Package ‘SimCorMultRes’

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Type Package

Title Simulates Correlated Multinomial Responses

Description
Simulates correlated multinomial responses conditional on a marginal model specification.

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Imports evd, methods, stats

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R.rsp, testthat

URL https://github.com/AnestisTouloumis/SimCorMultRes

BugReports https://github.com/AnestisTouloumis/SimCorMultRes/issues

License GPL-3

VignetteBuilder knitr, R.rsp

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Contents

`SimCorMultRes-package` *Simulating Correlated Multinomial Responses*

Description

Functions to simulate correlated multinomial responses (three or more nominal or ordinal response categories) and correlated binary responses subject to a marginal model specification.

Details

The simulated correlated binary or multinomial responses are drawn as realizations of a latent regression model for continuous random vectors with the correlation structure expressed in terms of the latent correlation.

For an ordinal response scale, the multinomial variables are simulated conditional on a marginal cumulative link model (`rmult.clm`), a marginal continuation-ratio model (`rmult.crm`) or a marginal adjacent-category logit model (`rmult.acl`).

For a nominal response scale, the multinomial responses are simulated conditional on a marginal baseline-category logit model (`rmult.bcl`).

Correlated binary responses are simulated using the function `rbin`.

The threshold approaches that give rise to the implemented marginal models are fully described in *Touloumis (2016)* and in the Vignette.

The formulae are easier to read from either the Vignette or the Reference Manual (both available here).

Author(s)

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References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

Emrich, L. J. and Piedmonte, M. R. (1991) A method for generating high-dimensional multivariate binary variates. *The American Statistician* **45**, 302–304.

Li, S. T. and Hammond, J. L. (1975) Generation of pseudorandom numbers with specified univariate distributions and correlation coefficients. *IEEE Transactions on Systems, Man and Cybernetics* **5**, 557–561.

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McFadden, D. (1974) *Conditional logit analysis of qualitative choice behavior*. New York: Academic Press, 105–142.

Touloumis, A. (2016) Simulating Correlated Binary and Multinomial Responses under Marginal Model Specification: The SimCorMultRes Package. *The R Journal* **8**, 79–91.

Touloumis, A., Agresti, A. and Kateri, M. (2013) GEE for multinomial responses using a local odds ratios parameterization. *Biometrics* **69**, 633–640.

Tutz, G. (1991) Sequential models in categorical regression. *Computational Statistics & Data Analysis* **11**, 275–295.

`rbin` *Simulating Correlated Binary Responses Conditional on a Marginal Model Specification*

Description

Simulates correlated binary responses assuming a regression model for the marginal probabilities.

Usage

```
rbin(clsize = clsize, intercepts = intercepts, betas = betas,
  xformula = formula(xdata), xdata = parent.frame(), link = "logit",
  cor.matrix = cor.matrix, rlatent = NULL)
```

Arguments

`clsize`	integer indicating the common cluster size.
`intercepts`	numerical (or numeric vector of length `clsize`) containing the intercept(s) of the marginal model.
`betas`	numerical vector or matrix containing the value of the marginal regression parameter vector associated with the covariates (i.e., excluding `intercepts`).
`xformula`	formula expression as in other marginal regression models but without including a response variable.
`xdata`	optional data frame containing the variables provided in `xformula`.
`link`	character string indicating the link function in the marginal model. Options include `'probit'`, `'logit'`, `'cloglog'`, `'cauchit'` or `'identity'`. Required when `rlatent = NULL`.
`cor.matrix`	matrix indicating the correlation matrix of the multivariate normal distribution when the NORTA method is employed (`rlatent = NULL`).
`rlatent`	matrix with `clsize` columns containing realizations of the latent random vectors when the NORTA method is not preferred. See details for more info.

Details

The formulae are easier to read from either the Vignette or the Reference Manual (both available here).

The assumed marginal model is

$$Pr(Y_{it} = 1|x_{it}) = F(\beta_{t0} + \beta_t^{'} x_{it})$$

where F is the cumulative distribution function determined by `link`. For subject i, Y_{it} is the t-th binary response and x_{it} is the associated covariates vector. Finally, β_{t0} and β_t are the intercept and regression parameter vector at the t-th measurement occasion.

The binary response Y_{it} is obtained by extending the approach of *Emrich and Piedmonte (1991)* as suggested in *Touloumis (2016).*

When $\beta_{t0} = \beta_0$ for all t, then `intercepts` should be provided as a single number. Otherwise, `intercepts` must be provided as a numeric vector such that the t-th element corresponds to the intercept at measurement occasion t.

`betas` should be provided as a numeric vector only when $\beta_t = \beta$ for all t. Otherwise, `betas` must be provided as a numeric matrix with `clsize` rows such that the t-th row contains the value of β_t. In either case, `betas` should reflect the order of the terms implied by `xformula`.

The appropriate use of `xformula` is `xformula = ~ covariates`, where `covariates` indicate the linear predictor as in other marginal regression models.

The optional argument `xdata` should be provided in "long" format.

The NORTA method is the default option for simulating the latent random vectors denoted by e_{it}^B in *Touloumis (2016).* To import simulated values for the latent random vectors without utilizing the NORTA method, the user can employ the `rlatent` argument. In this case, element (i, t) of `rlatent` represents the realization of e_{it}^B.

Value

Returns a list that has components:

`Ysim`	the simulated binary responses. Element (i,t) represents the realization of Y_{it}.
`simdata`	a data frame that includes the simulated response variables (y), the covariates specified by `xformula`, subjects' identities (id) and the corresponding measurement occasions (time).
`rlatent`	the latent random variables denoted by e_{it}^B in *Touloumis (2016).*

Author(s)

Anestis Touloumis

References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

Emrich, L. J. and Piedmonte, M. R. (1991) A method for generating high-dimensional multivariate binary variates. *The American Statistician* **45**, 302–304.

Li, S. T. and Hammond, J. L. (1975) Generation of pseudorandom numbers with specified univariate distributions and correlation coefficients. *IEEE Transactions on Systems, Man and Cybernetics* **5**, 557–561.

Touloumis, A. (2016) Simulating Correlated Binary and Multinomial Responses under Marginal Model Specification: The SimCorMultRes Package. *The R Journal* **8**, 79–91.

See Also

`rmult.bcl` for simulating correlated nominal responses, `rmult.clm`, `rmult.crm` and `rmult.acl` for simulating correlated ordinal responses.

Examples

```
## See Example 3.5 in the Vignette.
set.seed(123)
sample_size <- 5000
cluster_size <- 4
beta_intercepts <- 0
beta_coefficients <- 0.2
latent_correlation_matrix <- toeplitz(c(1, 0.9, 0.9, 0.9))
x <- rep(rnorm(sample_size), each = cluster_size)
simulated_binary_dataset <- rbin(clsize = cluster_size,
  intercepts = beta_intercepts, betas = beta_coefficients,
  xformula = ~x, cor.matrix = latent_correlation_matrix, link = "probit")
library(gee)
binary_gee_model <- gee(y ~ x, family = binomial("probit"), id = id,
  data = simulated_binary_dataset$simdata)
summary(binary_gee_model)$coefficients

## See Example 3.6 in the Vignette.
set.seed(8)
library(evd)
simulated_latent_variables1 <- rmvevd(sample_size, dep = sqrt(1 - 0.9),
  model = "log", d = cluster_size)
  simulated_latent_variables2 <- rmvevd(sample_size, dep = sqrt(1 - 0.9),
  model = "log", d = cluster_size)
simulated_latent_variables <- simulated_latent_variables1 -
  simulated_latent_variables2
simulated_binary_dataset <- rbin(clsize = cluster_size,
  intercepts = beta_intercepts, betas = beta_coefficients,
  xformula = ~x, rlatent = simulated_latent_variables)
binary_gee_model <- gee(y ~ x, family = binomial("logit"), id = id,
  data = simulated_binary_dataset$simdata)
summary(binary_gee_model)$coefficients
```

`rmult.acl` *Simulating Correlated Ordinal Responses Conditional on a Marginal Adjacent-Category Logit Model Specification*

Description

Simulates correlated ordinal responses assuming an adjacent-category logit model for the marginal probabilities.

Usage

```
rmult.acl(clsize = clsize, intercepts = intercepts, betas = betas,
  xformula = formula(xdata), xdata = parent.frame(),
  cor.matrix = cor.matrix, rlatent = NULL)
```

Arguments

`clsize`	integer indicating the common cluster size.
`intercepts`	numerical vector or matrix containing the intercepts of the marginal adjacent-category logit model.
`betas`	numerical vector or matrix containing the value of the marginal regression parameter vector.
`xformula`	formula expression as in other marginal regression models but without including a response variable.
`xdata`	optional data frame containing the variables provided in `xformula`.
`cor.matrix`	matrix indicating the correlation matrix of the multivariate normal distribution when the NORTA method is employed (`rlatent = NULL`).
`rlatent`	matrix with `(clsize * ncategories)` columns containing realizations of the latent random vectors when the NORTA method is not preferred. See details for more info.

Details

The formulae are easier to read from either the Vignette or the Reference Manual (both available here).

The assumed marginal adjacent-category logit model is

$$log \frac{Pr(Y_{it} = j | x_{it})}{Pr(Y_{it} = j + 1 | x_{it})} = \beta_{tj0} + \beta_t^{'} x_{it}$$

For subject i, Y_{it} is the t-th ordinal response and x_{it} is the associated covariates vector. Also β_{tj0} is the j-th category-specific intercept at the t-th measurement occasion and β_t is the regression parameter vector at the t-th measurement occasion.

The ordinal response Y_{it} is obtained by utilizing the threshold approach described in the Vignette. This approach is based on the connection between baseline-category and adjacent-category logit models.

When $\beta_{tj0} = \beta_{j0}$ for all t, then `intercepts` should be provided as a numerical vector. Otherwise, `intercepts` must be a numerical matrix such that row t contains the category-specific intercepts at the t-th measurement occasion.

`betas` should be provided as a numeric vector only when $\beta_t = \beta$ for all t. Otherwise, `betas` must be provided as a numeric matrix with `clsize` rows such that the t-th row contains the value of β_t. In either case, `betas` should reflect the order of the terms implied by `xformula`.

The appropriate use of xformula is xformula = ~ covariates, where covariates indicate the linear predictor as in other marginal regression models.

The optional argument xdata should be provided in "long" format.

The NORTA method is the default option for simulating the latent random vectors denoted by e^{O3}_{itj} in the Vignette. To import simulated values for the latent random vectors without utilizing the NORTA method, the user can employ the rlatent argument. In this case, row i corresponds to subject i and columns $(t-1)*\texttt{ncategories}+1, ..., t*\texttt{ncategories}$ should contain the realization of $e^{O3}_{it1}, ..., e^{O3}_{itJ}$, respectively, for $t = 1, \ldots, \texttt{clsize}$.

Value

Returns a list that has components:

- Ysim: the simulated nominal responses. Element (i,t) represents the realization of Y_{it}.
- simdata: a data frame that includes the simulated response variables (y), the covariates specified by xformula, subjects' identities (id) and the corresponding measurement occasions (time).
- rlatent: the latent random variables denoted by e^{O3}_{itj} in the Vignette.

Author(s)

Anestis Touloumis

References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

Li, S. T. and Hammond, J. L. (1975) Generation of pseudorandom numbers with specified univariate distributions and correlation coefficients. *IEEE Transactions on Systems, Man and Cybernetics* **5**, 557–561.

Touloumis, A. (2016) Simulating Correlated Binary and Multinomial Responses under Marginal Model Specification: The SimCorMultRes Package. *The R Journal* **8**, 79–91.

Touloumis, A., Agresti, A. and Kateri, M. (2013) GEE for multinomial responses using a local odds ratios parameterization. *Biometrics* **69**, 633–640.

See Also

rbin for simulating correlated binary responses, rmult.clm and rmult.crm for simulating correlated ordinal responses, and rmult.bcl for simulating nominal responses.

Examples

```
## See Example 3.4 in the Vignette.
beta_intercepts <- c(3, 1, 2)
beta_coefficients <- c(1, 1)
sample_size <- 500
cluster_size <- 3
```

```
set.seed(321)
x1 <- rep(rnorm(sample_size), each = cluster_size)
x2 <- rnorm(sample_size * cluster_size)
xdata <- data.frame(x1, x2)
identity_matrix <- diag(4)
equicorrelation_matrix <- toeplitz(c(1, rep(0.95, cluster_size - 1)))
latent_correlation_matrix <- kronecker(equicorrelation_matrix,
  identity_matrix)
simulated_ordinal_dataset <- rmult.acl(clsize = cluster_size,
  intercepts = beta_intercepts, betas = beta_coefficients,
  xformula = ~ x1 + x2, xdata = xdata,
  cor.matrix = latent_correlation_matrix)
suppressPackageStartupMessages(library("multgee"))
ordinal_gee_model <- ordLORgee(y ~ x1 + x2,
  data = simulated_ordinal_dataset$simdata, id = id, repeated = time,
  LORstr = "time.exch", link = "acl")
round(coef(ordinal_gee_model), 2)
```

`rmult.bcl` *Simulating Correlated Nominal Responses Conditional on a Marginal Baseline-Category Logit Model Specification*

Description

Simulates correlated nominal responses assuming a baseline-category logit model for the marginal probabilities.

Usage

```
rmult.bcl(clsize = clsize, ncategories = ncategories, betas = betas,
  xformula = formula(xdata), xdata = parent.frame(),
  cor.matrix = cor.matrix, rlatent = NULL)
```

Arguments

`clsize`	integer indicating the common cluster size.
`ncategories`	integer indicating the number of nominal response categories.
`betas`	numerical vector or matrix containing the value of the marginal regression parameter vector.
`xformula`	formula expression as in other marginal regression models but without including a response variable.
`xdata`	optional data frame containing the variables provided in `xformula`.
`cor.matrix`	matrix indicating the correlation matrix of the multivariate normal distribution when the NORTA method is employed (`rlatent = NULL`).
`rlatent`	matrix with `(clsize * ncategories)` columns containing realizations of the latent random vectors when the NORTA method is not preferred. See details for more info.

Details

The formulae are easier to read from either the Vignette or the Reference Manual (both available here).

The assumed marginal baseline category logit model is

$$log \frac{Pr(Y_{it} = j|x_{it})}{Pr(Y_{it} = J|x_{it})} = (\beta_{tj0} - \beta_{tJ0}) + (\beta^{'}_{tj} - \beta^{'}_{tJ})x_{it} = \beta^{*}_{tj0} + \beta^{*'}_{tj} x_{it}$$

For subject i, Y_{it} is the t-th nominal response and x_{it} is the associated covariates vector. Also β_{tj0} is the j-th category-specific intercept at the t-th measurement occasion and β_{tj} is the j-th category-specific regression parameter vector at the t-th measurement occasion.

The nominal response Y_{it} is obtained by extending the principle of maximum random utility (*McFadden, 1974*) as suggested in *Touloumis (2016)*.

`betas` should be provided as a numeric vector only when $\beta_{tj0} = \beta_{j0}$ and $\beta_{tj} = \beta_j$ for all t. Otherwise, `betas` must be provided as a numeric matrix with `clsize` rows such that the t-th row contains the value of $(\beta_{t10}, \beta_{t1}, \beta_{t20}, \beta_{t2}, ..., \beta_{tJ0}, \beta_{tJ})$. In either case, `betas` should reflect the order of the terms implied by `xformula`.

The appropriate use of `xformula` is `xformula = ~ covariates`, where `covariates` indicate the linear predictor as in other marginal regression models.

The optional argument `xdata` should be provided in "long" format.

The NORTA method is the default option for simulating the latent random vectors denoted by e^{NO}_{itj} in *Touloumis (2016)*. In this case, the algorithm forces `cor.matrix` to respect the assumption of choice independence. To import simulated values for the latent random vectors without utilizing the NORTA method, the user can employ the `rlatent` argument. In this case, row i corresponds to subject i and columns $(t-1) * \texttt{ncategories} + 1, ..., t * \texttt{ncategories}$ should contain the realization of $e^{NO}_{it1}, ..., e^{NO}_{itJ}$, respectively, for $t = 1, \ldots, \texttt{clsize}$.

Value

Returns a list that has components:

`Ysim`	the simulated nominal responses. Element (i,t) represents the realization of Y_{it}.
`simdata`	a data frame that includes the simulated response variables (y), the covariates specified by `xformula`, subjects' identities (id) and the corresponding measurement occasions (time).
`rlatent`	the latent random variables denoted by e^{NO}_{it} in *Touloumis (2016)*.

Author(s)

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References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

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Touloumis, A., Agresti, A. and Kateri, M. (2013) GEE for multinomial responses using a local odds ratios parameterization. *Biometrics* **69**, 633–640.

See Also

`rbin` for simulating correlated binary responses, `rmult.clm`, `rmult.crm` and `rmult.acl` for simulating correlated ordinal responses.

Examples

```
## See Example 3.1 in the Vignette.
betas <- c(1, 3, 2, 1.25, 3.25, 1.75, 0.75, 2.75, 2.25, 0, 0, 0)
sample_size <- 500
categories_no <- 4
cluster_size <- 3
set.seed(1)
x1 <- rep(rnorm(sample_size), each = cluster_size)
x2 <- rnorm(sample_size * cluster_size)
xdata <- data.frame(x1, x2)
equicorrelation_matrix <- toeplitz(c(1, rep(0.95, cluster_size - 1)))
identity_matrix <- diag(categories_no)
latent_correlation_matrix <- kronecker(equicorrelation_matrix,
  identity_matrix)
simulated_nominal_dataset <- rmult.bcl(clsize = cluster_size,
  ncategories = categories_no, betas = betas, xformula = ~ x1 + x2,
  xdata = xdata, cor.matrix = latent_correlation_matrix)
suppressPackageStartupMessages(library("multgee"))
nominal_gee_model <- nomLORgee(y ~ x1 + x2,
  data = simulated_nominal_dataset$simdata, id = id, repeated = time,
  LORstr = "time.exch")
round(coef(nominal_gee_model), 2)
```

`rmult.clm` *Simulating Correlated Ordinal Responses Conditional on a Marginal Cumulative Link Model Specification*

Description

Simulates correlated ordinal responses assuming a cumulative link model for the marginal probabilities.

Usage

```
rmult.clm(clsize = clsize, intercepts = intercepts, betas = betas,
  xformula = formula(xdata), xdata = parent.frame(), link = "logit",
  cor.matrix = cor.matrix, rlatent = NULL)
```

Arguments

`clsize`	integer indicating the common cluster size.
`intercepts`	numerical vector or matrix containing the intercepts of the marginal cumulative link model.
`betas`	numerical vector or matrix containing the value of the marginal regression parameter vector associated with the covariates (i.e., excluding `intercepts`).
`xformula`	formula expression as in other marginal regression models but without including a response variable.
`xdata`	optional data frame containing the variables provided in `xformula`.
`link`	character string indicating the link function in the marginal cumulative link model. Options include `'probit'`, `'logit'`, `'cloglog'` or `'cauchit'`. Required when `rlatent = NULL`.
`cor.matrix`	matrix indicating the correlation matrix of the multivariate normal distribution when the NORTA method is employed (`rlatent = NULL`).
`rlatent`	matrix with `clsize` columns containing realizations of the latent random vectors when the NORTA method is not preferred. See details for more info.

Details

The formulae are easier to read from either the Vignette or the Reference Manual (both available here).

The assumed marginal cumulative link model is

$$Pr(Y_{it} \leq j|x_{it}) = F(\beta_{tj0} + \beta_t^{'} x_{it})$$

where F is the cumulative distribution function determined by `link`. For subject i, Y_{it} is the t-th ordinal response and x_{it} is the associated covariates vector. Finally, β_{tj0} is the j-th category-specific intercept at the t-th measurement occasion and β_{tj} is the j-th category-specific regression parameter vector at the t-th measurement occasion.

The ordinal response Y_{it} is obtained by extending the approach of *McCullagh (1980)* as suggested in *Touloumis (2016)*.

When $\beta_{tj0} = \beta_{j0}$ for all t, then `intercepts` should be provided as a numerical vector. Otherwise, `intercepts` must be a numerical matrix such that row t contains the category-specific intercepts at the t-th measurement occasion.

`betas` should be provided as a numeric vector only when $\beta_t = \beta$ for all t. Otherwise, `betas` must be provided as a numeric matrix with `clsize` rows such that the t-th row contains the value of β_t. In either case, `betas` should reflect the order of the terms implied by `xformula`.

The appropriate use of `xformula` is `xformula = ~ covariates`, where `covariates` indicate the linear predictor as in other marginal regression models.

The optional argument xdata should be provided in "long" format.

The NORTA method is the default option for simulating the latent random vectors denoted by e^{O1}_{it} in *Touloumis (2016)*. To import simulated values for the latent random vectors without utilizing the NORTA method, the user can employ the rlatent argument. In this case, element (i, t) of rlatent represents the realization of e^{O1}_{it}.

Value

Returns a list that has components:

- Ysim — the simulated ordinal responses. Element (i,t) represents the realization of Y_{it}.
- simdata — a data frame that includes the simulated response variables (y), the covariates specified by xformula, subjects' identities (id) and the corresponding measurement occasions (time).
- rlatent — the latent random variables denoted by e^{O1}_{it} in *Touloumis (2016)*.

Author(s)

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References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

Li, S. T. and Hammond, J. L. (1975) Generation of pseudorandom numbers with specified univariate distributions and correlation coefficients. *IEEE Transactions on Systems, Man and Cybernetics* **5**, 557–561.

McCullagh, P. (1980) Regression models for ordinal data. *Journal of the Royal Statistical Society B* **42**, 109–142.

Touloumis, A. (2016) Simulating Correlated Binary and Multinomial Responses under Marginal Model Specification: The SimCorMultRes Package. *The R Journal* **8**, 79–91.

Touloumis, A., Agresti, A. and Kateri, M. (2013) GEE for multinomial responses using a local odds ratios parameterization. *Biometrics* **69**, 633–640.

See Also

rmult.bcl for simulating correlated nominal responses, rmult.crm and rmult.acl for simulating correlated ordinal responses and rbin for simulating correlated binary responses.

Examples

```
## See Example 3.2 in the Vignette.
set.seed(12345)
sample_size <- 500
cluster_size <- 4
beta_intercepts <- c(-1.5, -0.5, 0.5, 1.5)
beta_coefficients <- matrix(c(1, 2, 3, 4), 4, 1)
```

```
x <- rep(rnorm(sample_size), each = cluster_size)
latent_correlation_matrix <- toeplitz(c(1, 0.85, 0.5, 0.15))
simulated_ordinal_dataset <- rmult.clm(clsize = cluster_size,
  intercepts = beta_intercepts, betas = beta_coefficients, xformula = ~x,
  cor.matrix = latent_correlation_matrix, link = "probit")
head(simulated_ordinal_dataset$simdata, n = 8)

## Same sampling scheme except that the parameter vector is time-stationary.
set.seed(12345)
simulated_ordinal_dataset <- rmult.clm(clsize = cluster_size, betas = 1,
  xformula = ~x, cor.matrix = latent_correlation_matrix,
  intercepts = beta_intercepts, link = "probit")
## Fit a GEE model (Touloumis et al., 2013) to estimate the regression
## coefficients.
library(multgee)
ordinal_gee_model <- ordLORgee(y ~ x, id = id, repeated = time,
  link = "probit", data = simulated_ordinal_dataset$simdata)
coef(ordinal_gee_model)
```

`rmult.crm` *Simulating Correlated Ordinal Responses Conditional on a Marginal Continuation-Ratio Model Specification*

Description

Simulates correlated ordinal responses assuming a continuation-ratio model for the marginal probabilities.

Usage

```
rmult.crm(clsize = clsize, intercepts = intercepts, betas = betas,
  xformula = formula(xdata), xdata = parent.frame(), link = "logit",
  cor.matrix = cor.matrix, rlatent = NULL)
```

Arguments

`clsize`	integer indicating the common cluster size.
`intercepts`	numerical vector or matrix containing the intercepts of the marginal continuation-ratio model.
`betas`	numerical vector or matrix containing the value of the marginal regression parameter vector associated with the covariates (i.e., excluding `intercepts`).
`xformula`	formula expression as in other marginal regression models but without including a response variable.
`xdata`	optional data frame containing the variables provided in `xformula`.
`link`	character string indicating the link function of the marginal continuation-ratio model. Options include `'probit'`, `'logit'`, `'cloglog'` or `'cauchit'`. Required when `rlatent = NULL`.

`cor.matrix` matrix indicating the correlation matrix of the multivariate normal distribution when the NORTA method is employed (`rlatent = NULL`).

`rlatent` matrix with `clsize` rows and `ncategories` columns containing realizations of the latent random vectors when the NORTA method is not employed. See details for more info.

Details

The formulae are easier to read from either the Vignette or the Reference Manual (both available here).

The assumed marginal continuation-ratio model is

$$Pr(Y_{it} = j | Y_{it} \geq j, x_{it}) = F(\beta_{tj0} + \beta_t^{'} x_{it})$$

where F is the cumulative distribution function determined by `link`. For subject i, Y_{it} is the t-th multinomial response and x_{it} is the associated covariates vector. Finally, β_{tj0} is the j-th category-specific intercept at the t-th measurement occasion and β_{tj} is the j-th category-specific regression parameter vector at the t-th measurement occasion.

The ordinal response Y_{it} is determined by extending the latent variable threshold approach of *Tutz (1991)* as suggested in *Touloumis (2016)*.

When $\beta_{tj0} = \beta_{j0}$ for all t, then `intercepts` should be provided as a numerical vector. Otherwise, `intercepts` must be a numerical matrix such that row t contains the category-specific intercepts at the t-th measurement occasion.

`betas` should be provided as a numeric vector only when $\beta_t = \beta$ for all t. Otherwise, `betas` must be provided as a numeric matrix with `clsize` rows such that the t-th row contains the value of β_t. In either case, `betas` should reflect the order of the terms implied by `xformula`.

The appropriate use of `xformula` is `xformula = ~ covariates`, where `covariates` indicate the linear predictor as in other marginal regression models.

The optional argument `xdata` should be provided in "long" format.

The NORTA method is the default option for simulating the latent random vectors denoted by e^{O2}_{itj} in *Touloumis (2016)*. In this case, the algorithm forces `cor.matrix` to respect the local independence assumption. To import simulated values for the latent random vectors without utilizing the NORTA method, the user can employ the `rlatent` argument. In this case, row i corresponds to subject i and columns $(t-1) * \texttt{ncategories} + 1, ..., t * \texttt{ncategories}$ should contain the realization of $e^{O2}_{it1}, ..., e^{O2}_{itJ}$, respectively, for $t = 1, \ldots, \texttt{clsize}$.

Value

Returns a list that has components:

`Ysim` the simulated ordinal responses. Element (i,t) represents the realization of Y_{it}.

`simdata` a data frame that includes the simulated response variables (y), the covariates specified by `xformula`, subjects' identities (id) and the corresponding measurement occasions (time).

`rlatent` the latent random variables denoted by e^{O2}_{it} in *Touloumis (2016)*.

Author(s)

Anestis Touloumis

References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

Li, S. T. and Hammond, J. L. (1975) Generation of pseudorandom numbers with specified univariate distributions and correlation coefficients. *IEEE Transactions on Systems, Man and Cybernetics* **5**, 557–561.

Touloumis, A. (2016) Simulating Correlated Binary and Multinomial Responses under Marginal Model Specification: The SimCorMultRes Package. *The R Journal (forthcoming)*.

Tutz, G. (1991) Sequential models in categorical regression. *Computational Statistics & Data Analysis* **11**, 275–295.

See Also

`rmult.bcl` for simulating correlated nominal responses, `rmult.clm` and `rmult.acl` for simulating correlated ordinal responses and `rbin` for simulating correlated binary responses.

Examples

```
## See Example 3.3 in the Vignette.
set.seed(1)
sample_size <- 500
cluster_size <- 4
beta_intercepts <- c(-1.5, -0.5, 0.5, 1.5)
beta_coefficients <- 1
x <- rnorm(sample_size * cluster_size)
categories_no <- 5
identity_matrix <- diag(1, (categories_no - 1) * cluster_size)
equicorrelation_matrix <- toeplitz(c(0, rep(0.24, categories_no - 2)))
ones_matrix <- matrix(1, cluster_size, cluster_size)
latent_correlation_matrix <- identity_matrix +
  kronecker(equicorrelation_matrix, ones_matrix)
simulated_ordinal_dataset <- rmult.crm(clsize = cluster_size,
  intercepts = beta_intercepts, betas = beta_coefficients, xformula = ~x,
  cor.matrix = latent_correlation_matrix, link = "probit")
head(simulated_ordinal_dataset$Ysim)
```

`rnorta` *Simulating Random Vectors using the NORTA Method*

Description

Utility function to simulate random vectors with predefined marginal distributions via the NORTA method.

Usage

```
rnorta(R = R, cor.matrix = cor.matrix, distr = distr,
  qparameters = NULL)
```

Arguments

R	integer indicating the sample size.
cor.matrix	matrix indicating the correlation matrix of the multivariate normal distribution employed in the NORTA method.
distr	character string vector of length ncol(cor.matrix) naming the quantile functions of the desired marginal distributions.
qparameters	list of ncol(cor.matrix) lists indicating the parameter values of the quantile functions specified by distr.

Details

Checks are made to ensure that cor.matrix is a positive definite correlation matrix. The positive definiteness of cor.matrix is assessed via eigenvalues.

The t-th character string in distr indicates the quantile function of the t-th marginal distribution. See Distributions for the most common distributions. Quantile functions supported by other R packages are allowed provided that these packages have been uploaded first. However, note that no checks are made to ensure that the character strings in distr correspond to valid names of quantile functions.

If qparameters = NULL then the default parameter values for the quantile functions specified by distr are used. Otherwise, qparameters should be provided as a list of ncol(cor.matrix) lists such that the t-th list contains the desired parameter values of the t-th quantile function.

Value

Returns R random vectors of size ncol(cor.matrix) with marginal distributions specified by distr (and qparameters).

Author(s)

Anestis Touloumis

References

Cario, M. C. and Nelson, B. L. (1997) *Modeling and generating random vectors with arbitrary marginal distributions and correlation matrix*. Technical Report, Department of Industrial Engineering and Management Sciences, Northwestern University, Evanston, Illinois.

Li, S. T. and Hammond, J. L. (1975) Generation of pseudorandom numbers with specified univariate distributions and correlation coefficients. *IEEE Transactions on Systems, Man and Cybernetics* **5**, 557–561.

Touloumis, A. (2016) Simulating Correlated Binary and Multinomial Responses under Marginal Model Specification: The SimCorMultRes Package. *The R Journal* **8**, 79–91.

Examples

```
## An example with standard logistic as marginal distribution.
set.seed(1)
sample_size <- 1000
latent_correlation_matrix <- toeplitz(c(1, rep(0.8, 2)))
latent_correlation_matrix
common_marginal_distribution <- rep("qlogis", 3)
simulated_logistic_responses <- rnorta(R = sample_size,
  cor.matrix = latent_correlation_matrix,
  distr = common_marginal_distribution)

## The following lines exemplify the NORTA method.
set.seed(1)
simulated_normal_responses <- rsmvnorm(R = sample_size,
  cor.matrix = latent_correlation_matrix)
norta_simulated <- qlogis(pnorm(simulated_normal_responses))
all(simulated_logistic_responses == norta_simulated)

## Change the marginal distributions to standard normal, standard
## logistic and standard extreme value distribution.
set.seed(1)
different_marginal_distributions <- c("qnorm", "qlogis", "qgumbel")
simulated_logistic_responses <- rnorta(R = sample_size,
  cor.matrix = latent_correlation_matrix,
  distr = different_marginal_distributions)
cor(simulated_logistic_responses)
colMeans(simulated_logistic_responses)
apply(simulated_logistic_responses, 2, sd)

## Same as above but using parameter values other than the default ones.
set.seed(1)
qpars <- list(c(mean = 1, sd = 9), c(location = 2, scale = 1),
  c(loc = 3, scale = 1))
simulated_logistic_responses <- rnorta(R = sample_size,
  cor.matrix = latent_correlation_matrix,
  distr = different_marginal_distributions, qparameters = qpars)
cor(simulated_logistic_responses)
colMeans(simulated_logistic_responses)
apply(simulated_logistic_responses, 2, sd)
```

---

rsmvnorm — *Simulating Continuous Random Vectors from a Multivariate Normal Distribution*

Description

Utility function to simulate continuous random vectors from a multivariate normal distribution such that all marginal distributions are univariate standard normal.

Usage

```
rsmvnorm(R = R, cor.matrix = cor.matrix)
```

Arguments

R	integer indicating the sample size.
cor.matrix	matrix indicating the correlation matrix of the multivariate normal distribution.

Details

Checks are made to ensure that `cor.matrix` is a positive definite correlation matrix. The positive definiteness of `cor.matrix` is assessed via eigenvalues.

Value

Returns `R` random vectors of size `ncol(cor.matrix)`.

Author(s)

Anestis Touloumis

Examples

```
## Simulating 10000 bivariate random vectors with correlation parameter
## equal to 0.4.
set.seed(1)
sample_size <- 10000
correlation_matrix <- toeplitz(c(1, 0.4))
simulated_normal_responses <- rsmvnorm(R = sample_size,
  cor.matrix = correlation_matrix)
colMeans(simulated_normal_responses)
apply(simulated_normal_responses, 2, sd)
cor(simulated_normal_responses)
```

`simulation` *Simulated Correlation Parameters*

Description

A simulated dataset to explore the association between the correlation parameter of bivariate normally distributed random variables used in the intermediate step of the NORTA method and the correlation parameter of the resulting non-normal random responses generated by the NORTA method for all the threshold approached implemented in this package.

Usage

```
simulation
```

Format

A data frame with 100 rows and 4 columns:

rho numeric indicating the true value of the correlation parameter.

normal numeric indicating the simulated correlation parameter when the marginal distribution of each of the latent variables is normal.

logistic numeric indicating the simulated correlation parameter when the marginal distribution of each of the latent variables is logistic.

gumbel numeric indicating the simulated correlation parameter when the marginal distribution of each of the latent variables is Gumbel.

Examples

```
plot(rho - normal ~ rho, data = simulation, type = "l", col = "blue",
    ylim = c(0, 0.016),
    ylab = expression(rho - bar(rho)[sim]),
    xlab = expression(rho))
points(rho - logistic ~ rho, data = simulation, type = "l", col = "red")
points(rho - gumbel ~ rho, data = simulation, type = "l", col = "grey")
legend("topright", legend = c("Normal", "Logistic", "Gumbel"),
      col = c("blue", "red", "grey"), pch = "l" )
```

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