950 agecode1 3 06 0.085893525 1999 wtsample  
## 16 16 95800 agecode1 3 07 0.082327160 1916 wtsample  
## 17 17 95600 agecode1 3 08 0.082155287 1912 wtsample  
## 18 18 78450 agecode1 3 09 0.067417179 1569 wtsample  
## 19 19 67600 agecode1 3 10 0.058093069 1352 wtsample  
## 20 20 57950 agecode1 3 11 0.049800198 1159 wtsample  
## 21 21 50650 agecode1 3 12 0.043526834 1013 wtsample  
## 22 22 42250 agecode1 3 13 0.036308168 845 wtsample  
## 23 23 43850 agecode1 3 14 0.037683152 877 wtsample  
## 24 24 1083250 eth\_code5 4 01 0.930907060 21665 wtsample  
## 25 25 10450 eth\_code5 4 02 0.008980364 209 wtsample  
## 26 26 45600 eth\_code5 4 03 0.039187041 912 wtsample  
## 27 27 16300 eth\_code5 4 04 0.014007648 326 wtsample  
## 28 28 8050 eth\_code5 4 05 0.006917888 161 wtsample  
## 29 29 691300 econg 5 01 0.594078976 13826 wtsample  
## 30 30 28250 econg 5 02 0.024277059 565 wtsample  
## 31 31 444100 econg 5 03 0.381643965 8882 wtsample

# RUNNING 3A\_Distance\_Metrics.R

To calculate distance metrics, we first obtain one-way, and two-way frequency tables of categorical variables, and calculates proportions of each sub-category by each data source. Next, we combines master (benchmark) and admin freq tables. Then, we create domains for quality indicators, and finally compute distance metrics as part of quality indicators. We offer three different types of distance metrics. To allow comparison across the metrics, we standardise the calculations.

Users can also produce a summary table of three types of distance metrics, and visualise the results.

The preliminary step is to ensure we have benchmark data. Simply *source* the previous file as below to update it.

source("2\_Prep\_Wtsample\_Freq\_Table.R")

* Step 1: read admin data

df <- read\_csv("public\_release\_admin.csv")

## Rows: 1033664 Columns: 6  
## -- Column specification --------------------------------------------------------  
## Delimiter: ","  
## dbl (6): person\_id, geog1a, sex, agecode1, eth\_code5, econg  
##   
## i Use `spec()` to retrieve the full column specification for this data.  
## i Specify the column types or set `show\_col\_types = FALSE` to quiet this message.

tail(df)

## # A tibble: 6 x 6  
## person\_id geog1a sex agecode1 eth\_code5 econg  
## <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>  
## 1 1033659 1 1 6 1 1  
## 2 1033660 6 1 6 1 1  
## 3 1033661 6 2 5 1 1  
## 4 1033662 5 1 12 1 3  
## 5 1033663 1 2 9 1 1  
## 6 1033664 6 2 9 1 1

The last six observations of the example admin data is shown above. As the person id goes up to 1033664, we can see there are 1033664 observations in admin data. We also notice that geog1a is used instead of *geog1*. As such, we declare all five variables for tabulations.

var <- c("geog1a", "sex", "agecode1",  
 "eth\_code5", "econg")

* Step 2: Obtain admin freq tables.

As mentioned in the previous section, we obtain frequency table of categorical variables (count of categories). Users can run the pre-defined function, fn\_maxvar5\_freq\_table() to perform the task. The function[[3]](#footnote-3) automatically obtains counts and structure the output in long form, organised by each variable, and by its discrete category.

Once the frequency tables are obtained, we rename the object as Admin\_f\_table\_one. Let’s inspect Admin\_f\_table\_one.

head(Admin\_f\_table\_one)

## seq twdigits admin\_n admin\_perc oneway v by1 by2 by3 by4 by5  
## 1 1 101 137993 0.1334989 1 1 geog1a 01 oneway 0 0  
## 2 2 102 124051 0.1200110 1 1 geog1a 02 oneway 0 0  
## 3 3 103 131176 0.1269039 1 1 geog1a 03 oneway 0 0  
## 4 4 104 139867 0.1353119 1 1 geog1a 04 oneway 0 0  
## 5 5 105 142304 0.1376695 1 1 geog1a 05 oneway 0 0  
## 6 6 106 358273 0.3466049 1 1 geog1a 06 oneway 0 0

tail(Admin\_f\_table\_one)

## seq twdigits admin\_n admin\_perc oneway v by1 by2 by3 by4 by5  
## 340 340 404501 8557 0.0082783187 2 4 eth\_code5 04 econg 5 01  
## 341 341 404502 791 0.0007652390 2 4 eth\_code5 04 econg 5 02  
## 342 342 404503 5007 0.0048439338 2 4 eth\_code5 04 econg 5 03  
## 343 343 405501 3867 0.0037410609 2 4 eth\_code5 05 econg 5 01  
## 344 344 405502 255 0.0002466953 2 4 eth\_code5 05 econg 5 02  
## 345 345 405503 3420 0.0033086187 2 4 eth\_code5 05 econg 5 03

* Step 3: Merge admin + Weighted sample freq tables

fn\_merge\_one\_admin\_wtsample\_f\_table\_temp()

The code above executes merging two data sources.

* Step 4: Create domains of quality indicators

fn\_create\_domain\_temp()

To check the domains, we can use *Janitor* package’s tabyl function[[4]](#footnote-4). The function creates 15 domains, including five single variables’ domain, and ten bivariate domains.

display\_domain %>% tabyl(fct\_domain)

## fct\_domain n percent  
## geog1 6 0.017391304  
## sex 2 0.005797101  
## agecode1 14 0.040579710  
## eth\_code5 5 0.014492754  
## econg 3 0.008695652  
## geog1:sex 12 0.034782609  
## geog1:agecode1 84 0.243478261  
## geog1:eth\_code5 30 0.086956522  
## geog1:econg 18 0.052173913  
## sex:agecode1 28 0.081159420  
## sex:eth\_code5 10 0.028985507  
## sex:econg 6 0.017391304  
## agecode1:eth\_code5 70 0.202898551  
## agecode1:econg 42 0.121739130  
## eth\_code5:econg 15 0.043478261

* Step 5: Compute distance metrics

Run the functions to compute three types of distance metrics.

fn\_unstd\_distance\_metrics\_full()  
 fn\_unstd\_distance\_metrics\_tidy()

* Step 6: Standardise distance metrics
* Step 7: Reshape, then tidy

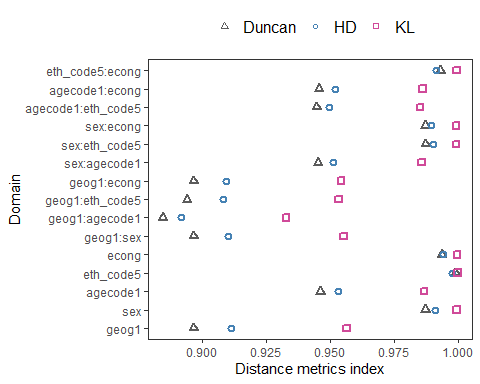
From wide form, the outputs have been reshaped to long form. Then, we keep standardised solutions. The results are as follows:

df <- distance\_metrics\_long  
 # std\_test(1-Duncan, 1-HD, 1-KL) only  
 df <- df %>% filter(std\_test\_use == 1)  
  
 df[1:9, c(1:2, 4:5, 9)]

## # A tibble: 9 x 5  
## domain\_id domain indicator index std\_test\_use  
## <int> <chr> <chr> <dbl> <dbl>  
## 1 1 geog1 Std\_Duncan 0.897 1  
## 2 1 geog1 Std\_t\_HD 0.911 1  
## 3 1 geog1 Std\_t\_KL 0.957 1  
## 4 2 sex Std\_Duncan 0.987 1  
## 5 2 sex Std\_t\_HD 0.991 1  
## 6 2 sex Std\_t\_KL 1.00 1  
## 7 3 agecode1 Std\_Duncan 0.946 1  
## 8 3 agecode1 Std\_t\_HD 0.953 1  
## 9 3 agecode1 Std\_t\_KL 0.987 1

* Step 8: Plot the distance metrics

plot(p)



# RUNNING 3B\_R-indicator.R

The file computes the overall R-indicator. Users can also proceed with computing **partial** R-indicators by category level, and variable level. The procedure can be computationally extensive. This is noted in the relevant section, so that users can allow some time to execute the code.

* Step 1: Benchmark mean population ready

Load auxiliary file and remove the last category From each categorical variable (group\_by(by1)), we identify the last category (max(by2)). We then remove the last category(dplyr::select(-c(lastcat\_, lastcat))). Here, we explicitly instruct R to use dplyr package to access select function[[5]](#footnote-5).

We only need meanpop. It is shown in a single column (column vector). Let’s check.

# Print meanpop  
 print(col\_auxiliary[1:nrow(col\_auxiliary), "meanpop"],  
 n = nrow(col\_auxiliary))

## # A tibble: 26 x 1  
## meanpop  
## <dbl>  
## 1 1   
## 2 0.0973   
## 3 0.128   
## 4 0.118   
## 5 0.150   
## 6 0.0793   
## 7 0.486   
## 8 0.0727   
## 9 0.0802   
## 10 0.0956   
## 11 0.107   
## 12 0.102   
## 13 0.0859   
## 14 0.0823   
## 15 0.0822   
## 16 0.0674   
## 17 0.0581   
## 18 0.0498   
## 19 0.0435   
## 20 0.0363   
## 21 0.931   
## 22 0.00898  
## 23 0.0392   
## 24 0.0140   
## 25 0.594   
## 26 0.0243

* Step 2: Keep wtsample distributions as row vectors

We then transpose meanpop to a single row format (row vector). Now, our benchmark data is ready. The next step is open the corresponding administrative data, and carry out computing R-indicators.

* Step 3: Compute R-indicators

Users can use pre-defined functions. Please allow a minute to execute.

aa <- read\_csv("public\_release\_admin.csv")  
  
 nrow(aa) # 1033664  
  
 df <- NULL  
 between <- NULL  
 partial <- NULL  
 partialtemp <- NULL  
 fn\_overall\_r\_indicator\_1()  
 fn\_overall\_r\_indicator\_2()  
 fn\_overall\_r\_indicator\_3()  
 fn\_overall\_r\_indicator\_4()

Users can consult Functions/2\_Functions\_R-indicators.R file for more details and operationalisation.

* Step 4: Save in Excel and inspect

At this stage, users can inspect accordingly. Let’s have a look. Here, we can see the overall R-indicator is estimated as 0.496 based on the administrative data (N=1033664). Looking at the variable-level R-indicator (see rows 4-8), geog1a was seen to have the greatest R-indicator (0.04) compared to econg (0.0002).

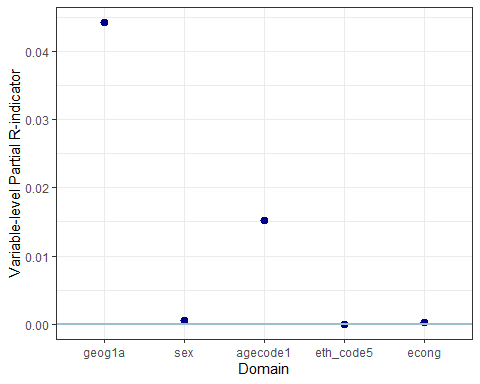
partial[1:17, c(1:2, 4, 8:10)]

## seq domain R\_indicator count n\_cat domain\_n  
## 1 1 Overall 0.49602631564 NA <NA> NA  
## 2 2 mrphatall 0.95977686066 NA <NA> NA  
## 3 3 resppop 1033664.00000000000 NA <NA> NA  
## 4 4 geog1a 0.04424182738 NA <NA> NA  
## 5 5 sex 0.00049830140 NA <NA> NA  
## 6 6 agecode1 0.01528411959 NA <NA> NA  
## 7 7 eth\_code5 0.00002968059 NA <NA> NA  
## 8 8 econg 0.00021170900 NA <NA> NA  
## 9 9 des1 NA 0 1 0  
## 10 10 geog1a\_1 0.08795013527 137993 1 1  
## 11 11 geog1a\_2 -0.01927948285 124051 2 1  
## 12 12 geog1a\_3 0.02190390676 131176 3 1  
## 13 13 geog1a\_4 -0.03661612887 139867 4 1  
## 14 14 geog1a\_5 0.13969467338 142304 5 1  
## 15 15 geog1a\_6 -0.12165434160 358273 6 1  
## 16 16 sex\_1 -0.01620055057 489228 1 2  
## 17 17 sex\_2 0.01535719891 544436 2 2

* Step 5: Scatterplot

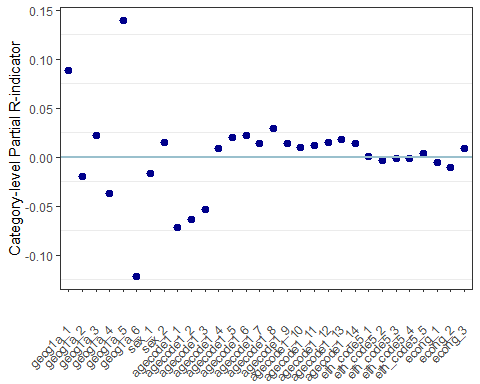
R-indicator by the variable level.

plot(p1)



R-indicator by the category-level.

plot(p2)



This concludes the manual. Thank you for taking the time reading the material. Please get in touch with any query or errata at [fanfurcada@gmail.com](mailto:fanfurcada@gmail.com).

If you need technical support, please consult Q & A.

# Q & A

## How do I know where to customise the code to suit my needs?

Unless indicated as “Customise as needed”, users can run the code as it is. Please consult each code file.

## How to use Starting path in multiple machines?

If users plan to use different machines, simply by changing the “starting path”, users can carry out the analysis with minimal disruption. To achieve this, please ensure to use the consistent master project folder name.

## What are the commonly used commands?

Most commonly used commands in the tidyverse package are:

arrange : sort variables.  
 bind\_rows: append multiple dataframes.  
 mutate : manipulate variables, and  
 create new variables based on old variables.  
 select : order, and keep(drop) variables of interest.  
 shell.exec: launch a software and opens the target file (Windows PC only)

## How to free up memory space and speed up RStudio?

You can remove objects that you no longer need.

# To remove objects except for certain objects  
 ls()  
 keepobjectslist <- c("a", "b", "c")  
 rm(list = ls()[!ls() %in% keepobjectslist])  
 ls()

## I get error messages when a pre-defined function is used.

Users can inspect the codes used in the function, and identify the issues. It is recommended NOT edit the function file directly, as the functions are used repeatedly, and the interlinked sections may not run as expected. Where preferable, users may copy the codes in the function, and use locally with minor tweaks.

## How do I modify pre-defined functions?

Users can modify 1\_Functions and 2\_Functions\_R-indicators.R under *Functions* folder.

# 1\_Functions.R  
fn\_output\_folder\_path <- function() {  
  
 currentdate <<- Sys.Date()  
 txtpath <<- "Output/01-Txt/"  
 figpath <<- "Output/02-Figure/"  
 xlsxpath <<- "./Output/03-ExcelOutput/"  
 Rdatapath <<- "Output/04-RData/"  
}

We can check how the output folder names are set as path to save the results during the analytical process.

fn\_output\_folder\_path()

Let’s run the function. We can see that xlsxpath is set as "./Output/03-ExcelOutput/".

xlsxpath

## [1] "./Output/03-ExcelOutput/"

Let’s customise the xlsxpath, by renaming the folder name. If we customise 1\_Functions.R file, we can edit the information enclosed in the brackets. Notice that we use <<- with functions so that the object created by a function will exist in the global R environment. This is very important.

Alternatively, We could ignore the pre-defined function and just write relevant lines of code and keep it in the main R script file. For instance, we could put output\_folder\_path at the top of the 2\_Prep\_Wtsample\_Freq\_Table.R. Here, we edited the xlsxpath. Notice that fn\_output\_folder\_path <- function() { } is removed.

xlsxpath\_2 <- "./Output/03-Excel/"  
   
 xlsxpath\_2

## [1] "./Output/03-Excel/"

#H---------------------------------------  
 ## > Step 1. Load Census data  
 #H--------------------------------------  
 # load("pop\_u\_short\_before\_sim\_5vars.RData")

Notice that we use <-. Using <<- is not necessary here. Users can remember the usage of <- and can modify the functions as appropriate, should the function incurs errors.

## Technial notes and programming strategies

When loop is used, base R functions were used (table, tapply, etc). For data manipulation, tidyverse package was used extensively. This strategy is partly to improve readability of the code.

To enhance users’ workflow, output files are programmed to launch using the pre-defined functions.

## Can I ignore Warning messages?

Some packages alert users with compatibility issues arising from old version. These can be ignored. For example,

library("fastDummies")  
 Warning message:  
 package 'fastDummies' was built under  
 R version 4.1.2  
  
 library(rlist)  
 Warning message:  
 package 'rlist' was built under R version 4.1.2

## Troubleshooting

### Unused argument error

For example, sim %>% select(geog1a) the select command can cause an error:

Error in select(., geog1a) :  
 unused arguments (geog1a)

This maybe due to the conflict in packages.

The error can be fixed by adding the name of the package used, dplyr, explicitly. sim %>% dplyr::select(geog1a)

### I get errors when computing…

Please inspect zero cells, and ensure 0 (numeric value) is entered for n and perc, as well as admin\_n and admin\_perc. Errors may occur with NA coding and data attributes(character, factor, numeric).

### I am experiencing slowness in computation.

R can be not responsive if memory is full. Please identify bottlenecks and remove them. It may be due to certain commands. For example, View(object) command could take a while if the object is huge in size. Unless one should inspect the data, suppress the View command to expedite the computation where possible.

It can also be the case that for loop functions can be slow as well. In some instances, removing objects may help as this procedure can free up memory space. See above commonly used commands for more information.

### Error: cannot allocate vector of size xxxx.x Gb

If matrix symbols have entered mistakenly, R shows an error message like this. Please double check whether there are any mistakes. For instance, one may have typed a\*b instead of a%\*%b. Users can type memory.limit() to check the current memory limit and increase as necessary.

## What version of R is used?

Tested with Windows PC. R version used: 4.1.1 RStudio version: RStudio 2022.07.2 Build 576

# References

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# Citation

Please cite this work as:

Shlomo, Natalie & Kim, Sook (2022). “Methodological advancements on the use of administrative data in Official Statistics - User Manual”, available at <https://github.com/sook-tusk/qualadmin>

1. As the var object is treated as global macro, the programme runs automatically using the information stored in global macro, and produces the results. [↑](#footnote-ref-1)
2. We also provide fn\_maxvar4\_freq\_table() for users who declare four categorical variables [↑](#footnote-ref-2)
3. We also provide fn\_maxvar4\_freq\_table() for users who declare four categorical variables [↑](#footnote-ref-3)
4. This is essentially almost identical to table(display\_domain$fct\_domain), but the approach tabyl produces percent by default [↑](#footnote-ref-4)
5. This is to avoid warnings messages from R when R searches for a particular function from two different packages. [↑](#footnote-ref-5)