## Package ‘BinNor’

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BinNor-package

A package for simultaneous generation of binary and normal data.

Description

Provides R functions for generating multiple binary and normal variables simultaneously given the marginal characteristics and association structure via combining well established results from the random number generation literature, based on the methodology proposed by Demirtas and Doganay (2012).

Details

Package: BinNor
Type: Package
Version: 2.3.2
Date: 2020-04-20
License: GPL-2
LazyLoad: yes

There are eight functions in this package. The functions lower.tri.to.corr.mat, validation.bin, validation.nor, validation.range and validation.nor are designed to prevent obvious specification errors and to validate the specified quantities. The most important functions are compute.sigma.star, jointly.generate.binary.normal and simulation. The function compute.sigma.star computes the matrix of tetrachoric correlations that will be used in the generation of multivariate normal data whose some components are dichotomized to obtain binary variables. The function jointly.generate.binary.normal generates mixed data, and the function simulation is capable of repeating this process many times and produces averages of some key statistical quantities across replications.

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References


compute.sigma.star  Computes intermediate correlation matrix

Description

This function computes the intermediate correlation matrix by assembling tetrachoric correlations for binary-binary combinations, biserial correlations for binary-normal combinations, and specified correlation for normal-normal combinations. If the resulting correlation matrix is not positive definite, a nearest positive matrix will be used.

Usage

compute.sigma.star(no.bin, no.nor, prop.vec.bin = NULL, corr.vec = NULL, corr.mat = NULL)

Arguments

- no.bin: Number of binary variables
- no.nor: Number of normal variables
- prop.vec.bin: Probability vector for binary variables
- corr.vec: Vector of elements below the diagonal of correlation matrix ordered columnwise
- corr.mat: Specified correlation matrix

Value

- sigma_star: A resulting intermediate correlation matrix \( \Sigma^* \)
- nonPD: If a resulting intermediate correlation matrix is non-positive definite, it is stored in this value. Otherwise it is NULL.
- PD: TRUE if \( \Sigma^* \) is positive definite, FALSE otherwise. A FALSE indicates that the nearest positive definite matrix is returned.
- eigenv: Eigenvalues of the \( \Sigma^* \) before the conversion

See Also

- validation.corr, nearPD, phi2tetra, is.positive.definite,
- jointly.generate.binary.normal, simulation

Examples

```R
cmat = lower.tri.to.corr.mat(corr.vec= c(0.16, 0.04, 0.38, 0.14, 0.47, 0.68),4) compute.sigma.star(no.bin=2, no.nor=2, prop.vec.bin=c(0.4,0.7), corr.vec=NULL,corr.mat=cmat)
```
**jointly.generate.binary.normal**

*Generates a mix of binary and normal data*

**Description**

Generates multiple binary and normal variables simultaneously given marginal characteristics and association structures.

**Usage**

```r
jointly.generate.binary.normal(no.rows, no.bin, no.nor, prop.vec.bin = NULL, mean.vec.nor = NULL, var.nor = NULL, sigma_star = NULL, corr.vec = NULL, corr.mat = NULL, continue.with.warning = TRUE)
```

**Arguments**

- `no.rows` Number of rows.
- `no.bin` Number of binary variables
- `no.nor` Number of normal variables
- `prop.vec.bin` Probability vector for binary variables
- `mean.vec.nor` Vector of means for normal variables
- `var.nor` Vector of variances for normal variables
- `sigma_star` Intermediate correlation matrix
- `corr.vec` Vector of elements below the diagonal of correlation matrix ordered columnwise
- `corr.mat` Specified correlation matrix
- `continue.with.warning` TRUE to proceed with the nearest positive definite $\Sigma^*$. FALSE to terminate program execution if $\Sigma^*$ is not positive definite

**Value**

- `data` A matrix of generated data.

**See Also**

- `compute.sigma.star`, `validation.corr`, `validation.bin`, `validation.nor`, `nearPD`, `simulation`, `rmvnorm`
lower.tri.to.corr.mat

Examples

no.rows=100
no.bin=2; no.nor=2
mean.vec.nor=c(3,1); var.nor=c(4,2)
prop.vec.bin=c(0.4,0.7)
corr.vec=c(0.16,0.04,0.38,0.14,0.47,0.68);

cmat = lower.tri.to.corr.mat(corr.vec,4)
sigma.star=compute.sigma.star(no.bin=2, no.nor=2, prop.vec.bin=c(0.4,0.7),
corr.mat=cmat)
mydata=jointly.generate.binary.normal(no.rows,no.bin,no.nor,prop.vec.bin,
mean.vec.nor,var.nor, sigma_star=sigma.star$sigma_star,
continue.with.warning=TRUE)

lower.tri.to.corr.mat  Converts a vector of correlations to a full correlation matrix

Description

This function creates full correlation matrix from the vector containing elements below the diagonal.

Usage

lower.tri.to.corr.mat(corr.vec = NULL, d)

Arguments

corr.vec A vector of elements below diagonal of correlation matrix. The elements must be ordered starting from first element below diagonal of the first column, then second element below diagonal of the first column and so on.
d Number of column in final correlation matrix.

Value

corr.mat Full correlation matrix

See Also

lower.tri

Examples

corr.vec=c(0.16,0.04,0.38,0.14,0.47,0.68)
lower.tri.to.corr.mat(corr.vec,4)
simulation

Repeats the data generation process in a simulation scheme

Description

Simulates many versions of mixed data, and reports averaged proportion, mean, variance and correlation estimates across replications.

Usage

simulation(seed = NULL, nsim, no.rows, no.bin, no.nor, mean.vec.nor = NULL, var.nor = NULL, prop.vec.bin = NULL, corr.vec = NULL, corr.mat = NULL, continue.with.warning = TRUE)

Arguments

- **seed**: A seed value for the random number generator. Seed value will be randomly generated unless specified.
- **nsim**: Number of simulation runs.
- **no.rows**: Number of rows.
- **no.bin**: Number of binary variables
- **no.nor**: Number of normal variables
- **prop.vec.bin**: Probability vector for binary variables
- **mean.vec.nor**: Vector of means for normal variables
- **var.nor**: Vector of variances for normal variables
- **corr.vec**: Vector of elements below the diagonal of correlation matrix ordered columnwise
- **corr.mat**: Specified correlation matrix
- **continue.with.warning**: TRUE to proceed with the nearest positive definite $\Sigma^*$. FALSE to terminate program execution if $\Sigma^*$ is not positive definite

See Also

compute.sigma.star, jointly.generate.binary.normal

Examples

simulation(nsim=10, no.rows=100, no.bin=2, no.nor=2, mean.vec.nor=c(3,1), var.nor=c(4,2), prop.vec.bin=c(0.4,0.7), corr.vec=c(0.16,0.04,0.38,0.14,0.47,0.68), corr.mat=NULL)
validation.bin

Validates the marginal specification of the binary part

Description

Checks whether the marginal specification of the binary part is valid and consistent.

Usage

validation.bin(no.bin, prop.vec.bin = NULL)

Arguments

no.bin Number of binary variates.
prop.vec.bin Probability vector for binary variables

Examples

## Not run: validation.bin (3, rep(0.6,4))
validation.bin (4, rep(0.6,4))

validation.corr

Validates the specified correlation matrix

Description

This function validates the correlation vector and/or matrix for appropriate dimension, symmetry, range, and positive definiteness. If both correlation matrix and correlation vector were supplied, it checks whether the matrix and vector are conformable.

Usage

validation.corr(no.bin, no.nor, prop.vec.bin = NULL,
corr.vec = NULL, corr.mat = NULL)

Arguments

no.bin Number of binary variables
no.nor Number of normal variables
prop.vec.bin Probability vector for binary variables
corr.vec Vector of elements below the diagonal of correlation matrix ordered columnwise
corr.mat Specified correlation matrix
See Also

validation.bin, validation.range

Examples

d=4
corr.vec=c(0.21, 0.61, 0.78, 0.10, 0.12, 0.65)
corr.mat=lower.tri.to.corr.mat(corr.vec, d)

validation.corr (no.bin=2, no.nor=2, prop.vec.bin=c(0.4, 0.7),
corr.vec, corr.mat=corr.mat)

validation.nor

Validates the marginal specification of the normal part

Description

This function checks whether mean and variance parameters for the normal part are valid and consistent.

Usage

validation.nor(no.nor, mean.vec.nor = NULL, var.nor = NULL)

Arguments

no.nor Number of normal variables
mean.vec.nor Vector of means for normal variables
var.nor Vector of variances for normal variables

validation.range

Checks if the correlation terms are within the feasible range

Description

This function checks if there are correlation range violations among binary-binary, binary-normal and normal-normal combinations.

Usage

validation.range(no.bin, no.nor, prop.vec.bin = NULL, corr.mat)
validation.range

Arguments

- no.bin: Number of binary variables
- no.nor: Number of normal variables
- prop.vec.bin: Probability vector for binary variables
- corr.mat: Specified correlation matrix

Examples

```r
cmat = lower.tri.to.corr.mat(corr.vec = c(0.16, 0.04, 0.38, 0.4, 0.47, 0.68), 4)
validation.range(no.bin = 2, no.nor = 2, prop.vec.bin = c(0.4, 0.7), corr.mat = cmat)
```
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