

Package ‘imputevalR’

December 13, 2021

Title Multiple Imputation of Missing Data Using Predictive Sampling

Version 0.1

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Description imputevalR contains methods to impute all missing values in a data frame for all variables in the data, so long as the variables are either continuous or binary. The imputation uses a Gibbs sampler to draw from the predictive distributions of the variable of interest, conditional on all other variables in the data set, and then returns a list of data.frames that can be used to complete pooled regression analyses (e.g. using Rubin's rules) or other sample statistics.

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Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.1.2

LazyData true

Imports Rcpp (>= 1.0.7),
methods,
Rdpack,
stats

Depends R (>= 2.10)

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compare_by_cell	<i>Squared error between means for all cells in a dataset</i>
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Description

Squared error between means for all cells in a dataset

Usage

```
compare_by_cell(df1, df2)
```

Arguments

df1	The first data.frame for which comparisons should be made
df2	The second data.frame for which comparisons should be made

Value

summed_diff, a named vector of length equal to the number of variables, where each value is the summed squared difference between the imputed and true value of a cell, for each variable

compare_by_column	<i>Squared error between means for all columns in a dataset</i>
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Description

Squared error between means for all columns in a dataset

Usage

```
compare_by_column(df1, df2)
```

Arguments

df1	The first data.frame for which comparisons should be made
df2	The second data.frame for which comparisons should be made

Value

squarediff, a named vector with length equal to the the number of variables for the comparison

`imputer`*Run Imputation Procedure and Generate Imputed Datasets*

Description

Takes in a data.frame with missing values, and runs the imputation algorithm to return 5 imputed data sets (stored as a list) with missing values replaced by imputed values.

Usage

```
imputer(df, nchains = 5, niter = 100)
```

Arguments

<code>df</code>	A data.frame for which imputed data sets should be generated
<code>nchains</code>	The number of chains (ie number of imputed datasets) to generate
<code>niter</code>	The number of iterations to complete in each chain (default is 100 for convergence)

Value

A list (returnSets) that contains 5 imputed data.frames with missing values that were originally in df with imputed values based on the predicted imputed values

Examples

```
data(nhanes)
miss_data <- imputevalR::makeNA(nhanes, proportionNA = 0.2)
imputed <- imputer(miss_data, nchains = 1, niter = 5)
```

`makeNA`*Set random values in a data frame to missing*

Description

Set random values in a data frame to missing

Usage

```
makeNA(df, proportionNA = 0.2)
```

Arguments

<code>df</code>	A data.frame for which NA values should be generated
<code>proportionNA</code>	The proportion of all cells that should be NA, across variables and observations

Value

A data.frame (df) that is the same dimensions as the input data frame, but now has values missing (set by proportionNA)

Examples

```
data(nhanes)
miss_data <- imputevalR::makeNA(nhanes, proportionNA = 0.2)
```

nhanes

Example NHANES data with no missing values

Description

A dataset containing sample variables from the National Health and Nutrition Examination Survey (2017-18) cycle

Usage

```
nhanes
```

Format

A data frame with 9254 rows and 21 variables:

gender gender (male/female)

age age in years

white Non-Hispanic White race/ethnicity

poverty ratio of household income to federal poverty level

weight measured weight in kg

height measured height in cm

bmi BMI in kg/m²

waist_circum waist circumference in cm

hip_circum hip circumference in cm

sbp1 systolic blood pressure (4 readings)

sbp2 systolic blood pressure (4 readings)

sbp3 systolic blood pressure (4 readings)

sbp4 systolic blood pressure (4 readings)

dbp1 diastolic blood pressure (4 readings)

dbp2 diastolic blood pressure (4 readings)

dbp3 diastolic blood pressure (4 readings)

dbp4 diastolic blood pressure (4 readings)

android_pfat percent fat in android

gynoid_pfat percent fat in gyneoid

selfreported_weight self rep weight in lbs

selfreported_ht self rep height in inches

Source

<https://www.cdc.gov/nchs/nhanes/index.htm>

pool_df	<i>Pool imputed data frame results</i>
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Description

Pool imputed data frame results

Usage

```
pool_df(dfList)
```

Arguments

dfList A list of imputed data.frames generated by imputer()

Value

finalDF: A single data.frame containing values that have been pooled across the imputations using Rubin's rules

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