

Package ‘EdSurvey’

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Description

Read in and analysis functions for education survey and assessment data from the National Center for Education Statistics (NCES) <<https://nces.ed.gov/>>, including National Assessment of Educational Progress (NAEP) data <<https://nces.ed.gov/nationsreportcard/>> and data from the International Assessment Database: OECD <<http://www.oecd.org/>>, including PISA, TALIS, PIAAC, and IEA <<http://www.iea.nl/>>, including TIMSS, TIMSS Advanced, PIRLS, ICCS, ICILS, and CivEd.

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EdSurvey-package	<i>Analysis of NCES Education Survey and Assessment Data</i>
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Description

The EdSurvey package uses appropriate methods for analyzing NCES datasets with a small memory footprint. Existing system control files, included with the data, are used to read in and format the data for further processing.

Details

To get started using EdSurvey, see the vignettes for tutorials and the statistical methodologies. Use `vignette("introduction", package="EdSurvey")` to see the vignettes.

The package provides functions called `readNAEP`, `readCivEDICCS`, `readICILS`, `readPIAAC`, `readPIRLS`, `read_ePIRLS`, `readPISA`, `readTALIS`, `readTIMSS`, `readTIMSSAdv`, and `readECLS_K2011` to read in NCES datasets. The functions `achievementLevels`, `cor.sdf`, `edsurveyTable`, `lm.sdf`, `logit.sdf`, `percentile`, and `gap` can then be used to analyze data. For advanced users, `getData` extracts the data of interest as a data frame for further processing.

achievementLevels *Achievement Levels*

Description

Returns achievement levels using weights and variance estimates appropriate for the `edsurvey.data.frame`.

Usage

```
achievementLevels(achievementVars = NULL, aggregateBy = NULL, data,
  cutpoints = NULL, returnDiscrete = TRUE, returnCumulative = FALSE,
  weightVar = NULL, jrrIMax = 1, omittedLevels = TRUE,
  defaultConditions = TRUE, recode = NULL, returnNumberOfPSU = FALSE,
  returnVarEstInputs = FALSE)
```

Arguments

achievementVars	character vector indicating variables to be included in the achievement levels table, potentially with a subject scale or subscale. When the subject scale or subscale is omitted, the default subject scale or subscale is used. You can find the default composite scale and all subscales using the function showPlausibleValues .
aggregateBy	character vector specifying variables to aggregate achievement levels by. The percentage column sums up to 100 for all levels of all variables specified here. When set to the default of NULL, the percentage column sums up to 100 for all levels of all variables specified in achievementVars.
data	an <code>edsurvey.data.frame</code>
cutpoints	numeric vector indicating cutpoints. Set to standard NAEP cutpoints for Basic, Proficient, and Advanced by default.
returnDiscrete	logical indicating if discrete achievement levels should be returned. Defaults to TRUE.
returnCumulative	logical indicating if cumulative achievement levels should be returned. Defaults to FALSE. The first and last categories are the same as defined for discrete levels.
weightVar	character string indicating the weight variable to use. Note that only the name of the weight variable needs to be included here, and any replicate weights will be automatically included. When this argument is NULL, the function uses the default. Use showWeights to find the default.
jrrIMax	numeric value. When using the jackknife variance estimation method, the V_{jrr} term (see Details) can be estimated with any positive number of plausible values and is estimated on the first of the lower of the number of available plausible values and <code>jrrIMax</code> . When <code>jrrIMax</code> is set to <code>Inf</code> , all plausible values will be used. Higher values of <code>jrrIMax</code> lead to longer computing times and more accurate variance estimates.

omittedLevels	a logical value. When set to the default value (TRUE), it drops those levels in all factor variables that are specified in achievementVars and aggregateBy. Use print on an edsurvey.data.frame to see the omitted levels.
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in an edsurvey.data.frame to subset the data. Use print on an edsurvey.data.frame to see the default conditions.
recode	a list of lists to recode variables. Defaults to NULL. Can be set as recode = list(var1= list(from=c("a", "b", "c"), to ="d")). See Examples.
returnNumberOfPSU	a logical value set to TRUE to return the number of primary sampling units (PSU)
returnVarEstInputs	a logical value set to TRUE to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for the computation of covariances between estimates.

Details

The achievementLevels function applies appropriate weights and the variance estimation method for each edsurvey.data.frame, with several arguments for customizing the aggregation and output of the analysis results. Namely, by using these optional arguments, users can choose to generate the percentage of students performing at each achievement level (discrete), generate the percentage of students performing at or above each achievement level (cumulative), calculate the percentage distribution of students by achievement level (discrete or cumulative) and selected characteristics (specified in aggregateBy), and compute the percentage distribution of students by selected characteristics within a specific achievement level.

Calculation of percentages: The details of the methods are shown in the vignette titled [Statistics](#) in “Estimation of Weighted Percentages When Plausible Values Are Present” and are used to calculate all cumulative and discrete probabilities.

When the requested achievement levels are discrete (returnDiscrete = TRUE), the percentage \mathcal{A} is the percentage of students (within the categories specified in aggregateBy) whose scores lie in the range $[cutPoints_i, cutPoints_{i+1})$, $i = 0, 1, \dots, n$. cutPoints is the score thresholds provided by the user with $cutPoints_0$ taken to be 0. cutPoints are set to NAEP standard cut-points for achievement levels by default. To aggregate by a specific variable, for example, dsex, specify dsex in aggregateBy and all other variables in achievementVars. To aggregate by subscale, specify the name of the subscale (e.g., num_oper) in aggregateBy and all other variables in achievementVars.

When the requested achievement levels are cumulative (returnCumulative = TRUE), the percentage \mathcal{A} is the percentage of students (within the categories specified in aggregateBy) whose scores lie in the range $[cutPoints_i, \infty)$, $i = 1, 2, \dots, n - 1$. The first and last categories are the same as defined for discrete levels.

Calculation of standard error of percentages: The method used to calculate the standard error of the percentages is described in the vignette titled [Statistics](#) in the sections “Estimation of the Standard Error of Weighted Percentages When Plausible Values Are Present, Using the Jackknife Method” and “Estimation of the Standard Error of Weighted Percentages When Plausible Values Are Not Present, Using the Taylor Series Method.” For “Estimation of the Standard Error of

Weighted Percentages When Plausible Values Are Present, Using the Jackknife Method,” the value of `jrrIMax` sets the value of m^* .

Value

A list containing up to two data frames, one for each of the discrete and cumulative achievement levels as determined by `returnDiscrete` and `returnCumulative`. The data.frame contains the following columns:

Level	one row for each level of the specified achievement cutpoints
Variables in achievementVars	one column for each variable in <code>achievementVars</code> and one row for each level of each variable in <code>achievementVars</code>
Percent	the percentage of students at or above each achievement level aggregated as specified by <code>aggregateBy</code>
StandardError	the standard error of the percentage, accounting for the survey sampling methodology. See the vignette titled Statistics .
N	the number of observations in the incoming data (the number of rows when <code>omittedLevels</code> and <code>defaultConditions</code> are set to FALSE).
wtdN	the weighted number of observations in the data
nPSU	the number of primary sampling units (PSUs) at or above each achievement level aggregated as specified by <code>aggregateBy</code> . Only returned with <code>returnNumberOfPSU=TRUE</code> .

Author(s)

Huade Huo, Ahmad Emad, and Trang Nguyen

References

Rubin, D. B. (1987). *Multiple imputation for nonresponse in surveys*. New York, NY: Wiley.

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# Discrete achievement levels
achievementLevels(achievementVars=c("composite"), aggregateBy=NULL, data=sdf)

# Discrete achievement levels with a different subscale
achievementLevels(achievementVars=c("num_oper"), aggregateBy=NULL, data=sdf)

# Cumulative achievement levels
achievementLevels(achievementVars=c("composite"), aggregateBy=NULL, data=sdf,
                  returnCumulative=TRUE)

# Cumulative achievement levels with a different subscale
achievementLevels(achievementVars=c("num_oper"), aggregateBy=NULL, data=sdf,
```

```

        returnCumulative=TRUE)

# Achievement levels as independent variables, by sex aggregated by composite
achievementLevels(achievementVars=c("composite", "dsex"), aggregateBy="composite",
                  data=sdf, returnCumulative=TRUE)

# Achievement levels as independent variables, by sex aggregated by sex
achievementLevels(achievementVars=c("composite", "dsex"), aggregateBy="dsex",
                  data=sdf, returnCumulative=TRUE)

# Achievement levels as independent variables, by race aggregated by race
achievementLevels(achievementVars=c("composite", "sdracem"),
                  aggregateBy="sdracem", data=sdf, returnCumulative=TRUE)

# Use customized cutpoints
achievementLevels(achievementVars=c("composite"), aggregateBy=NULL, data=sdf,
                  cutpoints = c("Customized Basic" = 200,
                                "Customized Proficient" = 300,
                                "Customized Advanced" = 400))

# Use recode to change values for specified variables:
achievementLevels(achievementVars=c("composite", "dsex", "b017451"),
                  aggregateBy = "dsex", sdf,
                  recode=list(b017451=list(from=c("Never or hardly ever",
                                                  "Once every few weeks",
                                                  "About once a week"),
                                          to="Infrequently"),
                              b017451=list(from=c("2 or 3 times a week",
                                                  "Every day"),
                                          to="Frequently")))

## End(Not run)

```

as.data.frame

Coerce to a Data Frame

Description

Function to coerce a `light.edsurvey.data.frame` to a `data.frame`.

Usage

```
## S3 method for class 'light.edsurvey.data.frame'
as.data.frame(x, ...)
```

Arguments

`x` a `light.edsurvey.data.frame`
`...` other arguments to be passed to [as.data.frame](#)

Value

a `data.frame`

Author(s)

Trang Nguyen

cbind

Combine R Objects by Rows or Columns

Description

Implements `cbind` and `rbind` for `light.edsurvey.data.frame` class. It takes a sequence of vector, matrix, `data.frame`, or `light.edsurvey.data.frame` arguments and combines by columns or rows, respectively.

Usage

```
cbind(..., deparse.level = 1)
```

```
rbind(..., deparse.level = 1)
```

Arguments

`...` one or more objects of class vector, `data.frame`, matrix, or `light.edsurvey.data.frame`

`deparse.level` integer determining under which circumstances column and row names are built from the actual arguments. See `cbind`.

Details

Because `cbind` and `rbind` are standard generic functions that do not use method dispatch, we set this function as generic, which means it overwrites `base::cbind` and `base::rbind` on loading. If none of the specified elements are of class `light.edsurvey.data.frame`, the function will revert to the standard base method. However, to be safe, you might want to explicitly use `base::cbind` when needed after loading the package.

Note that the returned object will contain attributes only from the first `light.edsurvey.data.frame` object in the call to `cbind.light.edsurvey.data.frame`.

Value

a matrix-like object like matrix or `data.frame`. Returns a `light.edsurvey.data.frame` if there is at least one `light.edsurvey.data.frame` in the list of arguments.

Author(s)

Trang Nguyen, Michael Lee, and Paul Bailey

See Also

cbind

contourPlot

*Overlaid Scatter and Contour Plots***Description**

Diagnostic plots for regressions can become too dense to interpret. This function helps by adding a contour plot over the points to allow the density of points to be seen, even when an area is entirely covered in points.

Usage

```
contourPlot(x, y, m = 30L, xrange, yrange, xkernel, ykernel,
            nlevels = 9L, ...)
```

Arguments

x	numeric vector of the x data to be plotted
y	numeric vector of the y data to be plotted
m	integer value of the number of x and y grid points
xrange	numeric vector of length two indicating x-range of plot; defaults to range(x)
yrange	numeric vector of length two indicating y-range of plot. defaults to range(y)
xkernel	numeric indicating the standard deviation of Normal x kernel to use in generating contour plot
ykernel	numeric indicating the standard deviation of Normal y kernel to use in generating contour plot
nlevels	integer with the number of levels of the contour plot
...	additional arguments to be passed to a plot call that generates the scatter plot and the contour plot

Author(s)

Yuqi Liao and Paul Bailey

Examples

```
## Not run:
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))
lm1 <- lm.sdf(composite ~ pared * dsex + sdracem, sdf)
# plot the results
contourPlot(x=lm1$fitted.values,
            y=lm1$residuals[,1], # use only the first plausible value
            m=30,
```

```

        xlab="fitted values",
        ylab="residuals",
        main="Figure 1")
# add a line indicating where the residual is zero
abline(0,0)

## End(Not run)

```

cor.sdf

*Bivariate Correlation***Description**

Computes the correlation of two variables on an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`. The correlation accounts for plausible values and the survey design.

Usage

```

cor.sdf(x, y, data, method = c("Pearson", "Spearman", "Polychoric",
  "Polyserial"), weightVar = "default", reorder = NULL,
  omittedLevels = TRUE, defaultConditions = TRUE, recode = NULL,
  condenseLevels = TRUE)

```

Arguments

<code>x</code>	a character variable name from the data to be correlated with <code>y</code>
<code>y</code>	a character variable name from the data to be correlated with <code>x</code>
<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
<code>method</code>	a character string indicating which correlation coefficient (or covariance) is to be computed. One of <code>Pearson</code> (default), <code>Spearman</code> , <code>Polychoric</code> , or <code>Polyserial</code> .
<code>weightVar</code>	character indicating the weight variable to use. See Details.
<code>reorder</code>	a list of variables to reorder. Defaults to <code>NULL</code> (no variables are reordered). Can be set as <code>reorder = list(var1 = c("a", "b", "c"), var2 = c("4", "3", "2", "1"))</code> . See Examples.
<code>omittedLevels</code>	a logical value. When set to the default value of <code>TRUE</code> , drops those levels of all factor variables that are specified in an <code>edsurvey.data.frame</code> . Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels.
<code>defaultConditions</code>	a logical value. When set to the default value of <code>TRUE</code> , uses the default conditions stored in an <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.
<code>recode</code>	a list of lists to recode variables. Defaults to <code>NULL</code> . Can be set as <code>recode = list(var1 = list(from = c("a", "b", "c"), to = "d"))</code> . See Examples.
<code>condenseLevels</code>	a logical value. When set to the default value of <code>TRUE</code> and either <code>x</code> or <code>y</code> is a categorical variable, the function will drop all unused levels and rank the levels of the variable before calculating the correlation. When set to <code>FALSE</code> , the numeric levels of the variable remain the same as in the codebook. See Examples.

Details

Note that the `getData` arguments and `recode.sdf` may be useful. (See Examples.) The correlation methods are calculated as described in the documentation for the `wCorr` package—see `browseVignettes(package="wCorr")`.

Value

An `edsurvey.cor` that has print and summary methods.

The class includes the following elements:

<code>correlation</code>	numeric estimated correlation coefficient
<code>Zse</code>	standard error of the correlation ($V_{imp} + V_{jrr}$). In the case of Pearson, this is calculated in the linear atanh space and so is not a standard error in the usual sense.
<code>correlates</code>	a vector of length two showing the columns for which the correlation coefficient was calculated
<code>variables</code>	correlates that are discrete
<code>order</code>	a list that shows the order of each variable
<code>method</code>	the type of correlation estimated
<code>Vjrr</code>	the jackknife component of the variance estimate. For Pearson, in the atanh space.
<code>Vimp</code>	the imputation component of the variance estimate. For Pearson, in the atanh space.
<code>weight</code>	the weight variable used
<code>npv</code>	the number of plausible values used
<code>njk</code>	the number of the jackknife replicates used

Author(s)

Paul Bailey; relies heavily on the `wCorr` package, written by Ahmad Emad and Paul Bailey

See Also

`cor` and `weightedCorr`

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# for two categorical variables any of the following work
c1_pears <- cor.sdf(x="b017451", y="b003501", data=sdf, method="Pearson",
  weightVar="origwt")
c1_spear <- cor.sdf(x="b017451", y="b003501", data=sdf, method="Spearman",
  weightVar="origwt")
c1_polyc <- cor.sdf(x="b017451", y="b003501", data=sdf, method="Polychoric",
```

```

weightVar="origwt")

c1_pears
c1_spear
c1_polyc

# for categorical variables, users can either keep the original numeric levels of the variables
# or condense the levels (default)
# The following call condenses the levels of the variable 'c046501'
cor.sdf(x="c046501", y="c044006", data=sdf)

# The following call keeps the original levels of the variable 'c046501'
cor.sdf(x="c046501", y="c044006", data=sdf, condenseLevels = FALSE)

# these take awhile to calculate for large datasets, so limit to a subset
sdf_dnf <- subset(sdf, b003601 == 1)

# for a categorical variable and a scale score any of the following work
c2_pears <- cor.sdf(x="composite", y="b017451", data=sdf_dnf, method="Pearson",
weightVar="origwt")
c2_spear <- cor.sdf(x="composite", y="b017451", data=sdf_dnf, method="Spearman",
weightVar="origwt")
c2_polys <- cor.sdf(x="composite", y="b017451", data=sdf_dnf, method="Polyserial",
weightVar="origwt")

c2_pears
c2_spear
c2_polys

# recode two variables
cor.sdf(x="c046501", y="c044006", data=sdf, method="Spearman", weightVar="origwt",
recode=list(c046501=list(from="0%",to="None"),
c046501=list(from=c("1-5%", "6-10%", "11-25%", "26-50%",
"51-75%", "76-90%", "Over 90%"),
to="Between 0% and 100%"),
c044006=list(from=c("1-5%", "6-10%", "11-25%", "26-50%",
"51-75%", "76-90%", "Over 90%"),
to="Between 0% and 100%)))

# reorder two variables
cor.sdf(x="b017451", y="sdracem", data=sdf, method="Spearman", weightVar="origwt",
reorder=list(sdracem=c("White", "Hispanic", "Black", "Asian/Pacific Island",
"Amer Ind/Alaska Natv", "Other"),
b017451=c("Every day", "2 or 3 times a week", "About once a week",
"Once every few weeks", "Never or hardly ever")))

# recode two variables and reorder
cor.sdf(x="pared", y="b013801", data=subset(sdf, !pared %in% "I Don't Know"),
method="Spearman", weightVar = "origwt",
recode=list(pared=list(from="Some ed after H.S.", to="Graduated H.S."),
pared=list(from="Graduated college", to="Graduated H.S."),
b013801=list(from="0-10", to="Less than 100"),
b013801=list(from="11-25", to="Less than 100"),

```

```

      b013801=list(from="26-100", to="Less than 100"),
      reorder=list(b013801=c("Less than 100", ">100"))

```

```
## End(Not run)
```

```
dim.edsurvey.data.frame
```

Dimensions of an edsurvey.data.frame or an edsurvey.data.frame.list

Description

Returns the dimensions of an `edsurvey.data.frame` or an `edsurvey.data.frame.list`.

Usage

```
## S3 method for class 'edsurvey.data.frame'
dim(x)
```

Arguments

`x` an `edsurvey.data.frame` or an `edsurvey.data.frame.list`

Value

For an `edsurvey.data.frame`, returns a numeric vector of length two, with the first element being the number of rows and the second element being the number of columns.

For an `edsurvey.data.frame.list`, returns a list of length two, where the first list element is named `nrow` and is a numeric vector containing the number of rows for each element of the `edsurvey.data.frame.list`. The second element is named `ncol` and is the number of columns for each element. This is done so that the `nrow` and `ncol` functions return meaningful results, even if nonstandard.

Author(s)

Paul Bailey

```
DoFCorrection
```

Degrees of Freedom

Description

Calculates the degrees of freedom for a statistic (or of a contrast between two statistics) based on the jackknife and imputation variance estimates.

Usage

```
DoFCorrection(varEstA, varEstB = varEstA, varA, varB = varA,
  method = c("WS", "JR"))
```

Arguments

varEstA	the varEstInput object returned from certain functions, such as <code>lm.sdf</code> when <code>returnVarEstInputs= TRUE</code>). The variable varA must be on this data. See Examples.
varEstB	similar to the varEstA argument. If left blank, both are assumed to come from varEstA. When set, the degrees of freedom are for a contrast between varA and varB, and the varB values are taken from varEstB.
varA	a character that names the statistic in the varEstA argument for which the degrees of freedom calculation is required.
varB	a character that names the statistic in the varEstB argument for which a covariance is required. When varB is specified, returns the degrees of freedom for the contrast between varA and varB.
method	a character that is either WS for the Welch-Satterthwaite formula or JR for the Johnson-Rust correction to the Welch-Satterthwaite formula

Details

This calculation happens under the notion that statistics have little variance within strata, and some strata will contribute fewer than a full degree of freedom.

Note that these functions are not vectorized, so varA and varB must contain exactly one variable name.

The method used to compute the degrees of freedom is in the vignette titled **Statistics** section “Estimation of Degrees of Freedom.”

Value

numeric; the estimated degrees of freedom

Author(s)

Paul Bailey

References

Johnson, E. G., & Rust, K. F. (1992). Population inferences and variance estimation for NAEP data. *Journal of Educational Statistics*, 17, 175–190.

Examples

```
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))
lm1 <- lm.sdf(composite ~ dsex + b017451, sdf, returnVarEstInputs=TRUE)
summary(lm1)
# this output agrees with summary of lm1 coefficient for dsex
DoFCorrection(lm1$varEstInputs,
              varA="dsexFemale",
              method="JR")
# second example, a covariance term requires more work
# first, estimate the covariance between two regression coefficients
```

```

# note that the variable names are parallel to what they are called in lm1 output
covFEveryDay <- varEstToCov(lm1$varEstInputs,
                           varA="dsexFemale",
                           varB="b017451Every day",
                           jkSumMultiplier=EdSurvey::getAttributes(sdf, "jkSumMultiplier"))
# second, find the difference and the SE of the difference
se <- lm1$coefmat["dsexFemale","se"] + lm1$coefmat["b017451Every day","se"] +
      -2*covFEveryDay
# third, calculate the t-statistic
tv <- (coef(lm1)["dsexFemale"] - coef(lm1)["b017451Every day"])/se
# fourth, calculate the p-value, which requires the estimated degrees of freedom
dofFEveryDay <- DoFCorrection(lm1$varEstInputs,
                              varA="dsexFemale",
                              varB="b017451Every day",
                              method="JR")

# finally, the p-value
2*(1-pt(abs(tv), df=dofFEveryDay))

```

downloadCivEDICCS

Instructions for Downloading and Unzipping CivED or ICCS Files

Description

Provides instructions to download CivED or ICCS data to be processed in readCivEDICCS.

Usage

```
downloadCivEDICCS(years = c(1999, 2009))
```

Arguments

years an integer vector indicating the study year. Valid years are 1999 and 2009.

Author(s)

Tom Fink

See Also

[readCivEDICCS](#)

Examples

```

## Not run:
#view instructions to manually download study data
downloadCivEDICCS()

## End(Not run)

```

downloadECLS_K *Download and Unzip ECLS_K Files*

Description

Uses an Internet connection to download ECLS_K data. Data come from nces.ed.gov zip files. This function works for 1998 and 2011 data.

Usage

```
downloadECLS_K(root, years = c(1998, 2011), cache = FALSE,
               verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the ECLS_K data should be stored. Files are placed in a subdirectory named ECLS_K/[year].
years	an integer vector of the assessment years to download. Valid years are 1998 and 2011.
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Author(s)

Tom Fink

See Also

[readECLS_K1998](#) and [readECLS_K2011](#)

Examples

```
## Not run:
# root argument will vary by operating system conventions
downloadECLS_K(years=c(1998, 2011), root = "C:/")

# cache=TRUE will download then process the datafiles
downloadECLS_K(years=c(1998, 2011), root = "C:/", cache = TRUE)

#set verbose=FALSE for silent output
#if year not specified, download all years
downloadECLS_K(root="C:/", verbose = FALSE)

## End(Not run)
```

`downloadICILS`*Instructions for Downloading and Unzipping ICILS Files*

Description

Provides instructions to download ICILS data to be processed in readICILS.

Usage

```
downloadICILS(years = c(2013))
```

Arguments

`years` an integer vector indicating the study year. Valid year is 2013 only.

Author(s)

Tom Fink

See Also

[readICILS](#)

Examples

```
## Not run:  
#view instructions to manually download study data  
downloadICILS()  
  
## End(Not run)
```

`downloadPIAAC`*Download and Unzip PIAAC Files*

Description

Uses a connection to download PIAAC data to a computer. Data come from the OECD website.

Usage

```
downloadPIAAC(root, round = 1, cache = FALSE, verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the PIAAC data should be stored. Files are placed in a folder named PIAAC/Round [round number].
round	a numeric value indicating the assessment round to download. Valid round is 1 only (2012/2014).
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Author(s)

Trang Nguyen

Examples

```
## Not run:
# Download all available data for PIAAC round 1 to "C:/PIAAC/Round 1" folder
# root argument will vary by operating system conventions
downloadPIAAC(root="C:/")

## End(Not run)
```

downloadPIRLS

Download and Unzip PIRLS Files

Description

Uses an Internet connection to download PIRLS data. Data come from timssandpirls.bc.edu zip files. This function works for 2001, 2006, 2011, and 2016 data.

Usage

```
downloadPIRLS(root, years = c(2001, 2006, 2011, 2016), cache = FALSE,
  verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the PIRLS data should be stored. Files are placed in a subdirectory named PIRLS/[year].
years	an integer vector of the assessment years to download. Valid years are 2001, 2006, 2011, and 2016.
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Author(s)

Tom Fink

See Also[readPIRLS](#)**Examples**

```
## Not run:
# root argument will vary by operating system conventions
downloadPIRLS(year=c(2006, 2011), root = "C:/")

# cache=TRUE will download then process the datafiles
downloadPIRLS(year=2011, root = "C:/", cache = TRUE)

#set verbose=FALSE for silent output
#if year not specified, download all years
downloadPIRLS(root="C:/", verbose = FALSE)

## End(Not run)
```

downloadPISA

Download and Unzips PISA Files

Description

Uses a connection to download PISA data to a computer. Data come from the OECD website.

Usage

```
downloadPISA(root, years = c(2000, 2003, 2006, 2009, 2012),
  database = c("INT", "CBA", "FIN"), cache = FALSE, verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the PISA data should be stored. Files are placed in a folder named PISA/[year].
years	an integer vector of the assessment years to download. Valid years are 2000, 2003, 2006, 2009, and 2012.
database	a character vector to indicate which database to download from. For 2012, three databases are available (INT = International, CBA = Computer-Based Assessment, and FIN = Financial Literacy). Defaults to INT.
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Details

The function uses `download.file` to download files from provided URLs. Some machines might require a different user agent in HTTP(S) requests. If the downloading gives an error or behaves unexpectedly (for example, a zip file cannot be unzipped or a data file is significantly smaller than expected), users can toggle `HTTPUserAgent` options to find one that works for their machines. One common alternative option is

```
options(HTTPUserAgent="Mozilla/5.0 (Windows NT 6.1; WOW64; rv:53.0) Gecko/20100101 Firefox/53.0").
```

Author(s)

Trang Nguyen

See Also

[readPISA](#), `download.file`, `options`

Examples

```
## Not run:
# Download PISA 2012 data (for all three databases)
downloadPISA(years = 2012, database = c("INT", "CBA", "FIN"), root="C:/")

# Download PISA 2009 and 2012 data (International Database only)
# to C:/PISA/2009 and C:/PISA/2012 folder respectively
downloadPISA(years = c(2009,2012), root="C:/")

## End(Not run)
```

downloadTALIS

Instructions for Downloading TALIS Files

Description

Provides instructions to download TALIS data to be processed in [readTALIS](#).

Usage

```
downloadTALIS(years)
```

Arguments

`years` a numeric value indicating the assessment year. Available years are 2008 and 2013.

Author(s)

Trang Nguyen

See Also[readTALIS](#)**Examples**

```
## Not run:
# Print out downloading instructions for TALIS 2008 database
downloadTALIS(2008)

## End(Not run)
```

downloadTIMSS	<i>Download and Unzip TIMSS Files</i>
---------------	---------------------------------------

Description

Uses an Internet connection to download TIMSS data. Data come from timssandpirls.bc.edu zip files. This function works for 2003, 2007, 2011, and 2015 data.

Usage

```
downloadTIMSS(root, years = c(2003, 2007, 2011, 2015), cache = FALSE,
  verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the TIMSS data should be stored. Files are placed in a subdirectory named TIMSS/[year].
years	an integer vector of the assessment years to download. Valid years are 2003, 2007, 2011, and 2015.
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Author(s)

Tom Fink

See Also[readTIMSS](#)

Examples

```
## Not run:
# root argument will vary by operating system conventions
downloadTIMSS(year=c(2015, 2011), root = "C:/")

# cache=TRUE will download then process the datafiles
downloadTIMSS(year=2015, root = "C:/", cache = TRUE)

#set verbose=FALSE for silent output
#if year not specified, download all years
downloadTIMSS(root="C:/", verbose = FALSE)

## End(Not run)
```

downloadTIMSSAdv

Download and Unzip TIMSS Advanced Files

Description

Uses an Internet connection to download TIMSS Advanced data. Data comes from timssand-pirls.bc.edu zip files. This function works for 1995, 2008, and 2015 data.

Usage

```
downloadTIMSSAdv(root, years = c(1995, 2008, 2015), cache = FALSE,
  verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the TIMSS Advanced data should be stored. Files are placed in a subdirectory named TIMSSAdv/[year].
years	an integer vector of the assessment years to download. Valid years are 1995, 2008, and 2015.
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Author(s)

Tom Fink

See Also

[readTIMSSAdv](#)

Examples

```
## Not run:
#root argument will vary by operating system conventions
downloadTIMSSAdv(year=c(2008, 2015), root = "C:/")

#cache=TRUE will download then process the datafiles
downloadTIMSSAdv(year=2015, root = "C:/", cache = TRUE)

#set verbose=FALSE for silent output
#if year not specified, download all years
downloadTIMSSAdv(root="C:/", verbose = FALSE)

## End(Not run)
```

download_ePIRLS

Download and Unzip ePIRLS Files

Description

Uses an Internet connection to download ePIRLS data. Data come from timssandpirls.bc.edu zip files. This function works for 2016 data.

Usage

```
download_ePIRLS(root, years = c(2016), cache = FALSE, verbose = TRUE)
```

Arguments

root	a character string indicating the directory where the ePIRLS data should be stored. Files are placed in a subdirectory named ePIRLS/[year].
years	an integer vector of the assessment years to download. Valid year is 2016 only.
cache	a logical value set to process and cache the text (.txt) version of files. This takes a very long time but saves time for future uses of the data. Default value is FALSE.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Author(s)

Tom Fink

See Also

[read_ePIRLS](#)

Examples

```
## Not run:
# root argument will vary by operating system conventions
download_ePIRLS(years=2016, root = "C:/")

# cache=TRUE will download then process the datafiles
download_ePIRLS(years=2016, root = "C:/", cache = TRUE)

#set verbose=FALSE for silent output
#if year not specified, download all years
download_ePIRLS(root="C:/", verbose = FALSE)

## End(Not run)
```

edsurvey.data.frame *EdSurvey Class Constructors*

Description

Two new classes in EdSurvey are described in this section: the `edsurvey.data.frame` and `light.edsurvey.data.frame`. The `edsurvey.data.frame` class stores metadata about survey data and data is stored on the disk (via the LaF package), allowing GB of data to be used easily on a machine otherwise inappropriate for manipulating large datasets. The `light.edsurvey.data.frame` is typically generated by the `getData` function and stores the data in a `data.frame`. Both of the classes use attributes to manage metadata and allow for correct statistics to be used in calculating results; the `getAttributes` acts as an accessor for these attributes, while `setAttributes` acts as a mutator for the attributes. As a convenience, `edsurvey.data.frame` implements the `$` function to extract a variable.

Usage

```
edsurvey.data.frame(userConditions, defaultConditions, data, dataSch,
  dataTch, dataListMeta, weights, pvvars, subject, year, assessmentCode,
  dataType, gradeLevel, achievementLevels, omittedLevels, fileFormat,
  fileFormatSchool, fileFormatTeacher, survey, country, psuVar, stratumVar,
  jkSumMultiplier, recodes = NULL, validateFactorLabels = FALSE,
  forceLower = TRUE)

## S3 method for class 'edsurvey.data.frame'
x$i

getAttributes(data, attribute = NULL)

setAttributes(data, attribute, value)
```

Arguments

`userConditions` a list of user conditions that includes subsetting or recoding conditions

defaultConditions	a list of default conditions that are often set for each survey
data	in the edsurvey.data.frame constructor, this is an LaF object that connects to the main data, often at the student level. For getAttributes and setAttributes, this argument is an edsurvey.data.frame or light.edsurvey.data.frame.
dataSch	an LaF object that connects to the school-level data (optional)
dataTch	an LaF object that connects to the teacher-level data (optional)
dataListMeta	a list that stores variables that can be used to link school-level and teacher-level data to the main data. See Details.
weights	a list that stores information regarding weight variables. See Details.
pvvars	a list that stores information regarding plausible values. See Details.
subject	a character that indicates subject domain of the given data
year	a character or numeric that indicates year of the given data
assessmentCode	a character that indicates the code of the assessment. Can be “National” or “International”.
dataType	a character that indicates the unit level of the main data. Examples include dQuoteStudent, “teacher”, “school”, “Adult Data”.
gradeLevel	a character that indicates grade level of the given data
achievementLevels	a list of achievement level categories and cutpoints
omittedLevels	a list of default omitted levels for the given data
fileFormat	a data.frame that stores codebook information for the main data. See Details.
fileFormatSchool	a data.frame that stores codebook information for the school-level data (if exists). See Details.
fileFormatTeacher	a data.frame that stores codebook information for the teacher-level data (if exists). See Details.
survey	a character that indicates the name of the survey
country	a character that indicates the country of the given data
psuVar	a character that indicates the PSU sampling unit variable. Ignored when weights have psuVar defined.
stratumVar	a character indicates the stratum variable. Ignored with weights have stratumVar defined.
jkSumMultiplier	a numeric value of the jackknife coefficient (used in calculating the jackknife replication estimation)
recodes	a list of variable recodes of the given data
validateFactorLabels	a Boolean that indicates whether the getData function needs to validate factor variables
forceLower	a Boolean; when set to TRUE, will automatically lowercase variable names

x	an edsurvey.data.frame
i	a character, the column name to extract
attribute	a character, name of an attribute to get or set
value	new value of the given attribute

Details

The `dataListMeta` argument is a list with an element `student` that is also a list. Each element of the `student` list is another dataset name (`teacher` or `school`) that indicates the variables used to link the student file to those files. The merge variables are shown with a caret character (“^”) between them. The first variable is the name of the merge variable on the student file, and the second variable is the name of the merge variable on the school file. When multiple variables are used to merge, a semicolon can separate pairs of variables; e.g., `student=list(school="varA^varY;varB^varZ")` would indicate that the student file can be merged to the school file using the `varA` and `varB` variables from the student file to merge it to `varY` and `varZ`, respectively, on the school file.

The `weight` list has an element named after each weight variable name that is a list with elements `jkbase` and `jksuffixes`. The `jkbase` variable is a single character indicating the jackknife replicate weight base name, while `jksuffixes` is a vector with one element for each jackknife replicate weight. When the two are pasted together, they should form the complete set of jackknife replicate weights. The `weights` argument can also have an attribute that is the default weight. If the primary sampling unit and stratum variables change by weight, they can also be defined on the weight list as `psuVar` and `stratumVar`. When this option is used, it overrides the `psuVar` and `stratumVar` on the `edsurvey.data.frame`, which can be left blank. A weight must define only one of `psuVar` and `stratumVar`.

The `pvvars` list has an element for each subject or subscale score that has plausible values. Each element is a list with a `varnames` element that indicates the column names of the plausible values and an `achievementLevel` argument that is a named vector of the achievement level cut points.

The `fileFormat` arguments are data frames that have the following columns:

variableName name of the variable. Changed to lower case by the constructor if `forceLower=TRUE`.

Start start column of the data

End end column of the data

Width number of characters wide the data is

Decimal power of 10 that the data should be divided by

Labels brief description of the variable

labelValues an caret (“^”) delimited list of label value pairs, each of which is equal delimited (“=”) as `code=value`. For example, the string “1=true^2=false^3=invalid” would result in values of 1 being labeled “true”, values 2 being labeled “false”, and values of 3 being labeled “invalid”.

dataType one of “character”, “numeric”, or “integer”

Weights Boolean set to `TRUE` to indicate that the column is a full sample (not replicate) weight column

Value

An object of class `edsurvey.data.frame` with the following elements:

Elements that store data connections and data codebooks

data an LaF object containing a connection to the student dataset on disk

dataSch an LaF object containing a connection to the school dataset on disk if exists. If not, will be NULL.

dataTch an LaF object containing a connection to the teacher dataset on disk if exists. If not, will be NULL.

fileFormat a `data.frame` containing the format of the file in the `data` parameter. See Details.

fileFormatSchool a `data.frame` containing the format of the file in the `dataSch` parameter. See Details.

fileFormatTeacher a `data.frame` containing the format of the file in the `dataTch` parameter. See Details.

Elements that store sample design and default subsetting information of the given survey data

userConditions a list containing all user conditions, set using the `subset.edsurvey.data.frame` method

defaultConditions the default subsample conditions

weights a list containing the weights. See Details.

stratumVar a character that indicates the default strata identification variable name in the data. Often used in Taylor series estimation.

psuVar a character that indicates the default PSU (sampling unit) identification variable name in the data. Often used in Taylor series estimation.

pvvars a list containing the plausible values. See Details.

achievementLevels default achievement cutoff scores and names. See Details.

omittedLevels the levels of the factor variables that will be omitted from the `edsurvey.data.frame`

Elements that store descriptive information of the survey

survey the type of survey data

subject the subject of the data

year the year of assessment

assessmentCode the assessment code

dataType the type of data (e.g., “student” or “school”)

gradeLevel the grade of the dataset contained in the `edsurvey.data.frame`

EdSurvey Classes

`edsurvey.data.frame` is an object that stores connection to data on the disk along with important survey sample design information.

`edsurvey.data.frame.list` is a list of `edsurvey.data.frame` objects. It is often used in trend or cross-regional analysis in the `gap` function. See `edsurvey.data.frame.list` for more information on how to create an `edsurvey.data.frame.list`. Users can also refer to the vignette titled [Using EdSurvey for Trend Analysis](#) for examples.

Besides `edsurvey.data.frame` class, EdSurvey package also implements `light.edsurvey.data.frame` class, which can be used by both EdSurvey and non-EdSurvey functions. More particularly, `light.edsurvey.data.frame` is a `data.frame` that also has basic survey and sample design information (i.e., plausible values and weights), which will be used for variance estimation in analytical functions. Because it is also a base R `data.frame`, users can also apply base R functions for data manipulation. vignette titled [getData](#) for more examples.

Many functions will remove attributes from a data frame, such as a `light.edsurvey.data.frame`, and the `rebindAttributes` function can add them back.

Users can get a `light.edsurvey.data.frame` object by using `getData` method with `addAttributes=TRUE`.

Basic Methods for EdSurvey Classes

Extracting a column from an `edsurvey.data.frame`

Users can extract a column from an `edsurvey.data.frame` object using `$` or `[]` like a normal data frame.

Extracting and updating attributes of an object of class `edsurvey.data.frame` or `light.edsurvey.data.frame`

Users can use `getAttributes` method to extract any of the attributes of an `edsurvey.data.frame` or `light.edsurvey.data.frame`. Note that a `light.edsurvey.data.frame` will not have attributes related to data connection because data has already been read in memory.

If users want to update an attribute (i.e., `omittedLevels`), users can use the `setAttributes` method.

Author(s)

Tom Fink, Trang Nguyen, and Paul Bailey

See Also

[rebindAttributes](#)

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# Run a base R function on a column of edsurvey.data.frame
table(sdf$dsex)

# Extract default omitted levels of NAEP primer data
getAttributes(sdf, "omittedLevels") # [1] "Multiple" NA "Omitted"
```

```
# Update default omitted levels of NAEP primer data
sdf <- setAttributes(sdf, "omittedLevels", c("Multiple", "Omitted", NA, "(Missing)"))
getAttributes(sdf, "omittedLevels") #[1] "Multiple" "Omitted" NA "(Missing)"
```

edsurvey.data.frame.list

EdSurvey Dataset Vectorization

Description

The `edsurvey.data.frame.list` function creates an `edsurvey.data.frame.list` object from a series of `edsurvey.data.frame` objects. `append.edsurvey.data.frame.list` creates an `edsurvey.data.frame.list` from two `edsurvey.data.frame` or `edsurvey.data.frame.list` objects.

An `edsurvey.data.frame.list` is useful for looking at data, for example, across time or graphically, and reduces repetition in function calls. The user may specify a variable that varies across the `edsurvey.data.frame` objects that is then included in further output.

Usage

```
edsurvey.data.frame.list(datalist, cov = NULL, labels = NULL)
```

```
append.edsurvey.data.frame.list(sdfA, sdfB, labelsA = NULL,
  labelsB = NULL)
```

Arguments

<code>datalist</code>	a list of <code>edsurvey.data.frame</code> s to be combined
<code>cov</code>	a character vector that indicates what varies across the <code>edsurvey.data.frame</code> objects. See Examples. Guessed if not supplied. For example, if several <code>edsurvey.data.frame</code> s for several different countries are supplied, then <code>cov</code> would be set to the country.
<code>labels</code>	a character vector that specifies labels. Must be the same length as <code>datalist</code> . Not needed if <code>cov</code> exists or can be guessed. See Examples.
<code>sdfA</code>	an <code>edsurvey.data.frame</code> or <code>edsurvey.data.frame.list</code> to be combined
<code>sdfB</code>	an <code>edsurvey.data.frame</code> or <code>edsurvey.data.frame.list</code> to be combined
<code>labelsA</code>	a character vector that specifies labels for <code>sdfA</code> when creating the new <code>edsurvey.data.frame.list</code> . <code>labelsA</code> would be ignored if <code>sdfA</code> is an <code>edsurvey.data.frame.list</code> with labels supplied.
<code>labelsB</code>	a character vector that specifies labels for <code>sdfB</code> when creating the new <code>edsurvey.data.frame.list</code> . <code>labelsB</code> would be ignored if <code>sdfB</code> is an <code>edsurvey.data.frame.list</code> with labels supplied.


```

                                "B locations",
                                "C locations",
                                "D locations"))

# this shows how these datasets will be described
sdf1$covs
## Not run:
# get the gaps between Male and Female for each data set
gap1 <- gap("composite", sdf1, dsex=="Male", dsex=="Female")
gap1

## End(Not run)

# make combine sdfA and sdfB
sdf11a <- edsurvey.data.frame.list(list(sdfA, sdfB),
                                   labels=c("A locations",
                                             "B locations"))

# combine sdfC and sdfD
sdf11b <- edsurvey.data.frame.list(list(sdfC, sdfD),
                                   labels=c("C locations",
                                             "D locations"))

# append to make sdf3 the same as sdf1
sdf13 <- append.edsurvey.data.frame.list(sdf11a, sdf11b)
identical(sdf1, sdf13) #TRUE

# append to make sdf4 the same as sdf1
sdf14 <- append.edsurvey.data.frame.list(
  append.edsurvey.data.frame.list(sdf11a, sdfC, labelsB = "C locations"),
  sdfD,
  labelsB = "D locations")
identical(sdf1, sdf14) #TRUE

```

edsurveyTable

EdSurvey Tables With Conditional Means

Description

Returns a summary table (as a data.frame) that shows the number of students, the percentage of students, and the mean value of the outcome (or left-hand side) variable by the predictor (or right-hand side) variable(s).

Usage

```

edsurveyTable(formula, data, weightVar = NULL, jrrIMax = 1,
              pctAggregationLevel = NULL, returnMeans = TRUE, returnSepct = TRUE,
              varMethod = c("jackknife", "Taylor"), drop = FALSE,

```

```
omittedLevels = TRUE, defaultConditions = TRUE, recode = NULL,
returnVarEstInputs = FALSE)
```

Arguments

formula	object of class formula, potentially with a subject scale or subscale on the left-hand side and variables to tabulate on the right-hand side. When the left-hand side of the formula is omitted and returnMeans is TRUE, then the default subject scale or subscale is used. You can find the default composite scale and all subscales using the function showPlausibleValues . Note that the order of the right-hand side variables affects the output.
data	object of class <code>edsurvey.data.frame</code> . See readNAEP for how to generate an <code>edsurvey.data.frame</code> .
weightVar	character string indicating the weight variable to use. Note that only the name of the weight variable needs to be included here, and any replicate weights will be automatically included. When this argument is NULL, the function uses the default. Use showWeights to find the default.
jrrIMax	integer indicating the maximum number of plausible values to include when calculating the variance term V_{jrr} (see the Details section of lm.sdf to see the definition of V_{jrr}). The default is Inf and results in all available plausible values being used in generating V_{jrr} . Setting this to 1 will make code execution faster but less accurate.
pctAggregationLevel	the percentage variable sums up to 100 for the first <code>pctAggregationLevel</code> columns. So when set to 0, the PCT column adds up to one across the entire sample. When set to 1, the PCT column adds up to one within each level of the first variable on the right-hand side of the formula; when set to 2, then the percentage adds up to 100 within the interaction of the first and second variable, and so on. Default is NULL, which will result in the lowest feasible aggregation level. See Examples section.
returnMeans	a logical value; set to TRUE (the default) to get the MEAN and SE(MEAN) columns in the returned table described in the Value section.
returnSepct	set to TRUE (the default) to get the SEPCT column in the returned table described in the Value section.
varMethod	a character set to “jackknife” or “Taylor” that indicates the variance estimation method to be used.
drop	a logical value. When set to the default value of FALSE, when a single column is returned, it is still represented as a <code>data.frame</code> and is not converted to a vector.
omittedLevels	a logical value. When set to the default value of TRUE, drops those levels of all factor variables that are specified in an <code>edsurvey.data.frame</code> . Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels.
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in an <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.

recode	a list of lists to recode variables. Defaults to NULL. Can be set as <code>recode = list(var1 = list(from = c("a", "b", "c"), to = "c"))</code> . See Examples.
returnVarEstInputs	a logical value set to TRUE to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for the computation of covariances between estimates.

Details

This method can be used to generate a simple one-way, two-way, or n -way table with unweighted and weighted n values and percentages. It also can calculate the average of the subject scale or subscale for students at each level of the cross-tabulation table.

A detailed description of all statistics is given in the vignette titled [Statistics](#).

Value

A table with the following columns:

RHS levels	one column for each right-hand side variable. Each row regards students who are at the levels shown in that row.
N	count of the number of students in the survey in the RHS levels
WTD_N	the weighted N count of students in the survey in RHS levels
PCT	the percentage of students at the aggregation level specified by <code>pctAggregationLevel</code> (see Arguments). See the vignette titled Statistics in the section “Estimation of Weighted Percentages” and its first subsection “Estimation of Weighted Percentages When Plausible Values Are Not Present.”
SE(PCT)	the standard error of the percentage, accounting for the survey sampling methodology. When <code>varMethod</code> is <code>jackknife</code> , the calculation of this column is described in the vignette titled Statistics in the section “Estimation of the Standard Error of Weighted Percentages When Plausible Values Are Not Present, Using the Jackknife Method.” When <code>varMethod</code> is set to <code>Taylor</code> , the calculation of this column is described in “Estimation of the Standard Error of Weighted Percentages When Plausible Values Are Not Present, Using the Taylor Series Method.”
MEAN	the mean assessment score for units in the RHS levels, calculated according to the vignette titled Statistics in the section “Estimation of Weighted Means When Plausible Values Are Present.”
SE(MEAN)	the standard error of the MEAN column (the mean assessment score for units in the RHS levels), calculated according to the vignette titled Statistics in the sections “Estimation of Standard Errors of Weighted Means When Plausible Values Are Present, Using the Jackknife Method” or “Estimation of Standard Errors of Weighted Means When Plausible Values Are Present, Using the Taylor Series Method,” depending on the value of <code>varMethod</code> .

When `returnVarEstInputs` is TRUE, two additional elements are returned. These are `meanVarEstInputs` and `pctVarEstInputs` and regard the MEAN and PCT columns, respectively. These two objects can be used for calculating covariances with [varEstToCov](#).

Author(s)

Paul Bailey and Ahmad Emad

References

Binder, D. A. (1983). On the variances of asymptotically normal estimators from complex surveys. *International Statistical Review*, 51(3), 279–292.

Rubin, D. B. (1987). *Multiple imputation for nonresponse in surveys*. New York, NY: Wiley.

Examples

```
## Not run:
# read in the example data (generated, not real student data)

sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# create a table that shows only the breakdown of dsex
edsurveyTable(composite ~ dsex, data=sdf, returnMeans=FALSE, returnSepct=FALSE)

# create a table with composite scores by dsex
edsurveyTable(composite ~ dsex, data=sdf)

# add a second variable
edsurveyTable(composite ~ dsex + b017451, data=sdf)

# add a second variable, do not omit any levels
edsurveyTable(composite ~ dsex + b017451 + b003501, data=sdf, omittedLevels=FALSE)

# add a second variable, do not omit any levels, change aggregation level
edsurveyTable(composite ~ dsex + b017451 + b003501, data=sdf, omittedLevels=FALSE,
              pctAggregationLevel=0)

edsurveyTable(composite ~ dsex + b017451 + b003501, data=sdf, omittedLevels=FALSE,
              pctAggregationLevel=1)

edsurveyTable(composite ~ dsex + b017451 + b003501, data=sdf, omittedLevels=FALSE,
              pctAggregationLevel=2)

# variance estimation using the Taylor series
edsurveyTable(composite ~ dsex + b017451 + b003501, data=sdf, varMethod="Taylor")

## End(Not run)
```

edsurveyTable2pdf

PDF File From an edsurveyTable

Description

Produces the LaTeX code and compiles as a PDF file from the edsurveyTable results.

Usage

```
edsurveyTable2pdf(data, formula, caption = NULL, filename = "",
  toCSV = "", returnMeans = TRUE, estDigits = 2, seDigits = 3)
```

Arguments

<code>data</code>	an <code>edsurveyTable</code> . See Value in edsurveyTable .
<code>formula</code>	a formula of the form <code>LHS ~ RHS</code> to cast the <code>edsurveyTable</code> results from long format to wide format. This formula takes the form <code>LHS ~ RHS</code> (e.g., <code>var1 + var2 ~ var3</code>). The order of the entries in the formula is essential.
<code>caption</code>	character vector of length one or two containing the table's caption or title. If length is two, the second item is the "short caption" used when LaTeX generates a <code>List of Tables</code> . Set to <code>NULL</code> to suppress the caption. Default value is <code>NULL</code> .
<code>filename</code>	a character string containing filenames and paths. By default (<code>filename = ""</code>), table will be saved in the working directory (<code>getwd()</code>). Use <code>filename = "CONSOLE"</code> to print LaTeX code in R console without generating a PDF file.
<code>toCSV</code>	a character string containing filenames and paths of <code>.csv</code> table output. <code>""</code> indicates no <code>.csv</code> output. <code>toCSV</code> is independent to <code>filename</code> , so both a <code>csv</code> file and PDF file would be generated if both <code>filename</code> and <code>toCSV</code> were specified.
<code>returnMeans</code>	a logical value set to <code>TRUE</code> (the default) to generate a PDF with the MEAN and SE(MEAN). It is set to <code>FALSE</code> to generate a PDF with the PCT and SE(PCT). See Value in edsurveyTable .
<code>estDigits</code>	an integer indicating the number of decimal places to be used for estimates. Negative values are allowed. See Details.
<code>seDigits</code>	an integer indicating the number of decimal places to be used for standard errors. Negative values are allowed. See Details.

Details

Rounding to a negative number of digits means rounding to a power of 10, so, for example, `estDigits = -2` rounds estimates to the nearest hundred.

Note

For more details, see the vignette titled [Producing LaTeX Tables From edsurveyTable Results With edsurveyTable2pdf](#).

Author(s)

Huade Huo

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))
```

```

# create a table with composite scores by dsex and b017451
est1 <- edsurveyTable(composite ~ dsex + b017451, sdf)

# create a table with csv output
edsurveyTable2pdf(data = est1,
                  formula = b017451~dsex,
                  toCSV = "C:/example table.csv",
                  filename = "C:/example table.pdf",
                  returnMeans = FALSE)

# create a pdf file using the default subject scale or subscale
# and keep two digits for estimates and three digits for SE after decimal point
edsurveyTable2pdf(est1, b017451~dsex,
                  returnMeans = TRUE, estDigits = 2, seDigits = 3)

# create a pdf file using the percentage of students at the
# aggregation level specified by \code{pctAggregationLevel}
# Output will be saved as "C:/example table.pdf"
edsurveyTable2pdf(est1,
                  b017451~dsex,
                  "C:/example table.pdf",
                  returnMeans = FALSE)

## End(Not run)

```

gap

Gap Analysis

Description

Compares the average levels of a variable between two groups that potentially share members.

Usage

```

gap(variable, data, groupA = "default", groupB = "default",
     percentiles = NULL, achievementLevel = NULL,
     achievementDiscrete = FALSE, targetLevel = NULL, weightVar = NULL,
     jrrIMax = 1, varMethod = c("jackknife", "Taylor"),
     omittedLevels = TRUE, defaultConditions = TRUE, recode = NULL,
     referenceDataIndex = 1, returnVarEstInputs = FALSE,
     returnSimpleDoF = FALSE, returnSimpleN = FALSE,
     returnNumberOfPSU = FALSE)

```

Arguments

variable	a character indicating the variable to be compared, potentially with a subject scale or subscale
data	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>

groupA	an expression or character expression that defines a condition for subset. This subset will be compared to groupB. If not specified, it will define a whole sample as in data.
groupB	an expression or character expression that defines a condition for subset. This subset will be compared to groupA. If not specified, it will define a whole sample as in data. If set to NULL, estimates for the second group will be dropped.
percentiles	a numeric vector. The gap function calculates the mean when this argument is omitted or set to NULL. Otherwise, the gap at the percentile given is calculated.
achievementLevel	the achievement level(s) at which percentages should be calculated
achievementDiscrete	a logical indicating if the achievement level specified in the achievementLevel argument should be interpreted as discrete so that just the percentage in that particular achievement level will be included. Defaults to FALSE so that the percentage at or above that achievement level will be included in the percentage.
targetLevel	a character string. When specified, calculates the gap in the percentage of students at targetLevel in variable. This is useful for comparing the gap in the percentage of students at a survey response level.
weightVar	a character indicating the weight variable to use. See Details.
jrrIMax	a numeric value; when using the jackknife variance estimation method, the V_{jrr} term (see Details) can be estimated with any positive number of plausible values and is estimated on the lower of the number of available plausible values and jrrIMax. When jrrIMax is set to Inf, all plausible values will be used. Higher values of jrrIMax lead to longer computing times and more accurate variance estimates.
varMethod	a character set to jackknife or Taylor that indicates the variance estimation method to be used
omittedLevels	a logical value. When set to the default value of TRUE, drops those levels of all factor variables. Use print on an edsurvey.data.frame to see the omitted levels.
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in edsurvey.data.frame to subset the data. Use print on an edsurvey.data.frame to see the default conditions.
recode	a list of lists to recode variables. Defaults to NULL. Can be set as recode = list(var1 = list(from = c("a", "b", "c"), to = "d")). See Examples.
referenceDataIndex	a numeric used only when data is an edsurvey.data.frame.list, indicating which dataset is the reference dataset that other datasets are compared with. Defaults to one.
returnVarEstInputs	a logical value; set to TRUE to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for the computation of covariances between estimates.

<code>returnSimpleDoF</code>	a logical value set to TRUE to return the degrees of freedom for some statistics (see Value section) that do not have a <i>t</i> -test; useful primarily for further computation
<code>returnSimpleN</code>	a logical value set to TRUE to add the count (<i>n</i> -size) of observations included in groups A and B in the percentage object
<code>returnNumberOfPSU</code>	a logical value set to TRUE to return the number of primary sampling units (PSU) used in calculation.

Details

This function calculates the gap between groupA and groupB (which may be omitted to indicate the full sample). The gap is calculated for one of four statistics:

the gap in means The mean score gap (in the score variable) identified in the `variable` argument. This is the default. The means and their standard errors are calculated using the methods described in the [lm.sdf](#) function documentation.

the gap in percentiles The gap between respondents at the percentiles specified in the `percentiles` argument. This is returned when the `percentiles` argument is defined. The mean and standard error are computed as described in the [percentile](#) function documentation.

the gap in achievement levels The gap in the percentage of students at (when `achievementDiscrete` is TRUE) or at or above (when `achievementDiscrete` is FALSE) a particular achievement level. This is used when the `achievementLevel` argument is defined. The mean and standard error are calculated as described in the [achievementLevels](#) function documentation.

the gap in a survey response The gap in the percentage of respondents responding at `targetLevel` to `variable`. This is used when `targetLevel` is defined. The mean and standard deviation are calculated as described in the [edsurveyTable](#) function documentation.

Value

The return type depends on if the class of the data argument is an `edsurvey.data.frame` or an `edsurvey.data.frame.list`. Both include the call (called `call`), a list called `labels`, an object named `percentage` that shows the percentage in groupA and groupB, and an object that shows the gap called `results`.

The labels includes the following elements:

definition the definitions of the groups

nFullData the *n*-size for the full dataset (before applying the definition)

nUsed the *n*-size for the data after the group is subsetted and other restrictions (such as omitted values) are applied

nPSU the number of PSUs used in calculation—only returned when `returnNumberOfPSU = TRUE`

The percentages are computed according to the vignette titled [Statistics](#) in the section “Estimation of Weighted Percentages When Plausible Values Are Not Present.” The standard errors are calculated according to “Estimation of the Standard Error of Weighted Percentages When Plausible Values

Are Not Present, Using the Jackknife Method.” Standard errors of differences are calculated as the square root of the typical variance formula

$$Var(A - B) = Var(A) + Var(B) - 2Cov(A, B)$$

where the covariance term is calculated as described in the vignette titled **Statistics** in the section “Estimation of Covariances.” These degrees of freedom are available only with the jackknife variance estimation. The degrees of freedom used for hypothesis testing are always set to the number of jackknife replicates in the data.

the data argument is an edsurvey.data.frame: When the data argument is an `edsurvey.data.frame`, `gap` returns an S3 object of class `gap`.

The percentage object is a numeric vector with the following elements:

pctA the percentage of respondents in groupA compared with the whole sample in data

pctAse the standard error on the percentage of respondents in groupA

dofA degrees of freedom appropriate for a *t*-test involving pctA. This value is returned only if `returnSimpleDoF=TRUE`.

pctB the percentage of respondents in groupB.

pctBse the standard error on the percentage of respondents in groupB

dofB degrees of freedom appropriate for a *t*-test involving pctA. This value is returned only if `returnSimpleDoF=TRUE`.

diffAB the value of pctA minus pctB

covAB the covariance of pctA and pctB; used in calculating `diffABse`.

diffABse the standard error of pctA minus pctB

diffABpValue the *p*-value associated with the *t*-test used for the hypothesis test that `diffAB` is zero.

dofAB degrees of freedom used in calculating `diffABpValue`

The results object is a numeric data frame with the following elements:

estimateA the mean estimate of groupA (or the percentage estimate if `achievementLevel` or `targetLevel` is specified)

estimateAse the standard error of `estimateA`

dofA degrees of freedom appropriate for a *t*-test involving meanA. This value is returned only if `returnSimpleDoF=TRUE`.

estimateB the mean estimate of groupB (or the percentage estimate if `achievementLevel` or `targetLevel` is specified)

estimateBse the standard error of `estimateB`

dofB degrees of freedom appropriate for a *t*-test involving meanB. This value is returned only if `returnSimpleDoF=TRUE`.

diffAB the value of `estimateA` minus `estimateB`

covAB the covariance of `estimateA` and `estimateB`. Used in calculating `diffABse`.

diffABse the standard error of `diffAB`

diffABpValue the *p*-value associated with the *t*-test used for the hypothesis test that `diffAB` is zero.

dofAB degrees of freedom used for the *t*-test on `diffAB`

If the gap was in achievement levels or percentiles and more than one percentile or achievement level is requested, then an additional column labeled `percentiles` or `achievementLevel` is included in the `results` object.

When `results` has a single row and when `returnVarEstInputs` is `TRUE`, the additional elements `varEstInputs` and `pctVarEstInputs` also are returned. These can be used for calculating covariances with [varEstToCov](#).

the data argument is an `edsurvey.data.frame.list`: When the data argument is an `edsurvey.data.frame.list`, `gap` returns an S3 object of class `gapList`.

The `results` object in the `edsurveyResultList` is a `data.frame`. Each row regards a particular dataset from the `edsurvey.data.frame`, and a reference dataset is dictated by the `referenceDataIndex` argument.

The percentage object is a `data.frame` with the following elements:

covs a data frame with a column for each column in the covs. See previous section for more details.

... all elements in the percentage object in the previous section

diffAA the difference in `pctA` between the reference data and this dataset. Set to NA for the reference dataset.

covAA the covariance of `pctA` in the reference data and `pctA` on this row. Used in calculating `diffAAse`.

diffAAse the standard error for `diffAA`.

diffAApValue the p -value associated with the t -test used for the hypothesis test that `diffAA` is zero

diffBB the difference in `pctB` between the reference data and this dataset. Set to NA for the reference dataset.

covBB the covariance of `pctB` in the reference data and `pctB` on this row. Used in calculating `diffBBse`.

diffBBse the standard error for `diffBB`

diffBBpValue the p -value associated with the t -test used for the hypothesis test that `diffBB` is zero

diffABAB the value of `diffAB` in the reference dataset minus the value of `diffAB` in this dataset. Set to NA for the reference dataset.

covABAB the covariance of `diffAB` in the reference data and `diffAB` on this row. Used in calculating `diffABABse`.

diffABABse the standard error for `diffABAB`

diffABABpValue the p -value associated with the t -test used for the hypothesis test that `diffABAB` is zero

The `results` object is a `data.frame` with the following elements:

... all elements in the `results` object in the previous section

diffAA the value of `groupA` in the reference dataset minus the value in this dataset. Set to NA for the reference dataset.

covAA the covariance of `meanA` in the reference data and `meanA` on this row. Used in calculating `diffAAse`.

diffAAse the standard error for `diffAA`.

diffAApValue the p -value associated with the t -test used for the hypothesis test that `diffAA` is zero

diffBB the value of `groupB` in the reference dataset minus the value in this dataset. Set to NA for the reference dataset.

covBB the covariance of `meanB` in the reference data and `meanB` on this row. Used in calculating `diffBBse`.

diffBBse the standard error for `diffBB`

diffBBpValue the p -value associated with the t -test used for the hypothesis test that `diffBB` is zero

diffABAB the value of `diffAB` in the reference dataset minus the value of `diffAB` in this dataset. Set to NA for the reference dataset.

covABAB the covariance of `diffAB` in the reference data and `diffAB` on this row. Used in calculating `diffABABse`.

diffABABse the standard error for `diffABAB`

diffABABpValue the p -value associated with the t -test used for the hypothesis test that `diffABAB` is zero

sameSurvey a logical value indicating if this line uses the same survey as the reference line. Set to NA for the reference line.

Author(s)

Paul Bailey and Trang Nguyen

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# find the mean score gap in the primer data between males and females
gap("composite", sdf, dsex=="Male", dsex=="Female")

# find the score gap of the quartiles in the primer data between males and females
gap("composite", sdf, dsex=="Male", dsex=="Female", percentile=50)
gap("composite", sdf, dsex=="Male", dsex=="Female", percentile=c(25, 50, 75))

# find the percent proficient (or higher) gap in the primer data between males and females
gap("composite", sdf, dsex=="Male", dsex=="Female",
    achievementLevel=c("Basic", "Proficient", "Advanced"))

# find the discrete achievement level gap--this is harder to interpret
gap("composite", sdf, dsex=="Male", dsex=="Female",
    achievementLevel="Proficient", achievementDiscrete=TRUE)

# find the percent talk about studies at home (b017451) never or hardly
# ever gap in the primer data between males and females
gap("b017451", sdf, dsex=="Male", dsex=="Female",
    targetLevel="Never or hardly ever")

# example showing how to compare multiple levels
```

```

gap("b017451",sdf, dsex=="Male", dsex=="Female", targetLevel="Infrequently",
  recode=list(b017451=list(from=c("Never or hardly ever",
    "Once every few weeks",
    "About once a week"),
    to=c("Infrequently"))))

# make subsets of sdf by scrpsu, "Scrambled PSU and school code"
sdfA <- subset(sdf, scrpsu %in% c(5,45,56))
sdfB <- subset(sdf, scrpsu %in% c(75,76,78))
sdfC <- subset(sdf, scrpsu %in% 100:200)
sdfD <- subset(sdf, scrpsu %in% 201:300)

sdf1 <- edsurvey.data.frame.list(list(sdfA, sdfB, sdfC, sdfD),
  labels=c("A locations", "B locations",
    "C locations", "D locations"))

gap("composite", sdf1, dsex=="Male", dsex=="Female", percentile=c(50))

## End(Not run)

```

getData

Read Data to a Data Frame

Description

Reads in selected columns to a `data.frame` or a `light.edsurvey.data.frame`. On an `edsurvey.data.frame`, the data are stored on disk.

Usage

```

getData(data, varnames = NULL, drop = FALSE, dropUnusedLevels = TRUE,
  omittedLevels = TRUE, defaultConditions = TRUE, formula = NULL,
  recode = NULL, includeNaLabel = FALSE, addAttributes = FALSE,
  returnJKreplicates = TRUE)

```

Arguments

<code>data</code>	an <code>edsurvey.data.frame</code> or a <code>light.edsurvey.data.frame</code>
<code>varnames</code>	a character vector of variable names that will be returned. When both <code>varnames</code> and a <code>formula</code> are specified, variables associated with both are returned. Set to <code>NULL</code> by default.
<code>drop</code>	a logical value. When set to the default value of <code>FALSE</code> , when a single column is returned, it is still represented as a <code>data.frame</code> and is not converted to a vector.
<code>dropUnusedLevels</code>	a logical value. When set to the default value of <code>TRUE</code> , drops unused levels of all factor variables.

omittedLevels	a logical value. When set to the default value of TRUE, drops those levels of all factor variables that are specified in an <code>edsurvey.data.frame</code> . Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels.
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in an <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.
formula	a formula. When included, <code>getData</code> returns data associated with all variables of the formula. When both <code>varnames</code> and a formula are specified, the variables associated with both are returned. Set to NULL by default.
recode	a list of lists to recode variables. Defaults to NULL. Can be set as <code>recode = list(var1 = list(from = c("a", "b", "c"), to = "d"))</code> . See Examples.
includeNaLabel	a logical value to indicate if NA (missing) values are returned as literal NA values or as factor levels coded as NA.
addAttributes	a logical value set to TRUE to get a <code>data.frame</code> that can be used in calls to other functions that usually would take an <code>edsurvey.data.frame</code> . This <code>data.frame</code> is also called <code>light.edsurvey.data.frame</code> . See Details section in edsurvey.data.frame for more information on <code>light.edsurvey.data.frame</code> .
returnJKreplicates	a logical value indicating if JK replicate weights should be returned. Defaults to TRUE.

Details

By default, an `edsurvey.data.frame` does not have data read into memory until `getData` is called and returns a `data.frame`. This structure allows `EdSurvey` to have a minimal memory footprint. To keep the footprint small, you need to limit `varnames` to just the necessary variables.

When `getData` is called, it returns a `data.frame`. When the `addAttributes` argument is set to TRUE, that `data.frame` has several attributes added to make it usable by the functions in the `EdSurvey` package (e.g., `lm.sdf`), and the class is a `light.edsurvey.data.frame`.

Note that if both `formula` and `varnames` are populated, the variables on both will be included.

See the vignette titled [getData](#) for long-form documentation on this function.

Value

When `addAttributes` is FALSE, returns a `data.frame` containing data associated with requested variables. When `addAttributes` is TRUE, returns a `light.edsurvey.data.frame`.

Author(s)

Tom Fink, Paul Bailey, and Ahmad Emad

See Also

[subset.edsurvey.data.frame](#) for how to remove rows from the output

Examples

```

# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# get two variables, without weights
df <- getData(data=sdf, varnames=c("dsex", "b017451"))
table(df)

# example of using recode
df2 <- getData(data=sdf, varnames=c("dsex", "t088301"),
               recode=list(t088301=list(from=c("Yes, available", "Yes, I have access"),
                                       to=c("Yes")),
                           t088301=list(from=c("No, have no access"),
                                       to=c("No"))))

table(df2)

# When readNAEP is called on a data file, it appends a default
# condition to the edsurvey.data.frame. You can see these conditions
# by printing the sdf
sdf

# As per the default condition specified, getData restricts the data to only
# Reporting Sample. This behavior can be changed as follows:
df2 <- getData(data=sdf, varnames=c("dsex", "b017451"), defaultConditions = FALSE)
table(df2)

# Similarly, the default behavior of omitting certain levels specified
# in the edsurvey.data.frame can be changed as follows:
df2 <- getData(data=sdf, varnames=c("dsex", "b017451"), omittedLevels = FALSE)
table(df2)

# the variable "c052601" is from the school-level data file; merging is handled automatically
# returns a light.edsurvey.data.frame using addAttributes=TRUE argument
gddat <- getData(data=sdf,
                varnames=c("composite", "dsex", "b017451", "c052601"),
                addAttributes = TRUE)

class(gddat)
# look at the first few lines
head(gddat)

```

getPlausibleValue *Get Plausible Value Variables*

Description

Gets the set of variables on an edsurvey.data.frame, a light.edsurvey.data.frame, or an edsurvey.data.frame.list associated with the given subject or subscale.

Usage

```
getPlausibleValue(var, data)
```

Arguments

var a character vector naming the subject scale or subscale
data an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`

Details

This function will return a set of plausible value names for variables that `hasPlausibleValue` returns as true.

Value

a character vector of the set of variable names for the plausible values

Author(s)

Michael Lee and Paul Bailey

See Also

[updatePlausibleValue](#), [showPlausibleValues](#)

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

getPlausibleValue(var="composite", data=sdf)
```

`getWeightJkReplicates` *Retrieve the Jackknife Replicate Weights*

Description

Returns the jackknife replicate weights on an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list` associated with a weight variable.

Usage

```
getWeightJkReplicates(var, data)
```

Arguments

var character indicating the name of the weight variable for which the jackknife replicate weights are desired
data an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`

Value

a character vector of the jackknife replicate weights

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

getWeightJkReplicates(var="origwt", data=sdf)
```

 glm.sdf

EdSurvey Generalized Linear Models

Description

Fits a logit or probit that uses weights and variance estimates appropriate for the `edsurvey.data.frame`, `light.edsurvey.data.frame`, or `edsurvey.data.frame.list`.

Usage

```
glm.sdf(formula, family = binomial(link = "logit"), data,
  weightVar = NULL, relevels = list(), jrrIMax = 1,
  omittedLevels = TRUE, defaultConditions = TRUE, recode = NULL,
  returnNumberOfPSU=FALSE, returnVarEstInputs = FALSE)
```

```
logit.sdf(formula, data, weightVar = NULL, relevels = list(),
  jrrIMax = 1, omittedLevels = TRUE, defaultConditions = TRUE,
  recode = NULL, returnNumberOfPSU = FALSE,
  returnVarEstInputs = FALSE)
```

```
probit.sdf(formula, data, weightVar = NULL, relevels = list(),
  jrrIMax = 1, omittedLevels = TRUE, defaultConditions = TRUE,
  recode = NULL, returnVarEstInputs = FALSE)
```

Arguments

formula	a formula for the linear model. See <code>glm</code> . For logit and probit, we recommend using the <code>I()</code> function to define the level used for success. (See Examples.)
family	the <code>glm.sdf</code> function currently fits only the binomial outcome models, such as logit and probit, although other link functions are available for binomial models. See the <code>link</code> argument in the help for <code>family</code> .
data	an <code>edsurvey.data.frame</code>

weightVar	character indicating the weight variable to use (see Details). The weightVar must be one of the weights for the edsurvey.data.frame. If NULL, uses the default for the edsurvey.data.frame.
relevels	a list; used when the user wants to change the contrasts from the default treatment contrasts to the treatment contrasts with a chosen omitted group. The name of each element should be the variable name, and the value should be the group to be omitted.
jrrIMax	the V_{jrr} term (see Details) can be estimated with any positive number of plausible values and is estimated on the lower of the number of available plausible values and jrrIMax. When jrrIMax is set to Inf, all plausible values will be used. Higher values of jrrIMax lead to longer computing times and more accurate variance estimates.
omittedLevels	a logical value. When set to the default value of TRUE, drops those levels of all factor variables that are specified in edsurvey.data.frame. Use print on an edsurvey.data.frame to see the omitted levels.
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in an edsurvey.data.frame to subset the data. Use print on an edsurvey.data.frame to see the default conditions.
recode	a list of lists to recode variables. Defaults to NULL. Can be set as recode=list(var1=list(from=c("a", "b", "c"), to="d")). See Examples.
returnNumberOfPSU	a logical value set to TRUE to return the number of primary sampling units (PSU)
returnVarEstInputs	a logical value set to TRUE to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for the computation of covariances between estimates.

Details

This function implements an estimator that correctly handles left-hand side variables that are logical, allows for survey sampling weights, and estimates variances using the jackknife replication method. The vignette titled [Statistics](#) describes estimation of the reported statistics.

The coefficients are estimated using the sample weights according to the section “Estimation of Weighted Means When Plausible Values Are Not Present” or the section “Estimation of Weighted Means When Plausible Values Are Present,” depending on if there are assessment variables or variables with plausible values in them.

How the standard errors of the coefficients are estimated depends on the presence of plausible values (assessment variables), But once it is obtained, the t statistic is given by

$$t = \frac{\hat{\beta}}{\sqrt{\text{var}(\hat{\beta})}}$$

where $\hat{\beta}$ is the estimated coefficient and $\text{var}(\hat{\beta})$ is its variance of that estimate.

Note that `logit.sdf` and `probit.sdf` are included for convenience only; they give the same results as a call to `glm.sdf` with the binomial family and the link function named in the function call (`logit`

or probit). By default, glm fits a logistic regression when family is not set, so the two are expected to give the same results in that case. Other types of generalized linear models are not supported.

Variance estimation of coefficients: All variance estimation methods are shown in the vignette titled [Statistics](#). When the predicted value does not have plausible values, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Not Present, Using the Jackknife Method.”

When plausible values are present, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Present, Using the Jackknife Method.”

Value

An `edsurveyGlm` with the following elements:

<code>call</code>	the function call
<code>formula</code>	the formula used to fit the model
<code>coef</code>	the estimates of the coefficients
<code>se</code>	the standard error estimates of the coefficients
<code>Vimp</code>	the estimated variance due to uncertainty in the scores (plausible values variables)
<code>Vjrr</code>	the estimated variance due to sampling
<code>M</code>	the number of plausible values
<code>nPSU</code>	the number of PSUs used in calculation
<code>varm</code>	the variance estimates under the various plausible values
<code>coefm</code>	the values of the coefficients under the various plausible values
<code>coefmat</code>	the coefficient matrix (typically produced by the summary of a model)
<code>weight</code>	the name of the weight variable
<code>npv</code>	the number of plausible values
<code>njk</code>	the number of jackknife replicates used
<code>varMethod</code>	always <code>jackknife</code>
<code>varEstInputs</code>	when <code>returnVarEstInputs</code> is <code>TRUE</code> , this element is returned. These are used for calculating covariances with varEstToCov .

Author(s)

Paul Bailey

See Also

`glm`

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# By default uses jackknife variance method using replicate weights
table(sdf$b013801)
logit1 <- logit.sdf(I(b013801 %in% c("26-100", ">100"))) ~ dsex + b017451, data=sdf)
# use summary to get detailed results
summary(logit1)
logit2 <- logit.sdf(I(composite >= 300) ~ dsex + b013801, data=sdf)
summary(logit2)

logit3 <- glm.sdf(I(composite >= 300) ~ dsex + b013801, data=sdf,
                 family=quasibinomial(link="logit"))
summary(logit3)

## End(Not run)
```

hasPlausibleValue	<i>Plausible Value Test</i>
-------------------	-----------------------------

Description

Returns a value indicating if this variable has associated plausible values in an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
hasPlausibleValue(var, data)
```

Arguments

<code>var</code>	a character indicating the variable in question
<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>

Details

Note that this function returns TRUE only when the variable passed to it is the name for a set of plausible values but not if it is an individual plausible value from such a set. Thus, on the NAEP Primer, `composite` has plausible values (and so TRUE would be returned by this function), but any of the plausible values or variable names defined in the actual data (such as `"mrpcm1"` or `"dsex"`) are not.

Value

a Boolean (or vector when `var` is a vector) indicating if each element of `var` has plausible values associated with it

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# True
hasPlausibleValue(var="composite", data=sdf)

# False
hasPlausibleValue(var="dsex", data=sdf)
```

isWeight

Weight Test

Description

Returns logical values indicating whether a vector of variables is a weight for an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
isWeight(var, data)
```

Arguments

<code>var</code>	a character vector of variables
<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>

Details

Note that this function returns TRUE only when the `var` element is the name of the weight used for making estimates but not if it is one of the individual jackknife replicates.

Value

a logical vector of values indicating if each element of `var` is a weight

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# True
isWeight(var="origwt", data=sdf)

# False
isWeight(var="dsex", data=sdf)
```

levelsSDF

Print Levels and Labels

Description

Retrieve the levels and labels of a variable from an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
levelsSDF(varnames, data)
```

Arguments

varnames	a vector of character strings to search for in the database connection object (data)
data	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# search variables in the sdf
levelsSDF(varnames="pared", data=sdf)

# search multiple variables
levelsSDF(varnames=c("pared", "ell3"), data=sdf)

# search multiple variables in a light.edsurvey.data.frame with recodes
df2 <- getData(data=sdf, varnames=c("dsex", "t088301"),
               recode=list(t088301=list(from=c("Yes, available", "Yes, I have access"),
                                       to=c("Yes")),
                           t088301=list(from=c("No, have no access"),
                                       to=c("No"))),
```

```

addAttributes=TRUE)
levelsSDF(varnames=c("dsex","t088301"), data=df2)

```

lm.sdf

*EdSurvey Linear Models***Description**

Fits a linear model that uses weights and variance estimates appropriate for the data.

Usage

```

lm.sdf(formula, data, weightVar = NULL, relelevels = list(),
        varMethod = c("jackknife", "Taylor"), jrrIMax = 1,
        omittedLevels = TRUE, defaultConditions = TRUE, recode = NULL,
        returnVarEstInputs = FALSE, returnNumberOfPSU = FALSE,
        standardizeWithSamplingVar = FALSE)

```

Arguments

formula	a formula for the linear model. See <code>lm</code> . If <code>y</code> is left blank, the default subject scale or subscale variable will be used. (You can find the default using showPlausibleValues .) If <code>y</code> is a variable for a subject scale or subscale (one of the names shown by showPlausibleValues), then that subject scale or subscale is used.
data	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
weightVar	a character indicating the weight variable to use (see Details). The <code>weightVar</code> must be one of the weights for the <code>edsurvey.data.frame</code> . If <code>NULL</code> , it uses the default for the <code>edsurvey.data.frame</code> .
relelevels	a list; used when the user wants to change the contrasts from the default treatment contrasts to the treatment contrasts with a chosen omitted group. The name of each element should be the variable name, and the value should be the group to be omitted.
varMethod	a character set to “jackknife” or “Taylor” that indicates the variance estimation method to be used. See Details .
jrrIMax	when using the jackknife variance estimation method, the V_{jrr} term (see Details) can be estimated with any positive number of plausible values and is estimated on the lower of the number of available plausible values and <code>jrrIMax</code> . When <code>jrrIMax</code> is set to <code>Inf</code> , all plausible values will be used. Higher values of <code>jrrIMax</code> lead to longer computing times and more accurate variance estimates.
omittedLevels	a logical value. When set to the default value of <code>TRUE</code> , drops those levels of all factor variables that are specified in an <code>edsurvey.data.frame</code> . Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels.

defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in an <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.
recode	a list of lists to recode variables. Defaults to NULL. Can be set as <code>recode=list(var1 = list(from= c("a", "b", "c"), to= "d"))</code> . See Examples.
returnVarEstInputs	a logical value set to TRUE to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for the computation of covariances between estimates.
returnNumberOfPSU	a logical value set to TRUE to return the number of primary sampling units (PSU).
standardizeWithSamplingVar	a logical value indicating if the standardized coefficients should have the variance of the regressors and outcome measured with sampling variance. Defaults to FALSE.

Details

This function implements an estimator that correctly handles left-hand side variables that are either numeric or plausible values and allows for survey sampling weights and estimates variances using the jackknife replication method. The vignette titled [Statistics](#) describes estimation of the reported statistics.

Regardless of the variance estimation, the coefficients are estimated using the sample weights according to the sections “Estimation of Weighted Means When Plausible Values Are Not Present” or “Estimation of Weighted Means When Plausible Values Are Present,” depending on if there are assessment variables or variables with plausible values in them.

How the standard errors of the coefficients are estimated depends on the value of `varMethod` and the presence of plausible values (assessment variables). But once it is obtained, the t statistic is given by

$$t = \frac{\hat{\beta}}{\sqrt{\text{var}(\hat{\beta})}}$$

where $\hat{\beta}$ is the estimated coefficient and $\text{var}(\hat{\beta})$ is the variance of that estimate.

The **coefficient of determination (R-squared value)** is similarly estimated by finding the average R-squared using the average across the plausible values.

Variance estimation of coefficients: All variance estimation methods are shown in the vignette titled [Statistics](#). When `varMethod` is set to `jackknife` and the predicted value does not have plausible values, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Not Present, Using the Jackknife Method.”

When plausible values are present and `varMethod` is `jackknife`, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Present, Using the Jackknife Method.”

When plausible values are not present and `varMethod` is `Taylor`, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Not Present, Using the Taylor Series Method.”

When plausible values are present and `varMethod` is “Taylor,” the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Present, Using the Taylor Series Method.”

Value

An `edsurvey.lm` with the following elements:

<code>call</code>	the function call
<code>formula</code>	the formula used to fit the model
<code>coef</code>	the estimates of the coefficients
<code>se</code>	the standard error estimates of the coefficients
<code>Vimp</code>	the estimated variance from uncertainty in the scores (plausible value variables)
<code>Vjrr</code>	the estimated variance from sampling
<code>M</code>	the number of plausible values
<code>varm</code>	the variance estimates under the various plausible values
<code>coefm</code>	the values of the coefficients under the various plausible values
<code>coefmat</code>	the coefficient matrix (typically produced by the summary of a model)
<code>r.squared</code>	the coefficient of determination
<code>weight</code>	the name of the weight variable
<code>npv</code>	the number of plausible values
<code>jrrIMax</code>	the <code>jrrIMax</code> value used in computation
<code>njk</code>	the number of jackknife replicates used; set to NA when Taylor series variance estimates are used
<code>varMethod</code>	one of <code>Taylor series</code> or <code>jackknife</code>
<code>residuals</code>	residuals from the average regression coefficients
<code>PV.residuals</code>	residuals from the by plausible value coefficients
<code>PV.fitted.values</code>	fitted values from the by plausible value coefficients
<code>B</code>	imputation variance covariance matrix, before multiplication by $(M+1)/M$
<code>U</code>	sampling variance covariance matrix
<code>rbar</code>	average relative increase in variance; see van Buuren (2012, eq. 2.29)
<code>nPSU</code>	number of PSUs used in calculation
<code>n0</code>	number of rows on <code>edsurvey.data.frame</code> before any conditions were applied
<code>nUsed</code>	number of observations with valid data and weights larger than zero
<code>data</code>	data used for the computation
<code>Xstdev</code>	standard deviations of regressors, used for computing standardized regression coefficients when <code>standardizeWithSamplingVar</code> is set to <code>FALSE</code> (the default)


```

                                "About once a week"),
                                to=c("Infrequently")),
                                b017451=list(from=c("2 or 3 times a week", "Every day"),
                                to=c("Frequently"))))
# Note: "Infrequently" is the dropped level for the recoded b017451
summary(lm4)

## End(Not run)

```

merge

EdSurvey Merge

Description

Takes a `data.frame` or a `light.edsurvey.data.frame` and merge, with a `light.edsurvey.data.frame`.

Usage

```

## S3 method for class 'light.edsurvey.data.frame'
merge(x, y, ...)

```

Arguments

<code>x</code>	a <code>light.edsurvey.data.frame</code> . The attributes of the resulting <code>light.edsurvey.data.frame</code> are taken from <code>x</code> .
<code>y</code>	either a <code>light.edsurvey.data.frame</code> or a <code>data.frame</code>
<code>...</code>	arguments to be passed to <code>merge</code>

Value

a `light.edsurvey.data.frame` with the same attributes as `x`

Author(s)

Trang Nguyen

See Also

`merge`

Examples

```

# Read in NAEP primer data
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))
lsdf <- getData(data=sdf, varnames=c("dsex", "b017451"), addAttributes = TRUE)
df <- data.frame(dsex = c("Male", "Female"), dsex2 = c("Boy", "Girl"))

# Merging a light.edsurvey.data.frame with a data.frame
# returns a light.edsurvey.data.frame object

```



```
merged_1sdf <- merge(1sdf,df, by = "dsex")
class(merged_1sdf) # "light.edsurvey.data.frame" "data.frame"
head(merged_1sdf) # shows merge results

# Merging behaves similarly to base::merge
df2 <- data.frame(dsex = c("Male","Female"), b017451 = c(1,2))
merged_1sdf2 <- merge(1sdf,df2, by = "dsex")
names(merged_1sdf2) # "dsex"      "b017451.x" "b017451.y"
head(merged_1sdf2) # shows merge results
```

mixed.sdf

EdSurvey Mixed-Effects Model

Description

Fits a linear or logistic weighted mixed-effects model.

Usage

```
mixed.sdf(formula, data, weightVars = NULL,
  weightTransformation = TRUE, recode = NULL,
  defaultConditions = TRUE, tolerance = 0.01, nQuad = NULL,
  verbose = 0, family = NULL, centerGroup = NULL,
  centerGrand = NULL, fast = FALSE, ...)
```

Arguments

formula	a formula for the multilevel regression or mixed model. See Examples and the vignette titled <i>Methods Used for Estimating Mixed-Effects Models in EdSurvey</i> for more details on how to specify a mixed model. If <i>y</i> is left blank, the default subject scale or subscale variable will be used. (You can find the default using showPlausibleValues .) If <i>y</i> is a variable for a subject scale or subscale (one of the names shown by showPlausibleValues), then that subject scale or subscale is used. For logistic models, we recommend using the <code>I()</code> function to define the level used for success. (See Examples.)
data	an <code>edsurvey.data.frame</code> or a <code>light.edsurvey.data.frame</code>
weightVars	character vector indicating weight variables for corresponding levels to use. The <code>weightVar</code> must be the weights for the <code>edsurvey.data.frame</code> . The weight variables must be in the order of level (from lowest to highest level).
weightTransformation	a logical value to indicate whether the function should standardize weights before using it in the multilevel model. If set to <code>TRUE</code> , the function will look up standard weight transformation methods often used for a specific survey. Weight transformation can be found in the vignette titled <i>Methods Used for Estimating</i>

Mixed-Effects Models in EdSurvey. If set to FALSE or if the survey of the specified data does not have a standard weight transformation method, raw weights will be used.

recode	a list of lists to recode variables. Defaults to NULL. Can be set as <code>recode=list(var1 = list(from= c("a", "b", "c"), to= "d"))</code> . See Examples in <code>lm.sdf</code> .
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in an <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.
tolerance	a numeric value to indicate how accurate the result is. See Details for more information. Defaults to 0.01.
nQuad	an integer to indicate the number of quadrature points in adaptive quadrature process. See documentation of <code>WeMix::mix</code> for more details on how <code>nQuad</code> affects the estimation. If no <code>nQuad</code> is set, function will start with <code>nQuad = 5</code> and increase until a stable result is reached. See details for further discussion.
verbose	an integer; when set to 1, it will print out the brief progress of the function <code>mix.sdf</code> . Users can use these traced messages for further diagnosis. When set to 2, it will print out the detailed progress, including temporary estimates during the optimization. Defaults to 0, which will run the function without output.
family	an element of family class; optionally used to specify generalized linear mixed models. Defaults to NULL, which runs mixed linear regression models. Another family option is <code>binomial(link="logit")</code> to run binomial mixed models.
centerGroup	a list in which the name of each element is the name of the aggregation level, and the element is a formula of variable names to be group mean centered. For example, to group mean center gender and age within the group student: <code>list("student"= ~gender+age)</code> . Defaults to NULL, which means predictors are not adjusted by group centering. See Examples in <code>mix</code> .
centerGrand	a formula of variable names to be grand mean centered. For example, to center the variable education by overall mean of education: <code>~education</code> . Defaults to NULL, which means predictors are not adjusted by grand centering.
fast	a logical value; when set to TRUE, use <code>c++</code> function for faster result. Defaults to FALSE.
...	other potential arguments to be used in <code>mix</code>

Details

If users do not specify the `nQuad` argument, the functions will use the `tolerance` argument to repeatedly run the mixed model and increment `nQuad` (starting at 5) until the percentage of difference between the log-likelihood of the new model and the old model is smaller than `tolerance`. If users provide `nQuad`, selecting a smaller value of `nQuad` can save processing time; however, it is recommended that users try incrementing `nQuad` to check whether the result is stable.

Note that if the outcome variable has plausible values, the previous setting will be applied to the estimation of all plausible values.

Value

A mixedSdfResults object with the following elements:

call	the original call used in mixed.sdf
formula	the formula used to fit the model
coef	the estimates of the coefficients
se	the standard error estimates of the coefficients
vars	variance components of the model
levels	the number of levels in the model
ICC	the intraclass correlation coefficient of the model
npv	the number of plausible values
ngroups	a data.frame that includes number of observations for each group

If the formula does not involve plausible values, the function will return the following additional elements:

lnlf	the likelihood function
lnl	the log-likelihood of the model

If the formula involves plausible values, the function will return the following additional elements:

Vimp	the estimated variance from uncertainty in the scores
Vjrr	the estimated variance from sampling

Author(s)

Trang Nguyen and Claire Kelley

References

Rabe-Hesketh, S., & Skrondal, A. (2006). Multilevel modelling of complex survey data. *Journal of the Royal Statistical Society: Series A (Statistics in Society)*, 169(4), 805–827.

See Also

mix and lm.sdf

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# subset to a smaller sample
sdf_subset <- subset(sdf, scrpsu < 500)

# fast is an argument from WeMix::mix that allows the function to run faster using c++
m1 <- mixed.sdf(composite ~ dsex + b017451 + (1|scrpsu), data=sdf_subset,
```

```

weightVar = c('origwt', 'srwt01'),
fast=TRUE, verbose=1)
summary(m1)

# Run multilevel logistic regression model
# nQuad is specified to be 7, which means
# the function will use 7 quadrature points for the integration
m2 <- mixed.sdf(I(composite >= 214) ~ (1|scrpsu),
               data=sdf_subset, family = binomial(link="logit"),
               weightVar = c('origwt', 'srwt01'),
               nQuad = 7, verbose=1)
summary(m2)

## End(Not run)

```

mvrlm.sdf

*Multivariate Regression***Description**

Fits a multivariate linear model that uses weights and variance estimates appropriate for the `edsurvey.data.frame`.

Usage

```

mvrlm.sdf(formula, data, weightVar = NULL, relevels = list(),
          jrrIMax = 1, omittedLevels = TRUE, defaultConditions = TRUE,
          recode = NULL, returnVarEstInputs = FALSE, estMethod = "OLS")

```

Arguments

<code>formula</code>	a Formula for the linear model. See Formula; left hand side (LHS) variables are separated with vertical pipes (<code> </code>). See Examples.
<code>data</code>	an <code>edsurvey.data.frame</code> or <code>edsurvey.data.frame.list</code>
<code>weightVar</code>	character indicating the weight variable to use (see Details). The <code>weightVar</code> must be one of the weights for the <code>edsurvey.data.frame</code> . If <code>NULL</code> , uses the default for the <code>edsurvey.data.frame</code> .
<code>relevels</code>	a list. Used when the user wants to change the contrasts from the default treatment contrasts to treatment contrasts with a chosen omitted group. To do this, the user puts an element on the list named the same name as a variable to change contrasts on and then makes the value for that list element equal to the value that should be the omitted group. (See Examples.)
<code>jrrIMax</code>	when using the jackknife variance estimation method, the V_{jrr} term (see Details) can be estimated with any positive number of plausible values and is estimated on the first of the lower of the number of available plausible values and <code>jrrIMax</code> . When <code>jrrIMax</code> is set to <code>Inf</code> , all of the plausible values will be used. Higher values of <code>jrrIMax</code> lead to longer computing times and more accurate variance estimates.

omittedLevels	a logical value. When set to the default value of TRUE, drops those levels of all factor variables that are specified in <code>edsurvey.data.frame</code> . Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels.
defaultConditions	a logical value. When set to the default value of TRUE, uses the default conditions stored in <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.
recode	a list of lists to recode variables. Defaults to NULL. Can be set as <code>recode = list(var1= list(from=c("a", "b", "c"), to = "d"))</code> . See Examples.
returnVarEstInputs	a logical value. Set to TRUE to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for computation of covariances between estimates.
estMethod	a character value indicating which estimation method to use. Default is 'OLS', other option is 'GLS'.

Details

This function implements an estimator that correctly handles multiple left hand side variables that are either numeric or plausible values, allows for survey sampling weights and estimates variances using the jackknife replication method. The vignette titled [Statistics](#) describes estimation of the reported statistics.

The **coefficients** are estimated using the sample weights according to the section “Estimation of Weighted Means When Plausible Values Are Not Present” or the section “Estimation of Weighted Means When Plausible Values Are Present,” depending on if there are assessment variables or variables with plausible values in them.

The **coefficient of determination (R-squared value)** is similarly estimated by finding the average R-squared using the sample weights for each set of plausible values.

Variance estimation of coefficients: All variance estimation methods are shown in the vignette titled [Statistics](#).

When the predicted value does not have plausible values, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Not Present, Using the Jackknife Method.”

When plausible values are present, the variance of the coefficients is estimated according to the section “Estimation of Standard Errors of Weighted Means When Plausible Values Are Present, Using the Jackknife Method.”

For more information on the specifics of multivariate regression, see the vignette titled [Multivariate Regression](#).

Value

An `edsurvey.mvr1m` with elements:

call	the function call
formula	the formula used to fit the model

coef	the estimates of the coefficients
se	the standard error estimates of the coefficients
Vimp	the estimated variance due to uncertainty in the scores (plausible values variables)
Vjrr	the estimated variance due to sampling
M	the number of plausible values
varm	the variance estimates under the various plausible values
coefm	the values of the coefficients under the various plausible values
coefmat	the coefficient matrix (typically produced by the summary of a model)
r.squared	the coefficient of determination
weight	the name of the weight variable
npv	number of plausible values
njk	the number of jackknife replicates used
varEstInputs	When returnVarEstInputs is TRUE, this element is returned. These are used for calculating covariances with varEstToCov .
residuals	residuals for each of the PV models
fitted.values	model fitted values
residCov	residual covariance matrix for dependent variables
residPV	residuals for each dependent variables
inputs	coefficient estimation input matrices
n0	full data n
nUsed	n used for model
B	imputation variance-covariance matrix, before multiplication by $(M+1)/M$
U	sampling variance-covariance matrix

Author(s)

Alex Lishinski and Paul Bailey

See Also

lm, lm.sdf

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# Use | symbol to separate dependent variables in the left hand side of formula
mvrlm.fit <- mvrlm.sdf(algebra | geometry ~ dsex + m072801, jrrIMax = 5, data = sdf)

# print method returns coefficients, as does coef method
```

```

mvrlm.fit
coef(mvrlm.fit)

# for more detailed results use summary:
summary(mvrlm.fit)

# Details of model can also be accessed through components of the returned object, for example:

# coefficients (1 column per dependent variable)
mvrlm.fit$coef
# coefficient table with standard errors and p-values (1 table per dependent variable)
mvrlm.fit$coefmat
# R-squared values (1 per dependent variable)
mvrlm.fit$r.squared
# Residual covariance matrix
mvrlm.fit$residCov

# Model residuals and other details are available as well

# show the structure of the residuals objects
str(mvrlm.fit$residuals)
str(mvrlm.fit$residPV)

# dependent variables can have plausible values or not (or a combination)

mvrlm.fit <- mvrlm.sdf(composite | mrps22 ~ dsex + m072801, data = sdf, jrrIMax = 5)
summary(mvrlm.fit)

mvrlm.fit <- mvrlm.sdf(algebra | geometry | measurement ~ dsex + m072801, data = sdf, jrrIMax = 5)
summary(mvrlm.fit)

mvrlm.fit <- mvrlm.sdf(mrps51 | mrps22 ~ dsex + m072801, data = sdf, jrrIMax = 5)
summary(mvrlm.fit)

# Hypotheses about coefficient restrictions can also be tested using the Wald test

mvr <- mvrlm.sdf(algebra | geometry ~ dsex + m072801, data = sdf)

hypothesis <- c("geometry_dsexFemale = 0", "algebra_dsexFemale = 0")

# test statistics based on the F and Chi-squared distribution are available
linearHypothesis(mvr, hypothesis = hypothesis, test = "F")
linearHypothesis(mvr, hypothesis = hypothesis, test = "Chisq")

## End(Not run)

```

Description

Converts coefficients from `edsurveyGlm` logit regression model to odds ratios.

Usage

```
oddsRatio(model)
```

Arguments

`model` an `edsurveyGlm` model

Value

An `oddsRatio.edsurveyGlm` object with the following elements:

OR	odds ratio coefficient estimates
2.5%	lower bound 95% confidence interval
97.5%	upper bound 95% confidence interval

percentile

EdSurvey Percentiles

Description

Calculates the percentiles of a numeric variable in an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
percentile(variable, percentiles, data, weightVar = NULL, jrrIMax = 1,
  varMethod = c("jackknife", "Taylor"), alpha = 0.05,
  omittedLevels = TRUE, defaultConditions = TRUE, recode = NULL,
  returnVarEstInputs = FALSE, returnNumberOfPSU = FALSE)
```

Arguments

<code>variable</code>	the character name of the variable to percentiles computed, typically a subject scale or subscale
<code>percentiles</code>	a numeric vector of percentiles in the range 0 to 100 (inclusive)
<code>data</code>	an <code>edsurvey.data.frame</code> or an <code>edsurvey.data.frame.list</code>
<code>weightVar</code>	a character indicating the weight variable to use. (See Details.)
<code>jrrIMax</code>	a numeric value; when using the jackknife variance estimation method, the V_{jrr} term (see Details) can be estimated with any positive number of plausible values and is estimated on the lower of the number of available plausible values and <code>jrrIMax</code> . When <code>jrrIMax</code> is set to <code>Inf</code> , all plausible values will be used. Higher values of <code>jrrIMax</code> lead to longer computing times and more accurate variance estimates.

<code>varMethod</code>	a character set to <code>jackknife</code> or <code>Taylor</code> that indicates the variance estimation method used when constructing the confidence intervals. The jackknife variance estimation method is always used to calculate the standard error.
<code>alpha</code>	a numeric value between 0 and 1 indicating the confidence level. An alpha value of 0.05 would indicate a 95 percent confidence interval and is the default.
<code>omittedLevels</code>	a logical value. When set to the default value of <code>TRUE</code> , drops those levels of all factor variables that are specified in <code>achievementVars</code> and <code>aggregatBy</code> . Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels.
<code>defaultConditions</code>	a logical value. When set to the default value of <code>TRUE</code> , uses the default conditions stored in an <code>edsurvey.data.frame</code> to subset the data. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the default conditions.
<code>recode</code>	a list of lists to recode variables. Defaults to <code>NULL</code> . Can be set as <code>recode=list(var1=list(from=c("a", "b", "c"), to="d"))</code> . See Examples.
<code>returnVarEstInputs</code>	a logical value set to <code>TRUE</code> to return the inputs to the jackknife and imputation variance estimates. This is intended to allow for the computation of covariances between estimates.
<code>returnNumberOfPSU</code>	a logical value set to <code>TRUE</code> to return the number of primary sampling units (PSU)

Details

Percentiles, their standard errors, and confidence intervals are calculated according to the vignette titled [Methods Used for Estimating Percentiles](#). Note that the standard errors and confidence intervals are based on separate formulas and assumptions.

The Taylor series variance estimation procedure is not relevant to percentiles because percentiles are not continuously differentiable.

Value

The return type depends on whether the class of the data argument is an `edsurvey.data.frame` or an `edsurvey.data.frame.list`.

the data argument is an `edsurvey.data.frame`: When the data argument is an `edsurvey.data.frame`, `percentile` returns an S3 object of class `percentile`. This is a `data.frame` with typical attributes (`names`, `row.names`, and `class`) and additional attributes as follows:

n0 number of rows on `edsurvey.data.frame` before any conditions were applied

nUsed number of observations with valid data and weights larger than zero

nPSU number of PSUs used in calculation

call the call used to generate these results

The columns of the `data.frame` are as follows:

percentile the percentile of this row

estimate the estimated value of the percentile

se the jackknife standard error of the estimated percentile

df degrees of freedom

confInt.ci_lower the lower bound of the confidence interval
confInt.ci_upper the upper bound of the confidence interval
nsmall number of units with more extreme results, averaged across plausible values

the data argument is an edsurvey.data.frame.list: When the data argument is an `edsurvey.data.frame.list`, `percentile` returns an S3 object of class `percentileList`. This is a `data.frame` with a `call` attribute.

The columns in the `data.frame` are identical to those in the previous section, but there also are columns from the `edsurvey.data.frame.list`.

covs A column for each column in the `covs` value of the `edsurvey.data.frame.list`. See Examples.

When `returnVarEstInputs` is `TRUE`, an attribute `varEstInputs` also is returned that includes the variance estimate inputs used for calculating covariances with `varEstToCov`.

Author(s)

Paul Bailey

References

Hyndman, R. J., & Fan, Y. (1996). Sample quantiles in statistical packages. *American Statistician*, 50, 361–365.

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# get the median of the composite
percentile("composite", 50, sdf)

## Not run:
# get several percentiles
percentile("composite", c(0,1,25,50,75,99,100), sdf)
# build an edsurvey.data.frame.list
sdfA <- subset(sdf, scrpsu %in% c(5,45,56))
sdfB <- subset(sdf, scrpsu %in% c(75,76,78))
sdfC <- subset(sdf, scrpsu %in% 100:200)
sdfD <- subset(sdf, scrpsu %in% 201:300)

sdf1 <- edsurvey.data.frame.list(list(sdfA, sdfB, sdfC, sdfD),
                                labels=c("A locations",
                                          "B locations",
                                          "C locations",
                                          "D locations"))

# this shows how these datasets will be described
sdf1$covs

percentile("composite", 50, sdf1)
percentile("composite", c(25, 50, 75), sdf1)
```

```
## End(Not run)
```

```
print.achievementLevels
```

```
Print AchievementLevels Results
```

Description

Prints details of discrete and cumulative achievement levels calculated using weights and variance estimates appropriate for the `edsurvey.data.frame`.

Usage

```
## S3 method for class 'achievementLevels'
print(x, printCall = TRUE,
      printDiscrete = TRUE, printCumulative = TRUE, ...)
```

Arguments

<code>x</code>	an <code>achievementLevels</code> object
<code>printCall</code>	a logical value; by default (TRUE), prints details about plausible values and weights used for calculating achievement levels
<code>printDiscrete</code>	a logical value; by default (TRUE), prints discrete achievement levels if they are present in <code>x</code>
<code>printCumulative</code>	a logical value; by default (TRUE), prints cumulative achievement levels if they are present in <code>x</code>
<code>...</code>	these arguments are not passed anywhere and are included only for compatibility

Author(s)

Huade Huo and Ahmad Emad

```
print.edsurvey.data.frame
```

```
EdSurvey Metadata Summary
```

Description

Prints metadata regarding an `edsurvey.data.frame` or an `edsurvey.data.frame.list`

Usage

```
## S3 method for class 'edsurvey.data.frame'
print(x, printColnames = FALSE, ...)
```

Arguments

`x` an `edsurvey.data.frame` or an `edsurvey.data.frame.list`

`printColnames` a logical value; set to `TRUE` to see all column names in the `edsurvey.data.frame` or the `edsurvey.data.frame.list`

`...` these arguments are not passed anywhere and are included only for compatibility

Author(s)

Michael Lee and Paul Bailey

`print.gap` *Gap Analysis Printing*

Description

Prints labels and a results vector of a gap analysis.

Usage

```
## S3 method for class 'gap'
print(x, ..., printPercentage = TRUE)

## S3 method for class 'gapList'
print(x, ..., printPercentage = TRUE)
```

Arguments

`x` an R object representing a gap of class `gap` or `gapList`

`...` these arguments are not passed anywhere and are included only for compatibility

`printPercentage` a logical value set to `TRUE` to request printing of the percentage in the groups. Defaults to `TRUE`.

Author(s)

Paul Bailey

readCivEDICCS *Connect to ICCS and CivED Data*

Description

Opens a connection to an ICCS or CivEd (1999) data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readCivEDICCS(path, countries, dataSet = c("student", "teacher"),
  gradeLvl = c("8", "9", "12"), forceReread = FALSE, verbose = TRUE)
```

Arguments

path	a character value of the full directory to the ICCS/CivED extracted SPSS (.sav) set of data
countries	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found on Wikipedia at https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes , or other online sources. Consult the <i>ICCS/CivED User Guide</i> to help determine what countries are included within a specific testing year of ICCS/CivED. To select all countries, use a wildcard value of <code>*</code> .
dataSet	a character value of either <code>student</code> or <code>teacher</code> to indicate which set of data is returned. The student-level and teacher-level datasets cannot both be returned at the same time, unlike other IEA datasets. Note: The CivED 1999 study also included student-to-teacher data for Grade 8. Specifying <code>dataSet="student"</code> and <code>gradeLvl=8</code> will include both the student and teacher data in the resulting <code>edsurvey.data.frame</code> .
gradeLvl	a character value of the grade level to return <ul style="list-style-type: none"> • 8 = eighth grade (the default if not specified) • 9 = ninth grade • 12 = 12th grade (for CivED 1999 only)
forceReread	a logical value to force rereading of all processed data. The default value of <code>FALSE</code> will speed up the <code>readCivEDICCS</code> function by using existing read-in data already processed.
verbose	a logical value to either print or suppress status message output. The default value is <code>TRUE</code> .

Details

Reads in the unzipped files downloaded from the international database(s) using the [IEA Study Data Repository](#). Data files require the SPSS data file (.sav) format using the default filenames.

When using the `getData` function with a CivED or ICCS study `edsurvey.data.frame`, the requested data variables are inspected, and it handles any necessary data merges automatically. Note

that the school data will always be returned merged to the student data, even if only school variables are requested. Only if a 1999 CivED Grade 8 edsurvey.data.frame with teacher data variables is requested by the getData call, will cause teacher data to be merged. Many students can be linked to many teachers, which varies widely between countries, and not all countries contain teacher data.

Calling the dim function for a CivED 1999 Grade 8 edsurvey.data.frame will result in the row count as if the teacher dataset was merged. This row count will be considered the full data N of the edsurvey.data.frame, even if no teacher data were included in an analysis. The column count returned by dim will be the count of unique column variables across all data levels.

Value

an edsurvey.data.frame for a single specified country or an edsurvey.data.frame.list if multiple countries specified

Author(s)

Tom Fink

See Also

[readNAEP](#), [readTIMSS](#), [getData](#), and [downloadCivEDICCS](#)

Examples

```
## Not run:
eng <- readCivEDICCS("C:/ICCS2009/Gr8", countries = c("eng"),
                    gradeLvl = 8, dataSet = "student")
gg <- getData(eng, c("famstruc", "totwgts", "civ"))
head(gg)
edsurveyTable(civ ~ famstruc, eng)

## End(Not run)
```

readECLS_K1998

Connect to ECLS-K 1998 Data

Description

Opens a connection to an ECLS-K 1998 data file and returns an edsurvey.data.frame with information about the file and data.

Usage

```
readECLS_K1998(path = getwd(),
              filename = "eclsk_98_99_k8_child_v1_0.dat",
              layoutFilename = "Layout_k8_child.txt", forceReread = FALSE,
              verbose = TRUE)
```

readECLS_K2011 *Connect to ECLS-K 2011 Data*

Description

Opens a connection to an ECLS-K 2011 data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readECLS_K2011(path = getwd(), filename = "childK4p.dat",
  layoutFilename = "ECLSK2011_K4PUF.sps", forceReread = FALSE,
  verbose = TRUE)
```

Arguments

<code>path</code>	a character value to the full directory path(s) to the ECLS-K 2010-2011 extracted fixed-width-format (.dat) set of data files
<code>filename</code>	a character value of the name of the fixed-width-file (.dat) data file in the specified path to be read
<code>layoutFilename</code>	a character value of the filename of either the ASCII text (.txt) layout file of the filename within the specified path, OR a character value of the filename of the SPSS syntax (.sps) layout file of the filename within the specified path
<code>forceReread</code>	a logical value to force re-reading of all processed data. The default value of FALSE will speed up the read function by using existing read-in data already processed.
<code>verbose</code>	a logical value that will determine if you want verbose output while the <code>readECLS-K2011</code> function is running to indicate processing progress. The default value is TRUE.

Details

Reads in the unzipped files downloaded from the ECLS-K 2010-2011 longitudinal database.

Value

an `edsurvey.data.frame` for the ECLS-K 2010-2011 longitudinal dataset

Author(s)

Tom Fink

See Also

[readECLS_K1998](#), [readNAEP](#), [getData](#), and [downloadECLS_K](#)

Examples

```
## Not run:
#read-in student file with defaults
eclsk_df <- readECLS_K2011(path="C:/ECLS_K/2011") #using defaults
d <- getData(eclsk_df, c("childid", "c1hgt1", "c1wgt1"))
summary(d)

#read-in with parameters specified
eclsk_df <- readECLS_K2011(path = "C:/ECLS_K2011",
                           filename = "childK4p.dat",
                           layoutFilename = "ECLSK2011_K4PUF.sps",
                           forceReread = FALSE,
                           verbose = TRUE)

## End(Not run)
```

readICILS

*Connect to ICILS Data***Description**

Opens a connection to an ICILS data file residing on the disk and returns an `edsurvey.data` frame with information about the file and data.

Usage

```
readICILS(path, countries, dataSet = c("student", "teacher"),
          forceReread = FALSE, verbose = TRUE)
```

Arguments

<code>path</code>	a character value to the full directory to the ICILS extracted SPSS (.sav) set of data
<code>countries</code>	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found on Wikipedia at https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes , or other online sources. Consult the <i>ICILS User Guide</i> to help determine what countries are included within a specific testing year of ICILS. To select all countries, use a wildcard value of <code>*</code> .
<code>dataSet</code>	a character value of either <code>student</code> (the default if not specified) or <code>teacher</code> to indicate which set of data is returned. The student-level and teacher-level datasets cannot both be returned at the same time, unlike other IEA datasets.
<code>forceReread</code>	a logical value to force rereading of all processed data. The default value of <code>FALSE</code> will speed up the <code>readICILS</code> function by using existing read-in data already processed.
<code>verbose</code>	a logical value to either print or suppress status message output. The default value is <code>TRUE</code> .

Details

Reads in the unzipped files downloaded from the ICILS international database(s) using the [IEA Study Data Repository](#). Data files require the SPSS data file (.sav) format using the default file-names.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Tom Fink

See Also

[readNAEP](#), [readTIMSS](#), and [getData](#)

Examples

```
## Not run:
pol <- readICILS("C:/ICILS/2013", countries = "pol", dataSet = "student")
gg <- getData(pol, c("idstud", "cil", "is1g18b"))
head(gg)
edsurveyTable(cil ~ is1g18b, pol)

## End(Not run)
```

readNAEP

Connect to NAEP Data

Description

Opens a connection to an NAEP data file residing on the disk and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readNAEP(path, defaultWeight = "origwt", defaultPvs = "composite",
  omittedLevels = c("Multiple", NA, "Omitted"), frPath = NULL)
```

Arguments

<code>path</code>	a character value indicating the full filepath location and name of the (.dat) data file
<code>defaultWeight</code>	a character value that indicates the default weight specified in the resulting <code>edsurvey.data.frame</code> . Default value is <code>origwt</code> if not specified.

defaultPvs	a character value that indicates the default plausible value specified in the resulting <code>edsurvey.data.frame</code> . Default value is <code>composite</code> if not specified.
omittedLevels	a character vector indicating which factor levels/labels should be excluded. When set to the default value of <code>c('Multiple', NA, 'Omitted')</code> , adds the vector to the <code>edsurvey.data.frame</code> .
frPath	a character value indicating the location of the <code>fr2</code> parameter layout file included with the data companion to parse the specified filepath data file

Details

The function uses the `frPath` file layout (`.fr2`) data to read in the fixed-width data file (`.dat`), and builds the `edsurvey.data.frame`.

Value

An `edsurvey.data.frame` containing the following elements:

<code>userConditions</code>	a list containing all user conditions set using the <code>subset.edsurvey.data.frame</code> method
<code>defaultConditions</code>	the default conditions to be applied to the <code>edsurvey.data.frame</code>
<code>data</code>	an LaF object containing a connection to the student dataset on disk
<code>dataSch</code>	an LaF object containing a connection to the school dataset on disk
<code>dataTch</code>	not applicable for NAEP data; returns <code>NULL</code>
<code>weights</code>	a list containing the weights found on the <code>edsurvey.data.frame</code>
<code>pvvar</code>	a list containing the plausible values found on the <code>edsurvey.data.frame</code>
<code>subject</code>	the subject of the dataset contained in the <code>edsurvey.data.frame</code>
<code>year</code>	the year of assessment of the dataset contained in the <code>edsurvey.data.frame</code>
<code>assessmentCode</code>	the code of the dataset contained in the <code>edsurvey.data.frame</code>
<code>dataType</code>	the type of data (whether student or school) contained in the <code>edsurvey.data.frame</code>
<code>gradeLevel</code>	the grade of the dataset contained in the <code>edsurvey.data.frame</code>
<code>achievementLevels</code>	default NAEP achievement cutoff scores
<code>omittedLevels</code>	the levels of the factor variables that will be omitted from the <code>edsurvey.data.frame</code>
<code>fileFormat</code>	a <code>data.frame</code> containing the parsed information from the student <code>.fr2</code> file associated with the data
<code>fileFormatSchool</code>	a <code>data.frame</code> containing the parsed information from the school <code>.fr2</code> file associated with the data
<code>fileFormatTeacher</code>	not applicable for NAEP data; returns <code>NULL</code>
<code>survey</code>	the type of survey data contained in the <code>edsurvey.data.frame</code>

Author(s)

Tom Fink and Ahmad Emad

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))
sdf

## Not run:
# To read in an NCES file first set the directory to the ~/Data subfolder,
# then read in the appropriate .dat file:
setwd("location/of/Data")
sdf <- readNAEP(path="M36NT2PM.dat")

# Or read in the .dat file directly through the folder pathway:
sdf <- readNAEP(path="location/of/Data/M36NT2PM.dat")

## End(Not run)
```

readPIAAC

Connect to PIAAC Data

Description

Opens a connection to a PIAAC data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readPIAAC(path, countries, forceReread = FALSE, verbose = TRUE)
```

Arguments

<code>path</code>	a character value to the full directory to the PIAAC .csv files and Microsoft Excel codebook
<code>countries</code>	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found in the PIAAC codebook or https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes . If files are downloaded using <code>downloadPIAAC</code> , a country dictionary text file can be found in the filepath. You can use <code>*</code> to indicate all countries available.
<code>forceReread</code>	a logical value to force rereading of all processed data. Defaults to <code>FALSE</code> . Setting <code>forceReread</code> to be <code>TRUE</code> will cause PIAAC data to be reread and increase processing time.
<code>verbose</code>	a logical value that will determine if you want verbose output while the function is running to indicate the progress. Defaults to <code>TRUE</code> .

Details

Reads in the unzipped .csv files downloaded from the PIAAC Database using the OECD Repository (<http://www.oecd.org/skills/piaac/>). Users can use `downloadPIAAC` to download all required files automatically.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Trang Nguyen

References

Organisation for Economic Co-operation and Development. (2016). *Technical report of the survey of adult skills (PIAAC)* (2nd ed.). Paris, France: Author. Retrieved from http://www.oecd.org/skills/piaac/PIAAC_Technical_Report_2nd_Edition_Full_Report.pdf

See Also

[getData](#) and [downloadPIAAC](#)

Examples

```
## Not run:
# The following call returns an edsurvey.data.frame to PIAAC for Canada
can <- readPIAAC("C:/PIAAC", countries = "can")

# Extract a data.frame with a few variables
gg <- getData(can, c("c_d05", "ageg10lfs"))
head(gg)

# Conduct a preliminary analysis on the edsurvey.data.frame
edsurveyTable(~ c_d05 + ageg10lfs, data = can)

## End(Not run)
```

readPIRLS

Connect to PIRLS Data

Description

Opens a connection to a PIRLS data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readPIRLS(path, countries, forceReread = FALSE, verbose = TRUE)
```

Arguments

path	a character value to the full directory to the PIRLS extracted SPSS (.sav) set of data
countries	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found on Wikipedia at https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes , or other online sources. Consult the <i>PIRLS User Guide</i> to help determine what countries are included within a specific testing year of PIRLS. To select all countries, use a wildcard value of *.
forceReread	a logical value to force rereading of all processed data. The default value of FALSE will speed up the readPIRLS function by using existing read-in data already processed.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Details

Reads in the unzipped files downloaded from the PIRLS international database(s) using the [IEA Study Data Repository](#). Data files require the SPSS data file (.sav) format using the default file-names.

A PIRLS `edsurvey.data.frame` includes three distinct data levels:

- student
- school
- teacher

When the `getData` function is called using a PIRLS `edsurvey.data.frame`, the requested data variables are inspected, and it handles any necessary data merges automatically. Note that the school data will always be returned merged to the student data, even if only school variables are requested. Only if teacher variables are requested by the `getData` call, will cause teacher data to be merged. Many students can be linked to many teachers, which varies widely between countries.

Please note that calling the `dim` function for a PIRLS `edsurvey.data.frame` will result in the row count as if the teacher dataset was merged. This row count will be considered the full data N of the `edsurvey.data.frame`, even if no teacher data were included in an analysis. The column count returned by `dim` will be the count of unique column variables across all three data levels.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Tom Fink

See Also[readNAEP](#), [readTIMSS](#), [getData](#), and [downloadPIRLS](#)**Examples**

```
## Not run:
nor <- readPIRLS("C:/PIRLS2011", countries = c("nor"))
gg <- getData(nor, c("itsex", "totwgt", "rrea"))
head(gg)
edsurveyTable(rrea ~ itsex, nor)

## End(Not run)
```

readPISA

*Connect to PISA Data***Description**

Opens a connection to a PISA data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readPISA(path, database = "INT", countries, cognitive = "score",
         forceReread = FALSE, verbose = TRUE)
```

Arguments

<code>path</code>	a character vector to the full directory path(s) to the PISA extracted fixed-width files and SPSS control files (.txt)
<code>database</code>	a character to indicate a selected database. Must be one of INT (general database that most people use), CBA (computer-based database in PISA 2012 only), or FIN (financial literacy database in PISA 2012 only). Defaults to INT.
<code>countries</code>	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found in the PISA codebook or https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes . If files are downloaded using downloadPISA , a country dictionary text file can be found in the filepath.
<code>cognitive</code>	one of none, score, or response. Default is score. The PISA database often has three student files: student questionnaire, cognitive item response, and scored cognitive item response. The first file is used as the main student file with student background information. Users can choose whether to merge score or response data into the main file or not (if none).

forceReread	a logical value to force rereading of all processed data. Defaults to FALSE. Setting forceReread to be TRUE will cause PISA data to be reread and increase processing time.
verbose	a logical value that will determine if you want verbose output while the function is running to indicate the progress. Defaults to TRUE.

Details

Reads in the unzipped files downloaded from the PISA database using the OECD Repository (<http://www.oecd.org/pisa/>). Users can use `downloadPISA` to download all required files. Student questionnaire files (with weights and plausible values) are used as main files, which are then merged with cognitive, school, and parent files (if available).

The average first-time processing time for one year and one database for all countries is 10–15 minutes. If forceReread is set to be FALSE, the next time this function is called will only take 5–10 seconds.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Trang Nguyen

References

Organisation for Economic Co-operation and Development. (2017). *PISA 2015 technical report*. Paris, France: Author. Retrieved from <http://www.oecd.org/pisa/data/2015-technical-report/>

See Also

[getData](#) and [downloadPISA](#)

Examples

```
## Not run:
# The following call returns an edsurvey.data.frame to
# PISA 2012 International Database for Singapore
sgp2012 <- readPISA(path = "C:/PISA/2012", database = "INT", countries = "sgp")

# Extract a data.frame with a few variables
gg <- getData(sgp2012, c("cnt", "read", "w_fstuwt"))
head(gg)

# Conduct a preliminary analysis on the edsurvey.data.frame
edsurveyTable(read ~ st04q01 + st20q01, data = sgp2012)

## End(Not run)
```

readTALIS *Connect to TALIS Data*

Description

Opens a connection to a TALIS data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readTALIS(path, countries, isced = "b", dataLevel = "teacher",
  forceReread = FALSE, verbose = TRUE)
```

Arguments

<code>path</code>	a character vector to the full directory path(s) to the TALIS SPSS files (.sav)
<code>countries</code>	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found in the TALIS codebook or you can use https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes . You can use * to indicate all countries available.
<code>isced</code>	a character value that is one of a, b, or c. a stands for <i>Primary Level</i> , b is for <i>Lower Secondary Level</i> ; and, c is for <i>Upper Secondary Level</i> . Default to b.
<code>dataLevel</code>	a character value that indicates which data level to be used. It can be <code>teacher</code> (the default) or <code>school</code> .
<code>forceReread</code>	a logical value to force rereading of all processed data. Defaults to FALSE. Setting <code>forceReread</code> to be TRUE will cause PISA data to be reread and increase processing time.
<code>verbose</code>	a logical value that will determine if you want verbose output while the function is running to indicate the progress. Defaults to TRUE.

Details

Reads in the unzipped files downloaded from the TALIS database using the OECD Repository (<http://www.oecd.org/skills/piaac/>). If `dataLevel` is set to be `teacher`, it treats the teacher data file as the main dataset and merges school data into teacher data for each country. If `dataLevel` is `school`, it uses only the school data file. To conduct a school-level analysis with teacher variables, it is recommended that users aggregate teacher-level data first before merging it to school files.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Trang Nguyen

References

Organisation for Economic Co-operation and Development. (2014a). *TALIS 2013 technical report*. Paris, France: Author. Retrieved from <http://www.oecd.org/education/school/TALIS-technical-report-2013.pdf>

See Also

[getData](#) and [downloadTALIS](#)

Examples

```
## Not run:
# The following call returns an edsurvey.data.frame to TALIS 2013
# for US teacher-level data at secondary level
usa2013 <- readTALIS(path = "C:/TALIS/2013", isced = "b",
                    dataLevel = "teacher", countries = "usa")

# Extract a data.frame with a few variables
gg <- getData(usa2013, c("tt2g05b", "tt2g01"))
head(gg)

# Conduct a preliminary analysis on the edsurvey.data.frame
edsurveyTable(tt2g05b ~ tt2g01, data = usa2013)

## End(Not run)
```

readTIMSS

Connect to TIMSS Data

Description

Opens a connection to a TIMSS data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
readTIMSS(path, countries, gradeLvl, forceReread = FALSE,
          verbose = TRUE)
```

Arguments

<code>path</code>	a character vector to the full directory path(s) to the TIMSS extracted SPSS (.sav) set of data
<code>countries</code>	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found on Wikipedia at https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes , or other online sources. Consult the <i>TIMSS User Guide</i> documentation to help determine what countries are included within a specific testing year of TIMSS and for country code definitions. To select all countries available, use a wildcard value of <code>*</code> .

gradeLvl	a character value to indicate the specific grade level you wish to return <ul style="list-style-type: none">• 4 = fourth grade (the default if not specified)• 8 = eighth grade
forceReread	a logical value to force rereading of all processed data. The default value of FALSE will speed up the readTIMSS function by using existing read-in data already processed.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Details

Reads in the unzipped files downloaded from the TIMSS international database(s) using the [IEA Study Data Repository](#). Data files require the SPSS data file (.sav) format using the default file-names.

A TIMSS `edsurvey.data.frame` includes three distinct data levels:

- student
- school
- teacher

When the `getData` function is called using a TIMSS `edsurvey.data.frame`, the requested data variables are inspected, and it handles any necessary data merges automatically. Note that the school data will always be returned merged to the student data, even if only school variables are requested. Only if teacher variables are requested by the `getData` call, will cause teacher data to be merged. Many students can be linked to many teachers, which varies widely between countries.

Please note that calling the `dim` function for a TIMSS `edsurvey.data.frame` will result in the row count as if the teacher dataset was merged. This row count will be considered the `full` data `N` of the `edsurvey.data.frame`, even if no teacher data were included in an analysis. The column count returned by `dim` will be the count of unique column variables across all three data levels.

Beginning in TIMSS 2015, a numeracy dataset was designed to assess mathematics at the end of the primary school cycle for countries where most children are still developing fundamental mathematics skills. The numeracy dataset is handled automatically for the user and is included within the fourth-grade dataset `gradeLvl=4`. Most numeracy countries have a 4th grade dataset in addition to their numeracy dataset, but some do not. For countries that have both a numeracy and 4th grade dataset, the two datasets are combined into one `edsurvey.data.frame` for that country. Data variables missing from either dataset are kept, with NA values inserted for the dataset records where that variable did not exist. Data variables common to both datasets are kept as a single data variable, with records retaining their original values from the source dataset. Consult the *TIMSS User Guide* for further information.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Tom Fink

See Also[readNAEP](#), [getData](#), and [downloadTIMSS](#)**Examples**

```
## Not run:
#single country specified
fin <- readTIMSS("C:/TIMSS2015", countries = c("fin"), gradeLvl = 4)
gg <- getData(fin, c("asbg01", "totwgt", "srea"))
head(gg)
edsurveyTable(srea ~ asbg01, fin)

#multiple countries returned as edsurvey.data.frame.list, specify all countries with '*' argument
timss2011 <- readTIMSS("C:/TIMSS2011", countries="*", gradeLvl = 8, verbose = TRUE)
#print out edsurvey.data.frame.list covariates
timss2011$covs

## End(Not run)
```

readTIMSSAdv

*Connect to TIMSS Advanced Data***Description**

Opens a connection to a TIMSS Advanced data file and returns an edsurvey.data.frame with information about the file and data.

Usage

```
readTIMSSAdv(path, countries, subject = c("math", "physics"),
  forceReread = FALSE, verbose = TRUE)
```

Arguments

path	a character value to the full directory to the TIMSS Advanced extracted SPSS (.sav) set of data
countries	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found on Wikipedia at https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes , or other online sources. Consult the <i>TIMSS Advanced User Guide</i> to help determine what countries are included within a specific testing year of TIMSS Advanced. To select all countries, use a wildcard value of *.
subject	a character value to indicate if you wish to import the math or physics dataset. Only one subject can be read in at a time.

forceReread	a logical value to force rereading of all processed data. The default value of FALSE will speed up the readTIMSSAdv function by using existing read-in data already processed.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Details

Reads in the unzipped files downloaded from the TIMSS Advanced international database(s) using the [IEA Study Data Repository](#). Data files require the SPSS data file (.sav) format using the default filenames.

A TIMSSAdvanced edsurvey.data.frame includes three distinct data levels:

- student
- school
- teacher

When the getData function is called using a TIMSS Advanced edsurvey.data.frame, the requested data variables are inspected, and it handles any necessary data merges automatically. Note that the school data will always be returned merged to the student data, even if only school variables are requested. Only if teacher variables are requested by the getData call, will cause the teacher data to be merged. Many students can be linked to many teachers, which varies widely between countries.

Please note that calling the dim function for a TIMSS Advanced edsurvey.data.frame will result in the row count as if the teacher dataset was merged. This row count will be considered the full data N of the edsurvey.data.frame, even if no teacher data were included in an analysis. The column count returned by dim will be the count of unique column variables across all three data levels.

Value

an edsurvey.data.frame for a single specified country or an edsurvey.data.frame.list if multiple countries specified

Author(s)

Tom Fink

See Also

[readNAEP](#), [readTIMSS](#), [getData](#), and [downloadTIMSSAdv](#)

Examples

```
## Not run:
swe <- readTIMSSAdv("C:/TIMSSAdvanced/Math/2015",
                  countries = c("swe"), subject = "math")
gg <- getData(swe, c("itsex", "totwgt", "malg"))
head(gg)
```

```
edsurveyTable(malg ~ itsex, swe)
## End(Not run)
```

read_ePIRLS *Connect to ePIRLS Data*

Description

Opens a connection to an ePIRLS data file and returns an `edsurvey.data.frame` with information about the file and data.

Usage

```
read_ePIRLS(path, countries, forceReread = FALSE, verbose = TRUE)
```

Arguments

path	a character value to the full directory to the ePIRLS extracted SPSS (.sav) set of data
countries	a character vector of the country/countries to include using the three-digit ISO country code. A list of country codes can be found on Wikipedia at https://en.wikipedia.org/wiki/ISO_3166-1#Current_codes , or other online sources. Consult the <i>ePIRLS User Guide</i> to help determine what countries are included within a specific testing year of ePIRLS. To select all countries, use a wildcard value of <code>*</code> .
forceReread	a logical value to force rereading of all processed data. The default value of FALSE will speed up the <code>read_ePIRLS</code> function by using existing read-in data already processed.
verbose	a logical value to either print or suppress status message output. The default value is TRUE.

Details

Reads in the unzipped files downloaded from the ePIRLS international database(s) using the [IEA Study Data Repository](#). Data files require the SPSS data file (.sav) format using the default file-names.

An ePIRLS `edsurvey.data.frame` includes three distinct data levels:

- student
- school
- teacher

When the `getData` function is called using a `ePIRLS edsurvey.data.frame`, the requested data variables are inspected, and it handles any necessary data merges automatically. Note that the school data will always be returned merged to the student data, even if only school variables are requested. Only if teacher variables are requested by the `getData` call, will cause teacher data to be merged. A student can be linked to many teachers, which varies widely between countries.

Please note that calling the `dim` function for an `ePIRLS edsurvey.data.frame` will result in the row count as if the teacher dataset was merged. This row count will be considered the `full data N` of the `edsurvey.data.frame`, even if no teacher data were included in an analysis. The column count returned by `dim` will be the count of unique column variables across all three data levels.

Value

an `edsurvey.data.frame` for a single specified country or an `edsurvey.data.frame.list` if multiple countries specified

Author(s)

Tom Fink

See Also

[readNAEP](#), [readTIMSS](#), [getData](#), and [download_ePIRLS](#)

Examples

```
## Not run:
usa <- read_ePIRLS("C:/ePIRLS/2016", countries = c("usa"))
gg <- getData(usa, c("itsex", "totwgt", "erea"))
head(gg)
edsurveyTable(erea ~ itsex, usa)

## End(Not run)
```

rebindAttributes

Copy Data Frame Attributes

Description

Many R functions strip attributes from data frame objects. This function assigns the attributes from the `'attributeData'` argument to the data frame in the `'data'` argument.

Usage

```
rebindAttributes(data, attributeData)
```

Arguments

`data` a `data.frame`

`attributeData` an `edsurvey.data.frame` or `light.edsurvey.data.frame` that contains the desired attributes

Value

a `data.frame` with a class of `light.edsurvey.data.frame` containing all of the elements of `data` and the attributes (except names and `row.names`) from `attributeData`

Author(s)

Trang Nguyen and Paul Bailey

Examples

```
## Not run:
require(dplyr)
PISA2012 <- readPISA(path = paste0(edsurveyHome, "PISA/2012"),
                    database = "INT",
                    countries = "ALB", verbose=TRUE)
ledf <- getData(data = PISA2012, varnames = c("cnt", "oecd", "wfstuwt",
                                             "st62q04", "st62q11",
                                             "st62q13", "math"),
               omittedLevels = FALSE, addAttributes = TRUE)

omittedLevels <- c('Invalid', 'N/A', 'Missing', 'Miss', 'NA', '(Missing)')
for (i in c("st62q04", "st62q11", "st62q13")) {
  ledf[,i] <- factor(ledf[,i], exclude=omittedLevels)
}

#after applying some dplyr functions, the "light.edsurvey.data.frame" becomes just "data.frame"
PISA2012_ledf <- ledf %>%
  rowwise() %>%
  mutate(avg_3 = mean(c(st62q04, st62q11, st62q13), na.rm = TRUE)) %>%
  ungroup() %>%
  rebindAttributes(PISA2012) # could also be called with ledf
class(PISA2012_ledf)
# again a light.edsurvey.data.frame
lma <- lm.sdf(math ~ avg_3, data=PISA2012_ledf)
summary(lma)

PISA2012_ledf <- ledf %>%
  rowwise() %>%
  mutate(avg_3 = mean(c(st62q04, st62q11, st62q13), na.rm = TRUE)) %>%
  ungroup() %>%
  rebindAttributes(ledf) # return attributes and make a light.edsurvey.data.frame
# again a light.edsurvey.data.frame
lma <- lm.sdf(math ~ avg_3, data=PISA2012_ledf)
summary(lma)
```



```
## End(Not run)
```

```
recode.sdf                      Recode Levels Within Variables
```

Description

Recodes variables in an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
recode.sdf(x, recode)
```

Arguments

`x` an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`
`recode` a list of recoding rules. See Examples for the format of recoding rules.

Value

an object of the same class as `x` with the `recode` added to it

Author(s)

Trang Nguyen and Paul Bailey

Examples

```
## Not run:
# filepath argument will vary by operating system conventions
usaG4.15 <- readTIMSS("C:/TIMSS2015", "usa", 4)
d <- getData(usaG4.15, "itsex")
summary(d) #show details: MALE/FEMALE
usaG4.15 <- recode.sdf(usaG4.15,
                      recode = list(itsex=list(from=c("MALE"),
                                                to=c("BOY")),
                                     itsex=list(from=c("FEMALE"),
                                                to=c("GIRL"))))

d <- getData(usaG4.15, "itsex") #apply recode
summary(d) #show details: BOY/GIRL

## End(Not run)
```

`rename.sdf`*Modify Variable Names*

Description

Renames variables in an `edsurvey.data.frame`, a `light.edsurvey.data.frame` or an `edsurvey.data.frame.list`. This function often is used when users want to conduct a gap analysis across years but variable names differ across two years of data.

Usage

```
rename.sdf(x, oldnames, newnames, avoid_duplicated = TRUE)
```

Arguments

<code>x</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
<code>oldnames</code>	a character vector of old variable names
<code>newnames</code>	a character vector of new variable names to replace the corresponding old names
<code>avoid_duplicated</code>	a logical value to indicate whether to avoid renaming the variable if the corresponding new name already exists in the data. Defaults to TRUE.

Details

Note that all variable names are coerced to lowercase to comply with the EdSurvey standard.

Value

an object of the same class as `x` with new variable names

Author(s)

Trang Nguyen

See Also

[gap](#)

Examples

```
## Not run:
usaG4.15 <- readTIMSS("C:/TIMSS2015", "usa", 4)
usaG4.15.renamed <- rename.sdf(usaG4.15,
                               c("itsex", "mmat"),
                               c("gender", "math_overall"))
lm1 <- lm.sdf(math_overall ~ gender, data = usaG4.15.renamed)
summary(lm1)

## End(Not run)
```

searchSDF	<i>EdSurvey Codebook Search</i>
-----------	---------------------------------

Description

Retrieves variable names and labels for an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list` using character string matching.

Usage

```
searchSDF(string, data, fileFormat = NULL, levels = FALSE)
```

Arguments

<code>string</code>	a character string to search for in the database connection object (<code>data</code>). Note that the function will search the student, school, and teacher datasets (if applicable) for a matching character string in the codebook.
<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
<code>fileFormat</code>	a character string indicating the data source to search for variables. The default <code>NULL</code> argument searches the student, school, and teacher codebooks.
<code>levels</code>	a logical value; set to <code>TRUE</code> to return a snapshot of the levels in an <code>edsurvey.data.frame</code>

Value

a `data.frame` that shows the variable names, labels, and levels (if applicable) from an `edsurvey.data.frame` or a `light.edsurvey.data.frame` based on a matching character string

Author(s)

Michael Lee

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPrimer"))

# search both the student and school files by a character string
searchSDF(string="book", data=sdf)

# search only the student files by a character string
searchSDF(string="algebra", data=sdf, fileFormat="student")

# search both the student and school files and return a glimpse of levels
searchSDF(string="value", data=sdf, levels=TRUE)

# save the search as an object to return a full data.frame of search
ddf <- searchSDF(string="value", data=sdf, levels=TRUE)
ddf
```

 showCodebook

Summary Codebook

Description

Retrieves variable names, variable labels, and value labels for an `edsurvey.data.frame`, `light.edsurvey.data.frame`, or `edsurvey.data.frame.list`.

Usage

```
showCodebook(data, fileFormat = NULL, labelLevels = FALSE,
             includeRecodes = FALSE)
```

Arguments

<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
<code>fileFormat</code>	a character string indicating the data source to search for variables. The default <code>NULL</code> argument searches all available codebooks in the database connection object.
<code>labelLevels</code>	a logical value; set to <code>TRUE</code> to return a snapshot of the label levels in an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code> . When set to <code>FALSE</code> (the default), label levels are removed.
<code>includeRecodes</code>	a logical value; set to <code>TRUE</code> to return value labels that have been recoded in an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code> . When set to <code>FALSE</code> (the default), only the original value labels are included in the returned data frame.

Value

a data frame that shows the variable names, variable labels, value labels, value levels (if applicable), and the file format data source from an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`

Author(s)

Michael Lee

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# return codebook information for all codebooks in an edsurvey.data.frame; commonly use View()
## Not run:
View(showCodebook(sdf))

## End(Not run)
```

```
# search both the student and school files, returning levels for variable values
showCodebook(sdf, c("student","school"), labelLevels = TRUE, includeRecodes = FALSE)

# return codebook information for the student file, excluding variable value levels,
# including recoded variables
sdf <- recode.sdf(sdf, recode = list(dsex = list(from = c("Male"), to = c("MALE"))))
showCodebook(sdf, c("student"), labelLevels = FALSE, includeRecodes = TRUE)

# return codebook information for the student file, including variable value levels
# and recoded variables
showCodebook(sdf, c("student","school"), labelLevels = FALSE, includeRecodes = TRUE)
```

showCutPoints	<i>Retrieve Achievement Level Cutpoints</i>
---------------	---

Description

Retrieves a summary of the achievement level cutpoints for a selected study represented in an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
showCutPoints(data)
```

Arguments

`data` an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# show the cut points
showCutPoints(data=sdf)
```

showPlausibleValues *Plausible Value Variable Names*

Description

Prints a summary of the subject scale or subscale and the associated variables for their plausible values for an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
showPlausibleValues(data, verbose = FALSE)
```

Arguments

<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
<code>verbose</code>	a logical value; set to <code>TRUE</code> to get the variable names for plausible values. Otherwise, prints only the subject scale or subscale names for variables that use plausible values.

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# show the plausible values
showPlausibleValues(data=sdf, verbose=TRUE)
```

showWeights *Retrieve Weight Variables*

Description

Prints a summary of the weights in an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
showWeights(data, verbose = FALSE)
```

Arguments

data	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>
verbose	a logical value; set to <code>TRUE</code> to print the complete list of jackknife replicate weights associated with each full sample weight; otherwise, prints only the full sample weights

Author(s)

Michael Lee and Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# show the weights
showWeights(sdf, TRUE)
```

subset	<i>EdSurvey Subset</i>
--------	------------------------

Description

Subsets an `edsurvey.data.frame`, an `edsurvey.data.frame.list`, or a `light.edsurvey.data.frame`.

Usage

```
## S3 method for class 'edsurvey.data.frame'
subset(x, subset, ..., inside = FALSE)
```

Arguments

x	an <code>edsurvey.data.frame</code> , an <code>edsurvey.data.frame.list</code> , or a <code>light.edsurvey.data.frame</code>
subset	a logical expression indicating elements or rows to keep
...	not used; included only for compatibility
inside	set to <code>TRUE</code> to prevent the substitute condition from being called on it. See Details.

Details

Note that any variables defined on condition that are not references to column names on the `edsurvey.data.frame` and are part of the environment where `subset.edsurvey.data.frame` was called will be evaluated in the environment from which `subset.edsurvey.data.frame` was called. Similar to the difficulty of using `subset` within a function call because of the call to `substitute` on condition, this function is difficult to use (with `inside` set to the default value of `FALSE`) inside another function call. See Examples for how to call this function from within another function.

Value

an object of the same class as x

Author(s)

Trang Nguyen and Paul Bailey

References

Wickham, H. (2014). *Advanced R*. Boca Raton, FL: Chapman & Hall/CRC.

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# table to compare to subsequent tables with subsets
edsurveyTable(composite ~ dsex, data=sdf, returnMeans=FALSE, returnSepct=FALSE)

# subset to just males
newsdf <- subset(x=sdf, subset= dsex == "Male")
# table of dsex after subset
edsurveyTable(composite ~ dsex, data=newsdf, returnMeans=FALSE, returnSepct=FALSE)

# Variable names that are not in the sdf get resolved in the parent frame.
# Practically, that means that the following two subset
# calls sdfM1 and sdfM2 do the same thing
male_var <- "Male"
sdfM1 <- subset(x=sdf, subset= dsex == male_var)
sdfM2 <- subset(x=sdf, subset= dsex == "Male")
table(getData(data=sdfM1, varnames="dsex"))
table(getData(data=sdfM2, varnames="dsex"))

# variable can also be resolved as members of lists
genders <- c("Male", "Female", "not a sex level")
sdfn <- subset(x=sdf, subset= dsex == genders[2])
table(getData(data=sdfn, varnames="dsex"))

# variables can also be subset using %in%
sdfM3 <- subset(x=sdf, subset= dsex %in% c("Male", "not a sex level"))
table(getData(data=sdfM3, varnames="dsex"))

# if you need to call a name on the sdf dynamically, you can use as.name
dsex_standin <- as.name("dsex")
sdfM4 <- subset(x=sdf, subset= dsex_standin == "Male")
table(getData(data=sdfM4, varnames="dsex"))

# Here is an example of how one might want to call
# subset from within a function or loop.
# First, define a few variables to use dynamically
rhs_vars <- c("dsex", "b017451")
lvls <- c("Male", "Female")
```



```

# create a parsed condition
cond <- parse(text=paste0(rhs_vars[1], " == \"",lvls[1],"\"))[[1]]

# when inside=TRUE a parsed condition can be passed to subset
dsdf <- subset(x=sdf, subset=cond, inside=TRUE)

# check the result
table(getData(data=dsdf, varnames="dsex"))

# returns data, but uses substantial memory
## Not run:
sdf[1:3, "origwt"]
head(sdf[c("origwt", "m145101")])
sdf[1:3, c("origwt", "m145101")]
sdf[1:3,]
head(sdf[,14])
head(sdf[,10:14])

# subset an edsurvey.data.frame.list
sdfA <- subset(sdf, scrpsu %in% c(5,45,56))
sdfB <- subset(sdf, scrpsu %in% c(75,76,78))
sdfC <- subset(sdf, scrpsu %in% 100:200)
sdfD <- subset(sdf, scrpsu %in% 201:300)

# construct an edsurvey.data.frame.list from these four datasets
sdf1 <- edsurvey.data.frame.list(list(sdfA, sdfB, sdfC, sdfD),
                                labels=c("A locations",
                                         "B locations",
                                         "C locations",
                                         "D locations"))

sdf12 <- subset(sdf1, dsex=="Male")
# the number of rows in each element of the sdf1
nrow(sdf1)
# the number of rows after subsetting each element to just the Males
nrow(sdf12)

## End(Not run)

```

summary2

Summarize edsurvey.data.frame Variables

Description

Summarizes edsurvey.data.frame variables.

Usage

```
summary2(data, variable, weightVar = attr(getAttributes(data, "weights"),
    "default"), omittedLevels = FALSE)
```

Arguments

<code>data</code>	an <code>edsurvey.data.frame</code> or <code>light.edsurvey.data.frame</code>
<code>variable</code>	character variable name
<code>weightVar</code>	character weight variable name. Default to be the default weight of data if exists. If the given survey data does not have a default weight, the function will produce unweighted statistics instead. Can be set to <code>NULL</code> to return unweighted statistics.
<code>omittedLevels</code>	a logical value. When set to <code>TRUE</code> , drops those levels of the specified variable. Use <code>print</code> on an <code>edsurvey.data.frame</code> to see the omitted levels. Defaults to <code>FALSE</code> .

Value

summary of weighted or unweighted statistics of a given variable in an `edsurvey.data.frame`

For categorical variables, summary results include:

N number of cases for each category. Weighted N is also produced if users choose to produce weighted statistics.

Percent percentage of each category. Weighted Percent is also produced if users choose to produce weighted statistics.

SE standard error of the percentage statistics

For continuous variables, summary results include:

N total number of cases (both valid and invalid cases)

Min. smallest value of the variable

1st Qu. first quantile of the variable

Median median value of the variable

Mean mean of the variable

3rd Qu. third quantile of the variable

Max. largest value of the variable

SD standard deviation or weighted standard deviation

NA's number of NA's in variable and in weight variables

Zero-weight's number of zero-weight cases if users choose to produce weighted statistics

If the weight option is chosen, the function produces weighted percentile and standard deviation. Refer to the vignette titled **Statistics** and the vignette titled **Percentile** for how the function calculates these statistics (with and without plausible values).

Author(s)

Trang Nguyen

See Also

[percentile](#)

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# print out summary of weighted statistics of a continuous variable
summary2(sdf, "composite")

## Not run:
# print out summary of weighted statistics of a variable including omitted levels
summary2(sdf, "b017451", omittedLevels = FALSE)

# print out summary of unweighted statistics of a variable
summary2(sdf, "composite", weightVar = NULL)

## End(Not run)
```

updatePlausibleValue *Update Plausible Value Variable Names*

Description

Changes the name used to refer to a set of plausible values from `oldVar` to `newVar` in an `edsurvey.data.frame`, a `light.edsurvey.data.frame`, or an `edsurvey.data.frame.list`.

Usage

```
updatePlausibleValue(oldVar, newVar, data)
```

Arguments

<code>oldVar</code>	a character value indicating the existing name of the variable
<code>newVar</code>	a character value indicating the new name of the variable
<code>data</code>	an <code>edsurvey.data.frame</code> , a <code>light.edsurvey.data.frame</code> , or an <code>edsurvey.data.frame.list</code>

Value

an object of the same class as the `data` argument, with the name of the plausible value updated from `oldVar` to `newVar`

Author(s)

Michael Lee and Paul Bailey

See Also

[getPlausibleValue](#) and [showPlausibleValues](#)

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package="NAEPprimer"))

# get the PVs before
showPlausibleValues(sdf)
sdf2 <- updatePlausibleValue("composite", "overall", sdf)
showPlausibleValues(sdf2)
lm1 <- lm.sdf(overall ~ b017451, data=sdf2)
summary(lm1)
```

varEstToCov

Covariance Estimation

Description

When the variance of a derived statistic (e.g., a difference) is required, the covariance between the two statistics must be calculated. This function uses results generated by various functions (e.g., a [lm.sdf](#)) to find the covariance between two statistics.

Usage

```
varEstToCov(varEstA, varEstB = varEstA, varA, varB = varA,
            jkSumMultiplier)
```

Arguments

varEstA	a list of two data.frames returned by a function after the returnVarEstInputs argument was turned on. The statistic named in the varA argument must be present in each data.frame.
varEstB	a list of two data.frames returned by a function after the returnVarEstInputs argument was turned on. The statistic named in the varA argument must be present in each data.frame. When the same as varEstA, the covariance is within one result.
varA	a character that names the statistic in the varEstA argument for which a covariance is required
varB	a character that names the statistic in the varEstB argument for which a covariance is required
jkSumMultiplier	when the jackknife variance estimation method—or balanced repeated replication (BRR) method—multiplies the final jackknife variance estimate by a value, set jkSumMultiplier to that value. For an edsurvey.data.frame, or a light.edsurvey.data.frame, the recommended value can be recovered with EdSurvey::getAttributes(myData, "jkSumMultiplier").

Details

Note that these functions are not vectorized, so varA and varB must contain exactly one variable name.

The method used to compute the covariance is in the vignette titled [Statistics](#) in the section “Estimation of Covariances.”

The method used to compute the degrees of freedom is in the vignette titled [Statistics](#) in the section “Estimation of Degrees of Freedom.”

Value

a numeric value; the jackknife covariance estimate

Author(s)

Paul Bailey

Examples

```
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# estimate a regression
lm1 <- lm.sdf(composite ~ dsex + b017451, sdf, returnVarEstInputs=TRUE)
summary(lm1)
# estimate the covariance between two regression coefficients
# note that the variable names are parallel to what they are called in lm1 output
covFEveryDay <- varEstToCov(lm1$varEstInputs,
                             varA="dsexFemale",
                             varB="b017451Every day",
                             jkSumMultiplier=EdSurvey::getAttributes(sdf, "jkSumMultiplier"))
# the estimated difference between the two coefficients
# note: unname prevents output from being named after the first coefficient
unname(coef(lm1)["dsexFemale"] - coef(lm1)["b017451Every day"])
# the standard error of the difference
# uses the formula SE(A-B) = sqrt(var(A) + var(B) - 2*cov(A,B))
sqrt(lm1$coefmat["dsexFemale", "se"]^2
      + lm1$coefmat["b017451Every day", "se"]^2
      - 2 * covFEveryDay)
```

waldTest

Wald Tests

Description

Tests on coefficient(s) of edsurveyGlm and edsurveyLm models.

Usage

```
waldTest(model, coefficients, H0 = NULL)
```

Arguments

model	an edsurveyGlm and edsurveyLm
coefficients	coefficients to be tested, by name or position in coef vector. See Details.
H0	reference values to test coefficients against, default = 0

Details

When plausible values are present, likelihood ratio tests cannot be used. However, the Wald test can be used to test estimated parameters in a model, with the null hypothesis being that a parameter(s) is equal to some value(s). In the default case where the null hypothesis value of the parameters is 0, if the test fails to reject the null hypothesis, removing the variables from the model will not substantially harm the fit of that model. Alternative null hypothesis values can also be specified with the H0 argument. See Examples.

Coefficients to test can be specified by an integer (or integer vector) corresponding to the order of coefficients in the summary output. Coefficients can also be specified using a character vector, to specify coefficient names to test. The name of a factor variable can be used to test all levels of that variable.

This test produces both chi-square and F-tests; their calculation is described in the vignette titled [Wald Test](#).

Value

An edsurveyWaldTest object with the following elements:

Sigma	coefficient covariance matrix
coefficients	indices of the coefficients tested
H0	null hypothesis values of coefficients tested
result	result object containing the values of the chi-square and F-tests
hypoMatrix	hypothesis matrix used for the Wald test

Author(s)

Alex Lishinski and Paul Bailey

References

- [UCLA IDRE - FAQ:How are the likelihood ratio, Wald, and Lagrange multiplier (score) tests different and/or similar?](<https://stats.idre.ucla.edu/other/mult-pkg/faq/general/faqhow-are-the-likelihood-ratio-wald-and-lagrange-multiplier-score-tests-different-and-or-similar/>)
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- Draper, N. R., & Smith, H. (1998). *Applied regression analysis*. New York, NY: Wiley.
- Fox, J. (1997). *Applied regression analysis, linear models, and related methods*. Thousand Oaks, CA: SAGE.
- Korn, E., & Graubard, B. (1990). Simultaneous testing of regression coefficients with complex survey data: Use of Bonferroni t statistics. *The American Statistician*, 44(4), 270–276.

Examples

```
## Not run:
# read in the example data (generated, not real student data)
sdf <- readNAEP(system.file("extdata/data", "M36NT2PM.dat", package = "NAEPprimer"))

# example with glm model
myLogit <- logit.sdf(dsex ~ b017451 + b003501, data = sdf, returnVarEstInputs = T)

# single coefficient integer
waldTest(model = myLogit, coefficients = 2)

# set of coefficients integer vector
waldTest(model = myLogit, coefficients = 2:5)

# specify levels of factor variable to test
waldTest(myLogit, c("b017451Every day", "b017451About once a week"))

# specify all levels of factor variable to test
waldTest(myLogit, "b017451")

# example with lm model
fit <- lm.sdf(composite ~ dsex + b017451, data = sdf, returnVarEstInputs = T)

waldTest(model = fit, coefficients = "b017451")

# examples with alternative (nonzero) null hypothesis values
waldTest(model = myLogit, coefficients = 2, H0 = 0.5)

waldTest(model = myLogit, coefficients = 2:5, H0 = c(0.5, 0.6, 0.7, 0.8))

waldTest(model = myLogit, coefficients = "b017451", H0 = c(0.5, 0.6, 0.7, 0.8))

waldTest(model = myLogit, coefficients = c("b017451Every day", "b017451About once a week"),
          H0 = c(0.1, 0.2))

## End(Not run)
```

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