

Package: dfertility (via r-universe)

August 15, 2024

Type Package

Title District level estimation of age-specific fertility

Version 0.1.666

Description This package estimates district-level estimates of age-specific fertility from nationally representative household survey data.

Depends R (>= 2.10)

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Encoding UTF-8

LazyData true

Roxygen list(markdown = TRUE)

RoxygenNote 7.1.2

Imports Matrix, countrycode, R6, TMB, ggplot2, here, magrittr, dplyr, parallel, mvtnorm, rdhs, sf, spdep, tidyverse, zip, zoo, methods, utils, rlang

Suggests tidyverse

LinkingTo RcppEigen, TMB

Repository <https://mrc-ide.r-universe.dev>

RemoteUrl <https://github.com/mrc-ide/dfertility>

RemoteRef master

RemoteSha e3b480d0fd14a828776c2a197c559cb2b5e76814

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area_populations	<i>Aggregate district level populations</i>
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Description

Aggregate district level populations to all levels of area hierarchy

Usage

```
area_populations(population, areas_wide, project, naomi_level)
```

Arguments

population	Age/sex/space/time stratified population
areas_wide	Area hierarchy in wide format

assign_cluster_area	<i>Aggregate district level cluster dataframe</i>
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Description

Aggregate a dataframe of survey clusters geolocated to an administrative to any higher administrative level

Usage

```
assign_cluster_area(clusters, areas_wide, area_level)
```

Arguments

<code>clusters</code>	Dataframe of geolocated survey clusters
<code>areas_wide</code>	Area hierarchy in wide format
<code>area_level</code>	Integer referring to administrative level, where 0 is national, 1 is provincial etc.

Value

Dataframe list of clusters by administrative area, one list item per survey

See Also

[get_areas](#)

`calculate_dhs_fertility`

Calculate fertility rates from DHS, MIS, AIS data

Description

Calculate fertility rates from DHS, MIS, AIS data

Usage

```
calculate_dhs_fertility(iso3, surveys, clusters, areas_wide)
```

`calculate_mics_fertility`

Calculate fertility rates from MICS data

Description

Calculate fertility rates from MICS data

Usage

```
calculate_mics_fertility(iso3, mics_wm, mics_births_to_women)
```

create_surveys_mics *Download MICS surveys*

Description

Download MICS surveys

Usage

```
create_surveys_mics(iso3, mics_indicators)
```

Arguments

iso3 ISO3 code

filter_mics *Filter MICS datasets*

Description

Filter MICS household, women, and birth history datasets to key variables, and rename to ensure consistent column names between surveys

Usage

```
filter_mics(dat, mics_indicators, survey_id_i)
```

fit_sample_tmb *Fit TMB model*

Description

Fit TMB model

Usage

```
fit_sample_tmb(data, par, random)
```

get_areas	<i>Make area dataframes</i>
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Description

Create area hierarchy dataframes in the Naomi package format. The function will always return the area hierarchy in long format, with arguments additionally to return full and wide hierarchies

Usage

```
get_areas(iso3_current, naomi_data_path, full = FALSE, wide = TRUE)
```

Arguments

iso3_current	A string of one or more ISO-3 codes
naomi_data_path	A path to directory containing input data files
full	Boolean to return the full area heirarchy
wide	Boolean to return the area hierarchy in wide format

get_boundaries	<i>Get populations</i>
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Description

Returns a dataframe of area boundaries

Usage

```
get_boundaries(iso3_current, naomi_data_path)
```

Arguments

iso3_current	A string of one or more ISO-3 codes
naomi_data_path	A path to directory containing input data files

`get_input_files` *Extract area, population, boundary files from target directory*

Description

Extract area, population, boundary files from target directory

Usage

```
get_input_files(iso3_current, naomi_data_path)
```

Arguments

`iso3_current` A string of one or more ISO-3 codes
`naomi_data_path` A path to directory containing input data files

Value

A named list of file paths

`get_populations` *Get populations*

Description

Returns a dataframe of population by district, five year age group, and sex

Usage

```
get_populations(iso3_current, naomi_data_path)
```

Arguments

`iso3_current` A string of one or more ISO-3 codes
`naomi_data_path` A path to directory containing input data files

<code>ir_by_area</code>	<i>Join individual recode survey datasets by area</i>
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Description

Join individual recode survey datasets by area

Usage

```
ir_by_area(ir, area_list, n, total)
```

Arguments

<code>ir</code>	Individual recode survey dataset
<code>area_list</code>	List of areas

<code>join_survey_areas</code>	<i>Join household datasets with area hierarchy</i>
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Description

Join household datasets with area hierarchy

Usage

```
join_survey_areas(fertility_mics_data, areas, warn = FALSE)
```

<code>make_adjacency_matrix</code>	<i>Neighbor list</i>
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Description

Neighbor list

Usage

```
make_adjacency_matrix(areas, model_level)
```

`make_asfr_inputs`*Transform survey datasets into inputs for calc_asfr***Description**

Transform survey datasets into inputs for calc_asfr

Usage

```
make_asfr_inputs(mics_survey_areas, mics_survey_data)
```

`make_model_frames_batch`*Make model frames for batch running***Description**

Create model frames and aggregation matrices for TMB model.

Usage

```
make_model_frames_batch(lvl_map, population, asfr, areas_list, project = 2020)
```

Arguments

<code>population</code>	Age/sex/space stratified population
<code>asfr</code>	ASFRs by district and time
<code>areas_list</code>	List of area files
<code>project</code>	Model will by default produce estimates up to year of last survey. Integer projection year if desired, else FALSE
<code>lvl</code>	Dataframe of district and province levels

make_model_frames_dev *Make model frames*

Description

Create model frames and aggregation matrices for TMB model.

Usage

```
make_model_frames_dev(  
  iso3,  
  population,  
  asfr,  
  areas,  
  naomi_level,  
  project = 2020  
)
```

Arguments

iso3	iso3 code for country
population	Age/sex/space stratified population
asfr	ASFRs by district and time
areas	Area hierarchy
naomi_level	Area level to produce estimates at
project	Model will by default produce estimates up to year of last survey. Integer projection year if desired, else FALSE

make_rw_structure_matrix

Make random walk structure matrices

Description

Make random walk structure matrices

Usage

```
make_rw_structure_matrix(x, order, adjust_diagonal = TRUE)
```

Arguments

x	Matrix size
order	Random walk order
adjust_diagonal	Add 1E-6 to the matrix diagonal to make the matrix proper. Default = TRUE

sample_tmb

Sample TMB

Description

Sample TMB

Usage

```
sample_tmb(  
  fit,  
  nsample = 1000,  
  rng_seed = NULL,  
  random_only = TRUE,  
  verbose = FALSE  
)
```

transform_mics

Transform MICS dataframes

Description

Convert lists by survey to lists by dataset type

Usage

```
transform_mics(mics_survey_data, mics_indicators)
```

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