

R documentation

of all in
'/home/dorer/projects/census/PovertyAssessmentToolkit/source/PAT/man'

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Builtin.marginals	<i>Builtin Marginal Tables for Marginal Adjustment</i>
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Value

Marginal table function (ends in .m) to be used in the marginal.tables component of a model.

List of tables

AgeRaceSex.m	Age by Race by Sex Age given by Age.code Race by Race.code
AgeSex.m	Age by Sex Age given by Age.code Race by Race.code
Race.m	Number of categories given by PAT.race.code

HispanicAgeSex.m	Hispanic yes/no Age given by PAT.age.code
Poverty2.m	Poverty 2 categories: Below/Above/Undefined
Poverty4.m	Poverty 4 categories Under 100_200, 200_300, 300+, Undefined
Employed.m	Employed, Unemployed, Not_in Under_16
HouseType.m	Non_inst, Inst (Institutional), House (Household)
DisabilityRace.m	Disability Yes, No, Other, Race give by Race.code
Education.m	Under_18, Less_than_high_school, High_school Some_College, College_degree
Education4.m	Less_than_high_school, High_school, Some_college, College_degree
Tenure.m	Rent, Own, Group_quarters.
MaritalStatus3.m	Married, Single Mother, Single Father.
MaritalStatus3a.m	Married, Single, Under_15_years
MaritalStatus7.m	Married, Widowed, Separated, Divorced Never_married, Other, Under_15_years.
HealthIns3.m	Insured, Uninsured, Military_inst (Military or Institutional Group Quarters)
FamilyType3.m	Married_head, Female_head, Male_head, Other

Author(s)

David Dorer

Builtin.models

*Builtin Models***Value**

BrooklineI.model

marginal tables: AgeRaceSex, MaritalStatus3, HispanicAgeSex, Education4, Employed, Tenure, DisabilityRace, Poverty2, HouseType, FamilyType3, Health-Ins3

extra variables: SPM3 (Supplemental Poverty Measure)

PennsylvaniaI.model

marginal.tables: AgeRaceSex, Poverty4, Employed, Education4, HispanicAgeSex, MaritalStatus3, Tenure.

extra variables: WIC, Age6a, EmployedHouse18.

Author(s)

David Dorer

Builtin.variables	<i>Builtin Derived PUMS variables</i>
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Value

PUMS variable function (ends in .v) to be used in the variables component of a model.

List of variables

Age.v	Age arguments AGEP and nages
WIC.v	Does person live in a household that receives WIC benefits.
Race.v	Race with arguments RAC1P and nraces
SNAP.v	Does person live in a household that receives SNAP benefits.
Hispanic.v	Is the person of Hispanic Ethnicity ("Hispanic" or "No")
Employed.v	Is the person employed ("Under_16", "Employed", "Unemployed", "Not_in")
Tenure.v	"Group_quarters", "Own", "Rent".
HousingCost.v	"Low", "High", "Vacant".
Poverty2.v	Poverty Threshold "Below", "Above", "Undefined".
Poverty4.v	Poverty Threshold "Below", "100_to_199", "200_to_299", "Over_300", "Undefined".
HealthIns3.v	"Insured", "Uninsured", "Military_inst" (Military or Institutional Group Quarters).
Disability.v	"Yes", "No", "Other" (See table B18101)
HouseType.m	Non_inst, Inst (Institutional), House (Household)
Education4.m	"Less_than_high_school", "High_school", "Some_college", "College_degree"
FamilyType3.v	"Married_head", "Female_head", "Male_head", "Other".
FamilyType.v	"Married_head", "Female_head", "Male_head", "Child_group", "18_years_and_over".
MaritalStatus3a.m	Married, Single, Under_15_years

MaritalStatus7.m Married, Widowed, Separated, Divorced
 Never_married, Other, Under_15_years.

Author(s)

David Dorer

loglinf3

Loglinear Model fit with specified marginal tables

Description

Wrapper R function for compiled C IPF subroutine.

Usage

```
loglin3f(seed.table, target.list, target.data, niter=5, maxdev=0.001, debug=0)
```

Arguments

seed	Starting seed table for fit.
target.list	list with index specifying variable is seed table. Note the marginal.tables function computes this list from the model.
target.data	list containing the data for the marginal tables specified by target.list
niter	Maximum number of iterations. Same as iter in the PAT.synth.data function.
maxdev	Maximum relative change in difference between target marginals and current fit margin between successive iteration.
debug	Verbose/debug level for messages. Default PAT.verbose()

Details

The arguments are the same as the IpF function in mipfp package.

Value

seed	Argument passed to function.
p.hat	Model fit scaled to sum to 1.
nlast	Last iteration.
ifault	Did the fit converge 0: yes 1: no
maxdev	Maximum deviation at last iteration.

Author(s)

David Dorer

References

<https://cran.r-project.org/web/packages/mipfp/mipfp.pdf> <https://www.jstatsoft.org/article/view/v086c02>

PAT.acs.table	<i>Download an ACS Detail (B, S, or DP) Table</i>
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Description

Download and ACS table for a given geography.

Usage

```
PAT.acs.table()
```

Arguments

group	ACS table name or "group" (API term). For example the table B01001 has title "Sex by Age." Check out tables at data.census.gov.
state	State FIPS code. Required.
county	County FIPS code. Character variable - optional defaults to "".
tract	County FIPS code. Character variable (6 characters) - optional defaults to ""
blockgroup	Blockgroup FIPS code. Character variable (1 character) defaults to "" (ignored)
puma	PUMA FIPS code. 5 characters - optional defaults to ""
place	Place FIPS code. Defaults to "" which will be ignored.
csd	County Subdivision FIPS code - defaults to "" which will be ignored
vintage	Defaults to PAT.acs.vintage().
period	Defaults to PAT.acs.period().
ddir	character variable indicating where to store downloaded tables - default datadir().
debug	Level of messages to print - default PAT.verbose().
cache.metadata	Cache level for metadata 0: no caching, download from Census website for every table. 1: cache metadata in file 2: cache metadata in both a file and in the .GlobalEnv or computer memory. Default 2 speeds downloading and decoding many tables. If you seem to be having difficulties with stale cached data use 0. Default PAT.cache.metadata().
cache.tables	Cache level for ACS tables. 0: download table from census for each table. 1: cache tables in a file (best). 2: cache table in both a file and in computer memory. With a run of many tracts 2 will quickly exhaust memory. For testing where you are repeatedly downloading the same table you might temporarily use 2 to save time.

Details

Download B, S, DP (SF1) tables attaches names for rows using metadata, table has 2 columns: "Est" and "MoE". Various geographies can be specified. The function sorts out the geography if you use too many geography values.

Value

list list with 2 components "data" (matrix with table rownames and 2 columns "Est" and "MoE") and "par" various parameters such as the table "object" name, folder where the table was stored, name of the filename of where the table was stored, date time of download, etc.

Author(s)

David Dorer

References

To find, examine and check data, see Census webpage data.census.gov

PAT.age.race	<i>Set age and race categories</i>
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Description

PAT.age.code(code) Set the PAT.age.code option.
 PAT.race.code(code) Set the PAT.race code option.

Usage

```
PAT.set.ages("7a")
PAT.set.races("5")
```

Arguments

code	Age code
code	Race code

With no argument return the Age or Race Code

Details

Without an argument return current setting for Age or Race code.

Value

The race or age code.
 or
 Sets the corresponding Global race or age code option.

Values for age code

"7a" 7 age categories
14 and under, 15 to 19, 20 to 34, 35 to 44,
45 to 54, 55 to 64, 65 and over.

"7b" 7 age categories
under 5, 5 to 9, 15 to 17, 18 to 24,
25 to 64, 65 and over

Values for race code

"3" 3 race categories:
White, Black, Other

"5" 5 race categories
White, Black, Asian, Other, Two.

"7" 7 race categories
white, black, native, asian, Hawaiian, Other, Two.

Author(s)

David Dorer

References

xxx

PAT.convert.spm

Convert Supplementel Poverty Measure US SAS Data

Description

Converts sas7bdat binary file downloaded from Census website.

Usage

PAT.convert.spm(vintage="2021")

Arguments

vintage	Vintage for SPM SAS file.
ddir	Output data folder/directory. Default datadir().
download	Should sas7bdat file be downloaded. Default PAT.download().

Details

The data file is very large, typically 900 MB and download and conversion can take 5-10 minutes depending the the speed of your connection and your processor speed. Requires package require("sas7bdat") which may need to be installed install.packages("sas7bdat") On Linux sas7bdat often throws an error and the conversion step fails. The Windows package doesn't seem to have this problem.

Value

Output U.S. file spm_pu_<vintage>.RData.
For download url=
https://www2.census.gov/programs-surveys/supplemental-poverty-measure/datasets/spm/spm_pu_<vintage>.sas7bdat

Author(s)

David Dorer

PAT.merge.synth.data	<i>Merge Synthetic Tract Files</i>
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Description

Zip individual tract files and merge into a single csv file.

Usage

```
PAT.merge.synth.data(state="25",ddir=datadir(),odir=outdir(),zip=TRUE,outtag="_test")
```

Arguments

state	State FIPS code.
ddir	Data directory for output zip and output csv file.
odir	Directory with input csv tract files.
zip	Zip tract files before merging.
outtag	Tag to append to file names.

Details

xxx

Value

zip file	Output zip file synth_data_<state><outtag>.zip contains individual tract csv files.
csv file	Merged csv data file synth_data_<state><outtag>.csv contains stacked individual tract csv files.

Author(s)

David Dorer

PAT.model

Statistical Models for Synthetic Data

Description

A model with PUMS variables and marginal tables.

Usage

See BrooklineI.model and PennsylvaniaI.model for examples

List Elements

name	character variable with name for model, e.g. "BrooklineI"
variables	A named list with functions that define the variables.
marginal.tables	A named list with functions that define the marginal tables. See PAT.acs.table for required arguments.
parameters	A named list with model parameters:
model	name of model. Same as name component.
nages	the number of age categories for model (see PAT.nage.codes)
nraces	the number of race categories for model (see PAT.nrace.codes)
geotype	type of geography "tract" or "blockgroup".
comment	comment (any text).
...	optional user defined parameters that can be passed to the model variables and marginal.tables functions.

Author(s)

David Dorer

PAT.root

Set root directory

Description

Set root folder/directory.

Usage

junk

Arguments

root	Value to set root directory. NA (default) return root directory.
default	Default Value to set root directory. Default for default getwd()

Details

xxx

Value

root	current value of root directory/folder
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Author(s)

David Dorer

PAT.synth.data

*Synthesize Data***Description**

Run a synthetic data model using a PUMA and set of marginal tables

Usage

```
PAT.synth.data(model="BrooklineIII",state="25",puma="03400",outtag="_test",iter=30)
```

Arguments

model	Model see vignette("CreatingModels")
state	State FIPS code
puma	PUMA FIPS code
outtag	A tag to include in some output file names
iter	Maximum number of IPF iterations
vers	Version of IPF program 1:Internal C code 2:mipfp package Ipfp function
maxdev	Termination criteria for relative change in marginal table deviation from one iteration to the next. Default 0.001
add	Add a small amount to every cell of seed/model table. This allows the data to take on non-zero weights even though the weight is zero in the PUMS data. Default 0.0
minwt	Records with weights < minwt will be dropped from the synthetic data. Default 0.001
update	Name of input checkpoint file. "" no input check point file Geographies in checkpoint file are skipped
county	Vector of county FIPS codes. Default character(0). For the default all the tracts in county will be run.
tract	Vector of tract FIPS codes Default character(0). For the default only tracts in tract will be run for county[1]

bdir	base directory for program run. Default PAT.root()
odir	Output directory. Tract synthetic files go here. Default PAT.root()/output/
ddir	Data directory. Cached ACS table go here. Various data files are stored here. For example the cross walk files. Default PAT.root()/data/
logfile	File to log messages. Default synth_<state>_checkpoint<outtag>.txt.
vintage	ACS vintage for marginal tables default: PAT.acs.vintage()
checkfile	Output check point file. As tract output files are written a record with the geography is appended to this file.
download	Should marginal tables be downloaded? Default PAT.download()
pums.vintage	PUMS/PUMA vintage default: PAT.pums.vintage()
pums.period	PUMS/PUMA period default: PAT.pums.period()
download	download files. Default: PAT.download().
key	Census key. Default PAT.census.key().
debug	Verbose/debug level for printing messages. Default PAT.verbose()
dump.seed	Dump the PUMA seed table/dataset in a file. 0 no dump 1 dump. Default 0

Value

A log of the run. The other effects are the output files.

synthetic data The output file go in odir with names synth_data_<state_FIPS>_<puma_FIPS>_<county_FIPS>_<tract_FIPS>.txt. The file has 3 initial comment lines with the output file name, last iteration, number of iterations (argument), convergence flag (1:yes 2:no) nages and nraces. The 4th row is a header record. Each data row has header variables:

state	State FIPS
puma	PUMA FIPS
county	County FIPS
tract	Tract FIPS
blockgroup	Blockgroup FIPS, set to "".
model	Model name
type	person or house.
ages	age parameter e.g. "9", "7", "7a".
races	race parameter, "3", "5", "7".
date	date-time when record was written.
variables ...	columns with the value of the model variables.

Check point file

synth_<state>_checkpoint<outtag>.txt

The file had the state, puma,count,tract and additional information. By using the check point file previously completed tracts will be skipped. Useful when you are doing an entire stage and your computer updates and reboots. The file is appended to so if you want to start over but still use the update feature delete the check point file before you start a clean run.

Tract files The synthetic data for each tract is put in a separate file in folder odir.

log file Folder logdir() file name log<outtag>.txt

Author(s)

David Dorer 8 Dec 2023 14:41

References

vignette("CreateModels") vignette("SynthesizeData")

PAT.test.model	Test PUMS model
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Usage

PAT.test.model(model="BrooklineIII")

Arguments

model	model or model name (with or without quotes)
state	state FIPS
puma	PUMA FIPS
key	census key (default PAT.census.key())
vintage	marginal tables vintage (default PAT.acs.vintage())
period	marginal tables period (default PAT.acs.period())
download	download data 1:download 0:used cached data (default PAT.download())
debug	debug/message level - higher more messages 0:no messages 1+: more messages (default PAT.verbose())

Details

Tests model to be used for synthesizing data.

Value

list with components:

marginals	marginal tables/targets ("B","S","DP" PUMA geography table)
variables	one-way PUMS/PUMA frequency (weighted) of PUMA model table for variables used in marginal tables
model.vars	one-way PUMS/PUMA frequency tables for variables not used in marginal tables ("carry along" variables)
total.pop	total population for marginal tables.
puma.population	total population for PUMA PUMA data (sum of PUMS weights)
par	various parameters state FIPS, PUMA FIPS, detail tables vintage & period, PUMS vintage & period.

Author(s)

David Dorer

References

PAT.model

PUMA.2012.Tract.2020 *PUMA (2012) to Tract (2020) Relationship File*

Description

PUMA to Tract Correlation/Relationship Correspondence Dataset

Usage

`data(PUMA.2020.Tract.2020)` `data(PUMA.2012.Tract.2020)`

Format

88865 observations (2012 PUMAs)
or 85395 observations (2020 PUMAs)

Value

A data.frame PUMA.2012.Tract.2020 or PUMA.2020.Tract.2020

Variable names

State	State FIPS code.
Puma	PUMA FIPS code.
County	County FIPS code.
Tract	Tract FIPS code.
StateAb	State Postal Abbreviation.
PumaName	County name.
PumaName	PUMA name.
Allocation	Fraction of Tract intersecting PUMA.

References

Missouri Census Data Center Geocorr Engine.
<https://mcdc.missouri.edu/applications/geocorr2022.html>
 Census Public Use Microdata Areas (PUMAs) webpage. [2020_Census_Tract_to_2020_PUMA.txt](#)

SPM_2021_25_data*Supplemental Poverty Measure Data for Massachusetts*

Description

Census 2021 SPM data with records for Massachusetts FIPS 25

Usage

```
data(SPM.2021.25.data)
```

Format

A data.frame with 67951 observations and 46 variables.

Value

An object SPM.2020.25.data

References

[Codebook](#)

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