

Package ‘RNHANES’

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Type Package

Title Facilitates Analysis of CDC NHANES Data

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URL <http://github.com/silentspringinstitute/RNHANES>

BugReports <https://github.com/silentspringinstitute/RNHANES/issues>

Description Tools for downloading and analyzing CDC NHANES data, with a focus on analytical laboratory data.

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LazyData TRUE

Depends R (>= 2.10)

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Suggests testthat, knitr, rmarkdown

VignetteBuilder knitr

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Contents

RNHANES-package	2
demography_filename	2
download_nhances_file	3
file_suffix	3
load_nhances_description	4

nhanes_analyze	4
nhanes_cycle_years	5
nhanes_data_files	5
nhanes_detection_frequency	6
nhanes_hist	7
nhanes_load_data	8
nhanes_load_demography_data	9
nhanes_quantile	10
nhanes_sample_size	11
nhanes_search	11
nhanes_survey	12
nhanes_survey_design	13
nhanes_variables	14
nhanes_vcov	15
process_file_name	16
validate_year	16

Index	17
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RNHANES-package

RNHANES simplifies downloading and analyzing NHANES data.

Description

RNHANES simplifies downloading and analyzing NHANES data.

demography_filename

Translates cycle years into the correct demography filename suffix, e.g. '2001-2002' returns 'B'

Description

Translates cycle years into the correct demography filename suffix, e.g. '2001-2002' returns 'B'

Usage

```
demography_filename(year)
```

Arguments

year	NHANES cycle, e.g. "2001-2002"
------	--------------------------------

Value

suffix character e.g. "B"

download_nhances_file *Download an NHANES data file from a given cycle*

Description

Download an NHANES data file from a given cycle

Usage

```
download_nhances_file(file_name, year, destination = tempdir(), cache = TRUE)
```

Arguments

file_name	file name
year	NHANES cycle
destination	directory to download the file into
cache	whether to cache the file

Value

path to the downloaded file

file_suffix *Returns the NHANES file suffix for the given year*

Description

Returns the NHANES file suffix for the given year

Usage

```
file_suffix(year)
```

Arguments

year	NHANES cycle year (e.g. "2001-2002")
------	--------------------------------------

Value

suffix character (e.g. "B" or "C")

load_nhances_description*Download an NHANES description file***Description**

Download an NHANES description file

Usage

```
load_nhances_description(file_name, year, destination = tempdir(),
                        cache = FALSE)
```

Arguments

<code>file_name</code>	file name
<code>year</code>	NHANES cycle
<code>destination</code>	directory to download the file into
<code>cache</code>	whether to cache the file

Value

data frame containing the file description

nhanes_analyze*Compute quantiles from NHANES weighted survey data***Description**

Compute quantiles from NHANES weighted survey data

Usage

```
nhanes_analyze(analysis_fun, nhanes_data, column, comment_column = "",
               weights_column = "", filter = NULL)
```

Arguments

<code>analysis_fun</code>	function to use to analyze each variable
<code>nhanes_data</code>	data frame containing NHANES data
<code>column</code>	column name of the variable to compute quantiles for
<code>comment_column</code>	comment column name of the variable
<code>weights_column</code>	name of the weights column
<code>filter</code>	logical expression used to subset the data

Value

a data frame

nhanes_cycle_years *List the valid NHANES cycle years*

Description

List the valid NHANES cycle years

Usage

```
nhanes_cycle_years()
```

Value

vector of NHANES cycle years

nhanes_data_files *List the NHANES data files*

Description

List the NHANES data files

Usage

```
nhanes_data_files(components = "all", destination = tempfile(),  
cache = TRUE)
```

Arguments

components	one of "all", "demographics", "dietary", "examination", "laboratory", "questionnaire"
destination	destinatino to save the file lists
cache	whether to cache the downloaded file lists so they don't have to be re-downloaded every time

Value

data frame of NHANES data files available to download

Examples

```
## Not run:

# Download a data frame of all the NHANES data files
files <- nhanes_data_files()

# Download a data frame of just the laboratory files
lab_files <- nhanes_data_files(component = "laboratory")

## End(Not run)
```

nhanes_detection_frequency

Compute detection frequencies of NHANES data

Description

Compute detection frequencies of NHANES data

Usage

```
nhanes_detection_frequency(nhanes_data, column, comment_column,
                           weights_column = "", filter = NULL)
```

Arguments

<i>nhanes_data</i>	data frame containing NHANES data
<i>column</i>	column names of the variables to compute detection frequencies for
<i>comment_column</i>	comment column names of the variables to compute detection frequencies for
<i>weights_column</i>	sample weight column
<i>filter</i>	logical expression used to subset the data

Value

named vector of detection frequencies

Examples

```
## Not run:
dat <- nhanes_load_data("UHG_G", "2011-2012", demographics = TRUE)

# Compute detection frequency
nhanes_detection_frequency(dat, c("URXUHG"), c("URDUHGLC"))

## End(Not run)
```

nhanes_hist*Plot a weighted histogram of an NHANES variable*

Description

Plot a weighted histogram of an NHANES variable

Usage

```
nhanes_hist(nhanes_data, column, comment_column, weights_column = "",  
            filter = "", transform = "", ...)
```

Arguments

nhanes_data	data frame containing NHANES data
column	column name of the variable to plot
comment_column	comment column of the variable to plot
weights_column	name of the weights column
filter	logical expression used to subset the data
transform	transformation to apply to the column. Accepts any function name, for example: "log"
...	parameters passed through to svyhist function

Value

a data frame

Examples

```
## Not run:  
dat <- nhanes_load_data("PFC_G", "2011-2012", demographics = TRUE)  
  
nhanes_hist(dat, "LBXPFOA")  
  
## End(Not run)
```

`nhanes_load_data` *Download NHANES data files.*

Description

Download NHANES data files.

Usage

```
nhanes_load_data(file_name, year, destination = tempdir(),
  demographics = FALSE, cache = TRUE, recode = FALSE,
  recode_data = FALSE, recode_demographics = FALSE,
  allow_duplicate_files = FALSE)
```

Arguments

<code>file_name</code>	NHANES file name (e.g. "EPH") or a vector of filenames (e.g c("EPH", "GHB"))
<code>year</code>	NHANES cycle year (e.g. "2007-2008") or a vector of cycle years
<code>destination</code>	directory to download the files to
<code>demographics</code>	include demographics data into the dataset
<code>cache</code>	whether to cache the file to disk
<code>recode</code>	whether to recode the data and demographics (overrides other parameters)
<code>recode_data</code>	whether to recode just the data
<code>recode_demographics</code>	whether to recode just the demographics
<code>allow_duplicate_files</code>	how to handle a request that has duplicate file names/cycle years. By default duplicates will be removed.

Details

If you supply vectors for both `file_name` and `year`, then the vectors are paired and each `file_name/year` pair is downloaded. For example, `file_name = c("EPH", "GHB")`, `year = c("2009-2010", "2011-2012")` will download "EPH_F.XPT" and "EPH_G.XPT". In other words, the function does not download every possible combination of `file_name` and `year`.

You can specify file names in several formats. In order of specificity: You can supply the complete filename: "EPH_F.XPT" You can supply the filename without an extension: "EPH_F" You can supply the filename without a suffix: "EPH", `year = "2009-2010"`

If you are loading the same file across multiple years, you must supply the filename without a suffix so that the correct suffix for each year can be used.

This function returns either a list or a data frame. If you load multiple files, the return value will always be a list. This is because the columns may not match in between files. If you load one file, the result will be a data frame.

Value

if file_name or year is a vector, returns a list containing a data frame for each file_name. If file_name and year are both singletons, then a data frame is returned.

Examples

```
## Not run:  
  
nhanes_load_data("UHG", "2011-2012")  
  
# Load data with demographics  
nhanes_load_data("UHG", "2011-2012", demographics = TRUE)  
  
# Download to /tmp directory and overwrite the file if it already exists  
nhanes_load_data("HDL_E", "2007-2008", destination = "/tmp", cache = FALSE)  
  
## End(Not run)
```

nhanes_load_demography_data

Download NHANES demography files for a specific cycle.

Description

Download NHANES demography files for a specific cycle.

Usage

```
nhanes_load_demography_data(year, destination = tempdir(), cache = FALSE)
```

Arguments

year	NHANES cycle year (e.g. "2011-2012")
destination	directory to download the file to
cache	whether load the file if it already exists on disk

Examples

```
## Not run:  
nhanes_load_demography_data("2011-2012")  
  
## End(Not run)
```

<i>nhanes_quantile</i>	<i>Compute quantiles from NHANES weighted survey data</i>
------------------------	---

Description

Compute quantiles from NHANES weighted survey data

Usage

```
nhanes_quantile(nhanes_data, column, comment_column = "",  
  weights_column = "", quantiles = seq(0, 1, 0.25), filter = NULL)
```

Arguments

nhanes_data	data frame containing NHANES data
column	column name of the variable to compute quantiles for
comment_column	comment column name of the variable for checking if computed quantiles are below the LOD
weights_column	name of the weights column
quantiles	numeric or vector numeric of quantiles to compute
filter	logical expression used to subset the data

Value

a data frame

Examples

```
## Not run:  
dat <- nhanes_load_data("UHG_G", "2011-2012", demographics = TRUE)  
  
# Compute 50th, 95th, and 99th quantiles  
nhanes_quantile(dat, "URXUHG", "URDUHGLC", "WTSA2YR", c(0.5, 0.95, 0.99))  
  
## End(Not run)
```

`nhanes_sample_size` *Compute the sample size of NHANES data*

Description

Compute the sample size of NHANES data

Usage

```
nhanes_sample_size(nhanes_data, column, comment_column = "",  
                   weights_column = "", filter = NULL)
```

Arguments

<code>nhanes_data</code>	data frame containing NHANES data
<code>column</code>	column name of the variable to compute quantiles for
<code>comment_column</code>	comment column name of the variable for checking if computed quantiles are below the LOD
<code>weights_column</code>	name of the weights column
<code>filter</code>	logical expression used to subset the data

Value

a data frame

Examples

```
## Not run:  
dat <- nhanes_load_data("UHG_G", "2011-2012", demographics = TRUE)  
  
nhanes_sample_size(dat, "URXUHG", "URDUHGLC")  
  
## End(Not run)
```

`nhanes_search` *Search the results from nhanes_variables or nhanes_data_files*

Description

Search the results from nhanes_variables or nhanes_data_files

Usage

```
nhanes_search(nhanes_data, query, ..., fuzzy = FALSE, ignore_case = TRUE,  
             max_distance = 0.2)
```

Arguments

<code>nhanes_data</code>	nhanes variable list, from <code>nhanes_variables</code> function, or data file list, from <code>nhanes_data_files</code>
<code>query</code>	regular expression search query
<code>...</code>	additional arguments to pass to <code>dplyr::filter</code>
<code>fuzzy</code>	whether to use fuzzy string matching for search (based on edit distances)
<code>ignore_case</code>	whether search query is case-sensitive
<code>max_distance</code>	parameter for tuning fuzzy string matching, 0-1

Value

data frame filtered by search query

Examples

```
## Not run:
nhanes_files <- nhanes_data_files()

# Search for data files about pesticides
nhanes_search(nhanes_files, "pesticides")

## End(Not run)
```

`nhanes_survey`

Apply a function from the survey package to NHANES data

Description

Apply a function from the survey package to NHANES data

Usage

```
nhanes_survey(survey_fun, nhanes_data, column, comment_column = "",  
  weights_column = "", filter = NULL, analyze = "values",  
  callback = NULL, ...)
```

Arguments

<code>survey_fun</code>	the survey package function (e.g. <code>svyquantile</code> or <code>svymean</code>)
<code>nhanes_data</code>	data frame containing NHANES data
<code>column</code>	column name of the variable to compute quantiles for
<code>comment_column</code>	comment column name of the variable
<code>weights_column</code>	name of the weights column
<code>filter</code>	logical expression used to subset the data

analyze	one of "values" or "comments", whether to apply the survey function to the value or comment column.
callback	optional function to execute on each row of the dataframe
...	other arguments to pass to the survey function

Details

This function provides a generic way to apply any function from the survey package to NHANES data. RNHANES provides specific wrappers for computing quantiles (nhanes_quantile) and detection frequencies (nhanes_detection_frequency), and this function provides a general way to use any survey function.

Value

a data frame

Examples

```
## Not run:
library(survey)

nhanes_data <- nhanes_load_data("EPH", "2011-2012", demographics = TRUE)

# Compute the mean of triclosan using the svymean function
nhanes_survey(svymean, nhanes_data, "URXTRS", "URDTRSLC", na.rm = TRUE)

# Compute the variance using svyvar
nhanes_survey(svyvar, nhanes_data, "URXTRS", "URDTRSLC", na.rm = TRUE)

## End(Not run)
```

nhanes_survey_design *Build survey objects for NHANES data*

Description

Build survey objects for NHANES data

Usage

```
nhanes_survey_design(nhanes_data, weights_column = "")
```

Arguments

nhanes_data	data frame containing NHANES data
weights_column	name of the weights column

Value

a survey design object

Examples

```
## Not run:
dat <- nhanes_load_data("UHG_G", "2011-2012", demographics = TRUE)

design <- nhanes_survey_design(dat, "WTSA2YR")

svymean(~RIDAGEYR, design)

svyglm(URXUHG ~ RIDAGEYR + RIAGENDR, design)

## End(Not run)
```

nhanes_variables

Load the NHANES comprehensive variable list

Description

Load the NHANES comprehensive variable list

Usage

```
nhanes_variables(components = "all", destination = tempfile(),
cache = TRUE)
```

Arguments

components	one of "all", "demographics", "dietary", "examination", "laboratory", "questionnaire"
destination	where to save the variable list
cache	whether to cache the downloaded variable list so it doesn't have to be re-downloaded every time
	Helper function for <i>nhanes_variables</i> function

Value

dat

Examples

```
## Not run:

# Download the comprehensive NHANES variable list
variables <- nhanes_variables()

# Download the variable list and cache it in a specific file
variables <- nhanes_variables(destination = "./nhanes_data")

## End(Not run)
```

nhanes_vcov

Extract variance/covariance matrix from parameters of svymean

Description

Extract variance/covariance matrix from parameters of svymean

Usage

```
nhanes_vcov(nhanes_data, columns, weights_column = "", filter = "")
```

Arguments

nhanes_data	data frame containing NHANES data
columns	columns to include in svymean for
weights_column	name of the weights column
filter	logical expression used to subset the data

Value

a data frame

Examples

```
## Not run:
dat <- nhanes_load_data("PFC_G", "2011-2012", demographics = TRUE)

nhanes_vcov(dat, c("LBXPFOA", "LBXPFOS"))

## End(Not run)
```

process_file_name	<i>Processes a file name to make sure it is valid and has the correct suffix and extension File names with an extension (e.g. ".XPT") are not altered</i>
-------------------	---

Description

Processes a file name to make sure it is valid and has the correct suffix and extension File names with an extension (e.g. ".XPT") are not altered

Usage

```
process_file_name(file_name, year, extension = ".XPT")
```

Arguments

file_name	name of the file
year	NHANES cycle year
extension	file extension

validate_year	<i>Check that the year is in the correct format e.g. '2001-2002' is correct and returns TRUE, '2001' is not correct and returns FALSE</i>
---------------	---

Description

Check that the year is in the correct format e.g. '2001-2002' is correct and returns TRUE, '2001' is not correct and returns FALSE

Usage

```
validate_year(year, throw_error = TRUE)
```

Arguments

year	the year or years to validate
throw_error	whether to throw an error if the year is invalid

Index

demography_filename, 2
download_nhances_file, 3

file_suffix, 3

load_nhances_description, 4

nhances_analyze, 4
nhances_cycle_years, 5
nhances_data_files, 5
nhances_detection_frequency, 6
nhances_hist, 7
nhances_load_data, 8
nhances_load_demography_data, 9
nhances_quantile, 10
nhances_sample_size, 11
nhances_search, 11
nhances_survey, 12
nhances_survey_design, 13
nhances_variables, 14
nhances_vcov, 15

process_file_name, 16

RNHANES (RNHANES-package), 2
rnhanes (RNHANES-package), 2
RNHANES-package, 2
rnhanes-package (RNHANES-package), 2

validate_year, 16