Package 'SURF'

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Type Package

Description

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Title Some Useful R Functions

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SURF-package Some Useful R Functions					

so I will package them for easy access/dissemination.

These are several useful functions that I find myself using often (read: frequently re-implementing),

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Details

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Author(s)

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Examples

```
## Simulate regression data:
testData <- simRegData(nObs = 100,
                       nVars = 10,
                       r2 = 0.5,
                       sigma = 0.2,
                       beta = matrix(c(0.25, rep(0.75, 10)))
## Impose missing data
missData <- imposeMissData(data</pre>
                                   = testData,
                           targets = list(mar = c("y", "x1"),
                                           mcar = c("x2", "x3"),
                                           mnar = c("x4", "x5")
                            preds = c("x8", "x9", "x10"),
                                   = list(mar = 0.2, mcar = 0.1, mnar = 0.1),
                                   = list(mar = 5, mnar = 2.5),
                           pattern = "random")
## Plot imputed vs. observed values:
data(testImps)
plotImps(impList = testImps$impList,
        rMat
              = testImps$rMat,
         typeVec = testImps$typeVec)
```

calcMode

Find the Modal Value of a Vector

Description

This function will find the most commonly occurring value (i.e., the mode) in a vector.

```
calcMode(x, discrete = TRUE)
```

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Arguments

x A vector for which to find the mode.

discrete A logical switch indicate whether x is a categorical (i.e., discrete = TRUE) or continuous (i.e., discrete = FALSE) variable.

Details

In the event of multiple modes, ties are broken randomly. When x is numeric, the return value will be a length one numeric vector, otherwise, the return value will be a length one character vector.

When discrete = FALSE the modal value is estimated via kernel density estimation by applying stats::density to x.

Value

The modal value in x.

Author(s)

Kyle M. Lang

Examples

```
## Find mode of an integer vector:
x1 <- sample(c(1 : 5), 100, TRUE)
calcMode(x1)

## Find mode of a factor:
x2 <- factor(sample(c(1 : 5), 100, TRUE))
calcMode(x2)

## Find mode of a character vector:
x3 <- sample(letters[1 : 5], 100, TRUE)
calcMode(x3)

## Find the mode of a continuous variable:
x4 <- rnorm(1000, 5, 2.5)
calcMode(x4, discrete = FALSE)</pre>
```

f2n

Safely Cast a Factor to a Numeric Vector

Description

This function will type-cast a factor to a numeric vector without mangling the levels when those levels are zero-indexed integers.

Usage

f2n(x)

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Arguments

Х

A factor to be converted.

Details

R indexes factor levels from 1, so type-casting factors that have zero-indexed integer labels using as.numeric can produce unanticipated results. This function will cast such factors to zero-indexed numeric vectors wherein the original factor *labels* are mapped to numeric values (rather than the inherent factor *levels* being so mapped).

Value

A numeric vector containing the type-cast version of x.

Author(s)

```
Kyle M. Lang
```

Examples

```
## Generate a factor:
x1 <- factor(c(0 : 5))
x1

## Naive type-casting:
as.numeric(x1)

## Safe type-casting:
f2n(x1)</pre>
```

imposeMissData

Impose Missing Data

Description

Impose missing data according to MAR, MCAR, and MNAR mechanisms.

imposeMissData 5

Arguments

data A data.frame wherein missing data are to be imposed.

targets A named list with slots "mar", "mcar", and "mnar" containing character vectors

giving the column labels for variables onto which missing at random, missing completely at random, and missing not at random data, respectively, shall be

imposed.

preds A character vector giving the column labels for predictors of the MAR missing-

ness.

pm A named list with slots "mar", "mcar", and "mnar" containing real numbers

in [0, 1) giving the proportions of missing at random, missing completely at

random, and missing not at random data, respectively, to generate.

snr A named list with slots "mar" and "mnar" containing real numbers giving the

signal-to-noise ratios of the probit regression models used to impose missing at

random and missing not at random data, respectively.

pattern A (possibly named) character vector indicating in what parts of the missing data

predictors' distributions MAR and MNAR missing data should be imposed. See

Details section for more information.

Details

MCAR missing data is imposed by generating a random Bernoulli flag variable for each target variable with probability of success equal to pm\$mcar.

MAR missing data is imposed via a noisy probit regression model wherein the weighted sum of the columns listed in preds are used to predict the response propensity.

MNAR missing data is imposed via the same procedure as MAR missing data but the missing data predictor is taken to be the target variable itself.

Legal keywords for the pattern argument are: "low" = impose missing in the negative tail of the predictor, "high" = impose missing in the positive tail of the predictor, "center" = impose missing in the center of the predictor, "tails" = impose missing, symmetrically, in both tails of the predictor, "random" = randomly sample from the previous four keywords.

The pattern argument can be given as a length-one character vector, in which case the specified keyword is applied to all applicable variables. Otherwise, the pattern argument must be a named vector with length equal to length(c(targets\$mar, targets\$mnar)) and names matching all elements of c(targets\$mar, targets\$mnar).

Value

A two-element list with the following slots:

data: The incomplete version of data

pattern: A character vector showing which pattern was used for each target variable.

Note

Due to the stochastic nature of the missing data simulation, the actual proportions of missing data will only equal the values provided for pm asymptotically.

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Author(s)

Kyle M. Lang

See Also

simRegData

Examples

```
## Simulate some data:
testData <- simRegData(nObs = 100,
                       nVars = 10,
                       r2 = 0.5,
                       sigma = 0.2,
                       beta = matrix(c(0.25, rep(0.75, 10)))
## Impose missing data:
missData <- imposeMissData(data</pre>
                                    = testData,
                           targets = list(mar = c("y", "x1"),
                                           mcar = c("x2", "x3"),
                                           mnar = c("x4", "x5")
                                           ),
                            preds = c("x8", "x9", "x10"),
                                   = list(mar = 0.2, mcar = 0.1, mnar = 0.1),
                                   = list(mar = 5, mnar = 2.5),
                            pattern = "random")
```

plotImps

Plot Imputed vs. Observed Values

Description

This function will generate plots of imputed versus observed values in multiply imputed data. These plots can be examined to "sanity-check" the imputation procedure.

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Arguments

impList A list of multiply imputed datasets.

rMat A logical pattern matrix flagging missing values in the orginal data used to gen-

erate the imputed datasets in impList. Note that TRUE flags missing cells and

FALSE flags observed cells.

typeVec A character vector with length(typeVec) = ncol(impList[[1]]) giving the

measurement levels of the variables in impList. Two values are recognized: "cat" = a categorical variable (i.e., nominal or ordinal) and "con" = a continuous

variable (i.e., interval or ratio).

targetVar An optional character vector giving the column names for variables to plot.

When targetVar = NULL all variables with imputed values are plotted.

interactive A logical flag: Should the cycle through all plotted variables by prompting the

user to continue after generating each plot?

Value

Used for its side-effects.

Author(s)

```
Kyle M. Lang
```

Examples

rangeNorm	Range Normalize a Vector

Description

This function will standardize a vector so that all transformed values exist between user-defined minimum and maximum values.

```
rangeNorm(x, oldMin = min(x), oldMax = max(x), newMin = 0.0, newMax = 1.0)
```

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Arguments

Х	A numeric vector to be standardized.
oldMin	An optional real number giving the minimum possible value of x.
oldMax	An optional real number giving the maximum possible value of x.
newMin	An optional real number giving the minimum possible value of the transformed input vector.
newMax	An optional real number giving the maximum possible value of the transformed input vector.

Value

A numeric vector containing the range-normalized version of x.

Author(s)

```
Kyle M. Lang
```

Examples

```
x0 <- runif(100, -3, 5)
x1 <- rangeNorm(x = x0)
range(x1)</pre>
```

simCovData

Simulate Data with Known Covariance Structure

Description

This function will simulation simple data with a known covariance structure.

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Arguments

n0bs	An integer giving the number of rows to simulate.
sigma	Either a numeric matrix giving the covariance matrix of the predictor variables or a length-one numeric vector with value in [-1.0, 1.0]. In the latter case, sigma gives the correlation between the predictors (i.e., the degree of collinearity).
nVars	An integer giving the number of (possibly latent) predictor variables to simulate. Defaults to ncol(sigma).
means	A numeric vector of predictor means. Recycled when length(means) = 1 to match nVars, otherwise length(means) must equal nVars.
scales	A numeric vector of predictor scales. Recycled when length(scales) = 1 to match nVars, otherwise length(scales) must equal nVars. Ignored when sigma is a matrix.

Details

If sigma is a length-one vector, a covariance matrix is constructed from the values provided for sigma and scales. In this case, sigma gives the *correlation* between predictors (not the covariance) and scales gives the predictors' standard deviations.

Value

An nObs by nVars data.frame of simulated data.

Note

The column labels of the simulated data will be paste0("x", 1 : nVars).

Author(s)

```
Kyle M. Lang
```

See Also

imposeMissData simRegData

Examples

```
## Specify 'sigma' as a full covariance matrix:
sigma      <- matrix(0.3, 10, 10)
diag(sigma) <- 1.0

testData <- simCovData(nObs = 100, sigma = sigma)

## Specify 'sigma' as an inter-predictor correlation:
testData <- simCovData(nObs = 100, nVars = 10, sigma = 0.2)</pre>
```

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simRegData	Simulate Regression Data	

Description

This function will simulation regression data with known R-Squared, inter-predictor correlation, and latent grouping structure among the predictors.

Usage

indicators.

Arguments

n0bs	An integer giving the number of rows to simulate.		
r2	A real number in [0, 1]. The R-Squared of the data generating model. That is, what proportion of variability in the outcome should be explained by the predictors.		
sigma	Either a numeric matrix giving the covariance matrix of the predictor variables or a length-one numeric vector with value in [-1.0, 1.0]. In the latter case, sigma gives the correlation between the predictors (i.e., the degree of collinearity).		
beta	An numeric matrix of regression coefficients with $dim(beta) = c(nVars, 1)$. Note that the first element is taken to be the intercept.		
nVars	An integer giving the number of (possibly latent) predictor variables to simulate. Defaults to ncol(sigma).		
means	A numeric vector of predictor means. Recycled when length(means) = 1 to match nVars, otherwise length(means) must equal nVars.		
scales	A numeric vector of predictor scales. Recycled when length(scales) = 1 to match nVars, otherwise length(scales) must equal nVars. Ignored when sigma is a matrix.		
itemsPerPred	An integer giving the number of observed items used to define each latent predictor. When itemsPerPred = 1, no latent structure is imposed on the predictors.		
predReliability	**-**		
	A real number in [0, 1]. When itemsPerPred > 1, predReliability defines the proportion of reliable variance among the indicators of each latent predictor. That is, the proportion of shared variance among each latent predictor's observed		

).

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Details

If sigma is a length-one vector, a covariance matrix is constructed from the values provided for sigma and scales. In this case, sigma gives the *correlation* between predictors (not the covariance) and scales gives the predictors' standard deviations.

Value

An nObs by nVars * itemsPerPred + 1 data.frame of simulated data.

Note

The column labels of the simulated data will be c("y", paste0("x", 1 : nVars * itemsPerPred)).

Author(s)

Kyle M. Lang

See Also

imposeMissData simCovData

Examples

testImps

List of Example Data Elements

Description

This data object is a list of data elements meant to demonstrate usage of the plotImps function.

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Usage

```
data("testImps")
```

Format

A list with the following three elements.

impList: A list containing 10 multiply imputed datasets.

rMat: A logical pattern matrix flagging missing values in the data that were imputed to generate impList.

typeVec: A character vector giving the measurement level of each variable in impList.

See Also

plotImps

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