Package ‘reReg’

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Title  Recurrent Event Regression
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Description
A comprehensive collection of practical and easy-to-use tools for regression analysis of recurrent events, with or without the presence of a (possibly) informative terminal event. The modeling framework is based on a joint frailty scale-change model, that includes models described in Wang et al. (2001) <doi:10.1198/016214501753209031>, Huang and Wang (2004) <doi:10.1198/016214504000001033>, Xu et al. (2017) <doi:10.1080/01621459.2016.1173557>, and Xu et al. (2019) <doi:10.5705/SS.202018.0224> as special cases. The implemented estimating procedure does not require any parametric assumption on the frailty distribution. The package also allows the users to specify different model forms for both the recurrent event process and the terminal event.

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Author  Sy Han (Steven) Chiou [aut, cre],
         Chiung-Yu Huang [aut]
Maintainer  Sy Han (Steven) Chiou <schiou@utdallas.edu>
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reReg-package

reReg: Recurrent Event Regression

Description

The package offers a comprehensive collection of practical and easy-to-use tools for analyzing recurrent event data, with or without the presence of a (possibly) correlated terminal event. The modeling framework is based on a joint frailty scale-change model, that encompasses many existing models, including the popular Cox-type models, as special cases and accommodates informative censoring through a subject-specific frailty. The implemented estimating procedure does not require any parametric assumption on the frailty distribution. The package allows the users to specify different model forms for both the recurrent event process and the terminal event. The package also includes tools for visualization of recurrent events and simulation from the regression models.

Author(s)

Maintainer: Sy Han (Steven) Chiou <schiou@utdallas.edu>

Authors:

- Chiung-Yu Huang <ChiungYu.Huang@ucsf.edu>

References


See Also

Useful links:

- https://github.com/stc04003/reReg
- Report bugs at https://github.com/stc04003/reReg/issues

---

**basebind**

*Function used to combine baseline functions in one plot*

**Description**

Combine different plots into one.

**Usage**

```r
basebind(..., legend.title, legend.labels, control = list())
```

**Arguments**

- `...` ggplot objects created by plotting reReg objects.
- `legend.title` an optional character string to specify the legend title.
- `legend.labels` an optional character string to specify the legend labels.
- `control` a list of control parameters.

**Examples**

```r
data(simDat)
fm <- Recur(t.stop, id, event, status) ~ x1 + x2
fit1 <- reReg(fm, subset = x1 == 0, data = simDat, B = 200)
fit2 <- reReg(fm, subset = x1 == 1, data = simDat, B = 200)
basebind(plot(fit1), plot(fit2))
```
**plot.Recur**

*Produce Event Plot or Mean Cumulative Function Plot*

**Description**

Plot the event plot or the mean cumulative function (MCF) from an Recur object.

**Usage**

```r
## S3 method for class 'Recur'
plot(
  x,
  mcf = FALSE,
  event.result = c("increasing", "decreasing", "asis"),
  event.calendarTime = FALSE,
  mcf.adjustRiskset = TRUE,
  mcf.conf.int = FALSE,
  control = list(),
  ...
)
```

**Arguments**

- `x`: an object of class Recur returned by the Recur() function. See ?Recur for creating Recur objects.
- `mcf`: an optional logical value indicating whether the mean cumulative function (MCF) will be plotted instead of the event plot. When `mcf = TRUE`, the `mcf` is internally called. See `mcf` for details.
- `event.result`: an optional character string that is passed to the plotEvents() function as the result argument. See plotEvents. This argument is used to specify whether the event plot is sorted by the subjects' terminal time. The available options are:
  - `increasing`: sort the terminal time from in ascending order (default). This places longer terminal times on top.
  - `decreasing`: sort the terminal time from in descending order. This places shorter terminal times on top.
  - `none`: present the event plots as is, without sorting by the terminal times.
- `event.calendarTime`: an optional logical value indicating whether to plot in calendar time. When `event.calendarTime = FALSE` (default), the event plot will have patient time on the x-axis.
- `mcf.adjustRiskset`: an optional logical value that is passed to the mcf() function as the adjustRiskset argument. This argument indicates whether risk set size will be adjusted. If `mcf.adjustRiskset = TRUE`, subjects leave the risk set after terminal times as in the Nelson-Aalen estimator. If `mcf.adjustRiskset = FALSE`, subjects remain in the risk set after terminal time.
mcf.conf.int  an optional logical value that is passed to the mcf() function as the conf.int argument. See mcf for details.
control      a list of control parameters. See Details.
...          additional graphical parameters to be passed to methods.

Details

The argument control consists of options with argument defaults to a list with the following values:
xlab customizable x-label, default value is "Time".
ylab customizable y-label, default value is "Subject" for event plot and "Cumulative mean" for MCF plot.
main customizable title, the default value is "Recurrent event plot" when mcf = FALSE and "Sample cumulative mean function plot" when mcf = TRUE.
terminal.name customizable label for terminal event, the default value is "Terminal event".
recurrent.name customizable legend title for recurrent event, the default value is "Recurrent events".
recurrent.types customizable label for recurrent event type, the default value is NULL.
alpha    between 0 and 1, controls the transparency of points.

The xlab, ylab and main parameters can be specified outside of the control list.

Value

A ggplot object.

References


See Also

Recur, plotEvents, mcf

Examples

data(simDat)
reObj <- with(simDat, Recur(t.start %to% t.stop, id, event, status))

## Event plots:
plot(reObj)
plot(reObj, event.result = "decreasing")

## With (hypothetical) multiple event types
simDat$event2 <- with(simDat, ifelse(t.stop > 10 & event > 0, 2, event))
reObj2 <- with(simDat, Recur(t.start %to% t.stop, id, event2, status))
plot(reObj2)

## With (hypothetical) calendar times
simDat2 <- simDat
simDat2$t.start <- as.Date(simDat2$t.start + simDat2$x2 * 5, origin = "20-01-01")
simDat2$t.stop <- as.Date(simDat2$t.stop + simDat2$x2 * 5, origin = "20-01-01")
reObj3 <- with(simDat2, Recur(t.start %to% t.stop, id, event, status))
plot(reObj3, event.calendarTime = TRUE)

## MCF plots
plot(reObj, mcf = TRUE)
plot(reObj, mcf = TRUE, mcf.adjustRiskset = FALSE)

library(reReg)
data(simDat)
reObj <- with(simDat, Recur(t.start %to% t.stop, id, event, status))
summary(reObj)

---

**plot.reReg**

Plot the Baseline Cumulative Rate Function and the Baseline Cumulative Hazard Function

### Description

Plot the baseline cumulative rate function and the baseline cumulative hazard function (if applicable) for an `reReg` object.

### Usage

```r
## S3 method for class 'reReg'
plot(
x, 
baseline = c("both", "rate", "hazard"),
smooth = FALSE,
newdata = NULL,
frailty = NULL,
showName = FALSE,
control = list(),
...
)
```

### Arguments

- **x**: an object of class `reReg`, returned by the `reReg` function.
- **baseline**: a character string specifying which baseline function to plot.
  - `baseline = "both"` plot both the baseline cumulative rate and the baseline cumulative hazard function (if applicable) in separate panels within the same display (default).
  - `baseline = "rate"` plot the baseline cumulative rate function.
baseline = "hazard" plot the baseline cumulative hazard function.

smooth an optional logical value indicating whether to add a smooth curve obtained from a monotone increasing P-splines implemented in package scam.

newdata an optional data frame contains variables to include in the calculation of the cumulative rate function. If omitted, the baseline rate function will be plotted.

frailty an optional vector to specify the shared frailty for newdata. If newdata is given and frailty is not specified, the

showName an optional logical value indicating whether to label the curves when newdata is specified.

control a list of control parameters. See Details.

... additional graphical parameters to be passed to methods.

Details

The argument control consists of options with argument defaults to a list with the following values:

xlab customizable x-label, default value is "Time".

ylab customizable y-label, default value is empty.

main customizable title, default value are "Baseline cumulative rate and hazard function" when baseline = "both", "Baseline cumulative rate function" when baseline = "rate", and "Baseline cumulative hazard function" when baseline = "hazard".

Value

A ggplot object.

See Also

reReg

Examples

data(simDat)
fm <- Recur(t.start %to% t.stop, id, event, status) ~ x1 + x2

fit <- reReg(fm, data = simDat, B = 0)
plot(fit)
plot(fit, xlab = "Time (days)", smooth = TRUE)

## Predicted cumulative rate and hazard given covariates
newdata <- expand.grid(x1 = 0:1, x2 = mean(simDat$x2))
plot(fit, newdata = newdata, showName = TRUE)
plotEvents 

Produce Event Plots

Description

Plot the event plot for an Recur object. The usage of the function is similar to that of plot.Recur() but with more flexible options.

Usage

plotEvents(
  formula,
  data,
  result = c("increasing", "decreasing", "asis"),
  calendarTime = FALSE,
  control = list(),
  ...
)

Arguments

 formula a formula object, with the response on the left of a "~" operator, and the predictors on the right. The response must be a recurrent event survival object as returned by function Recur().

 data an optional data frame in which to interpret the variables occurring in the "formula".

 result an optional character string specifying whether the event plot is sorted by the subjects’ terminal time. The available options are

 increasing sort the terminal time from in ascending order (default). This places longer terminal times on top.

decreasing sort the terminal time from in descending order. This places shorter terminal times on top.

 none present the event plots as is, without sorting by the terminal times.

 calendarTime an optional logical value indicating whether to plot in calendar time. When calendarTime = FALSE (default), the event plot will have patient time on the x-axis.

 control a list of control parameters. See Details.

 ... graphical parameters to be passed to methods. These include xlab, ylab, main, and more. See Details.

Details

The argument control consists of options with argument defaults to a list with the following values:

xlab customizable x-label, default value is "Time".
**ylab** customizable y-label, default value is "Subject" for event plot and "Cumulative mean" for MCF plot.

**main** customizable title, the default value is "Recurrent event plot" when `mcf = FALSE` and "Sample cumulative mean function plot" when `mcf = TRUE`.

**terminal.name** customizable label for terminal event, the default value is "Terminal event".

**recurrent.name** customizable legend title for recurrent event, the default value is "Recurrent events".

**recurrent.types** customizable label for recurrent event type, the default value is `NULL`.

**alpha** between 0 and 1, controls the transparency of points.

The `xlab`, `ylab` and `main` parameters can be specified outside of the `control` list.

### Value

A ggplot object.

### See Also

`Recur`, `plot.Recur`

### Examples

```r
data(simDat)
plotEvents(Recur(t.start %to% t.stop, id, event, status) ~ 1, data = simDat, 
    xlab = "Time in days", ylab = "Subjects arranged by terminal time")

## Separate plots by x1
plotEvents(Recur(t.start %to% t.stop, id, event, status) ~ x1, data = simDat)

## For multiple recurrent events
simDat$x3 <- ifelse(simDat$x2 < 0, "x2 < 0", "x2 > 0")
simDat$event <- simDat$event * sample(1:3, nrow(simDat), TRUE)
plotEvents(Recur(t.start %to% t.stop, id, event, status) ~ x1 + x3, data = simDat)
```

---

**Description**

This function provides the plotting options for the `plotEvents()` function.

**Usage**

```r
plotEvents.control(
    xlab = NULL,
    ylab = NULL,
    main = NULL,
    terminal.name = NULL,
```
plotEvents.control

recurrent.name = NULL,
recurrent.type = NULL,
legend.position = "top",
base_size = 12,
cex = NULL,
width = NULL,
bar.color = NULL,
recurrent.color = NULL,
recurrent.shape = NULL,
recurrent.stroke = NULL,
terminal.color = NULL,
terminal.shape = NULL,
terminal.stroke = NULL,
not.terminal.color = NULL,
not.terminal.shape = NULL,
aloha = 0.7
)

Arguments

xlab a character string indicating the label for the x axis. The default value is "Time".

ylab a character string indicating the label for the y axis. The default value is "Subject".

main a character string indicating the title of the plot.

terminal.name a character string indicating the label for the terminal event displayed in the legend. The default value is "Terminal event".

recurrent.name a character string indicating the label for the recurrent event displayed in the legend. The default value is "Recurrent events".

recurrent.type a factor indicating the labels for the different recurrent event types. This option is only available when there are more than one types of recurrent events. The default value is "Recurrent events 1", "Recurrent events 2", ....

legend.position a character string specifies the position of the legend. The available options are "none", "left", "right", "bottom", "top", or a two-element numeric vector specifies the coordinate of the legend. This argument is passed to the ggplot theme environment. The default value is "top".

base_size a numerical value to specify the base font size, given in pts. This argument is passed to the ggplot theme environment. The default value is 12.

cex a numerical value specifies the size of the points.

width a numerical value specifies the width of the event plot. By ggplot default, set to 90% of the resolution of the data.

bar.color a numerical value or a character string specifies color for lines. Default to gray.

recurrent.color a numerical value or a character string specifies color for recurrent events. Default to green.
recurrent.shape
   a numerical value or a character string specifies shape for recurrent events. Default to circle.
recurrent.stroke
   a numerical value or a character string specifies stroke for recurrent events. Default to circle.
terminal.color
   a numerical value or a character string specifies color for terminal events. Default to red.
terminal.shape
   a numerical value or a character string specifies shape for terminal events. Default to triangle.
terminal.stroke
   a numerical value or a character string specifies stroke for terminal events. Default to triangle.
not.terminal.color
   a numerical value or a character string specifies color for non-terminal events. Non-terminal events are not plotted at default.
not.terminal.shape
   a numerical value or a character string specifies shape for terminal events. Non-terminal events are not plotted at default.
alpha
   a numerical value specifies the transparency of the points.

See Also

plotEvents

plotHaz(x, newdata = NULL, frailty = NULL, showName = FALSE, type = c("unrestricted", "bounded", "scaled"), smooth = FALSE, control = list(), ...)

plotHaz
   Plot the Baseline Cumulative Hazard Function for the Terminal Time

Description

Plot the baseline cumulative hazard function for an reReg object. The 95% confidence interval on the baseline cumulative rate function

Usage

plotHaz(
   x,  
   newdata = NULL,  
   frailty = NULL,  
   showName = FALSE,  
   type = c("unrestricted", "bounded", "scaled"),  
   smooth = FALSE,  
   control = list(),  
   ...
)


Arguments

- **x**: an object of class `reReg`, returned by the `reReg` function.
- **newdata**: an optional data frame contains variables to include in the calculation of the cumulative rate function. If omitted, the baseline rate function will be plotted.
- **frailty**: an optional vector to specify the shared frailty for `newdata`. If `newdata` is given and `frailty` is not specified, the
- **showName**: an optional logical value indicating whether to label the curves when `newdata` is specified.
- **type**: a character string specifying the type of rate function to be plotted. Options are "unrestricted", "scaled", "bounded". See Details.
- **smooth**: an optional logical value indicating whether to add a smooth curve obtained from a monotone increasing P-splines implemented in package `scam`.
- **control**: a list of control parameters.
- **...**: graphical parameters to be passed to methods. These include `xlab`, `ylab`, `main`, and more. See Details.

Details

The argument `control` consists of options with argument defaults to a list with the following values:

- **xlab**: customizable x-label, default value is "Time".
- **ylab**: customizable y-label, default value is empty.
- **main**: customizable title, default value is "Baseline cumulative hazard function".

These arguments can also be passed down without specifying a `control` list.

Value

A `ggplot` object.

See Also

- `reReg`
- `plot.reReg`

Examples

```r
data(simDat)
fm <- Recur(t.start %to% t.stop, id, event, status) ~ x1 + x2

fit <- reReg(fm, data = simDat, model = "cox|cox", B = 0)
## Plot both the baseline cumulative rate and hazard function
plot(fit)
## Plot baseline cumulative hazard function
plotHaz(fit)
plotHaz(fit, smooth = TRUE)
```
Description

Plot the baseline cumulative rate function for an reReg object.

Usage

plotRate(
  x,
  newdata = NULL,
  frailty = NULL,
  showName = FALSE,
  type = c("unrestricted", "bounded", "scaled"),
  smooth = FALSE,
  control = list(),
  ...
)

Arguments

x an object of class reReg, usually returned by the reReg function.
newdata an optional data frame contains variables to include in the calculation of the cumulative rate function. If omitted, the baseline rate function will be plotted.
frailty an optional vector to specify the shared frailty for newdata. If newdata is given and frailty is not specified, the
showName an optional logical value indicating whether to label the curves when newdata is specified.
type a character string specifying the type of rate function to be plotted. Options are "unrestricted", "scaled", "bounded". See Details.
smooth an optional logical value indicating whether to add a smooth curve obtained from a monotone increasing P-splines implemented in package scam.
control a list of control parameters.
... graphical parameters to be passed to methods. These include xlab, ylab, main, and more. See Details.

Details

The plotRate() plots the estimated baseline cumulative rate function depending on the identifiability assumption. When type = "unrestricted" (default), the baseline cumulative rate function is plotted under the assumption $E(Z) = 1$. When type = "scaled", the baseline cumulative rate function is plotted under the assumption $\Lambda(\min(Y^*, \tau)) = 1$. When type = "bounded", the
baseline cumulative rate function is plotted under the assumption $\Lambda(\tau) = 1$. See ?reReg for the specification of the notations and underlying models.

The argument control consists of options with argument defaults to a list with the following values:

- **xlab** customizable x-label, default value is "Time".
- **ylab** customizable y-label, default value is empty.
- **main** customizable title, default value is "Baseline cumulative rate function".

These arguments can also be specified outside of the control list.

**Value**
A ggplot object.

**See Also**
reReg plot.reReg

**Examples**

data(simDat)
fm <- Recur(t.start %to% t.stop, id, event, status) ~ x1 + x2
fit <- reReg(fm, data = simDat, model = "cox|cox", B = 0)
## Plot both the baseline cumulative rate and hazard function
plot(fit)
## Plot baseline cumulative rate function
plotRate(fit)
plotRate(fit, smooth = TRUE)

---

**reReg**

*Fits Semiparametric Regression Models for Recurrent Event Data*

**Description**

Fits a general (joint) semiparametric regression model for the recurrent event data, where the rate function of the underlying recurrent event process and the hazard function of the terminal event can be specified as a Cox-type model, an accelerated mean model, an accelerated rate model, or a generalized scale-change model. See details for model specifications.

**Usage**

```r
reReg(
  formula,
  data,
  subset,
  model = "cox",
  B = 0,
  se = c("boot", "sand"),
  control = list()
)
```
Arguments

- **formula**: a formula object, with the response on the left of a "~" operator, and the predictors on the right. The response must be a recurrent event survival object as returned by function `Recur`.
- **data**: an optional data frame in which to interpret the variables occurring in the "formula".
- **subset**: an optional logical vector specifying a subset of observations to be used in the fitting process.
- **model**: a character string specifying the underlying model. The available functional form for the rate function and the hazard function include a Cox-type model, an accelerated mean model, an accelerated rate model, or a generalized scale-change model, and can be specified via "cox", "am", "ar", or "gsc", respectively. The rate function and hazard function separated by "|". See Details.
- **B**: a numeric value specifies the number of bootstraps for variance estimation. When \( B = 0 \), variance estimation will not be performed.
- **se**: a character string specifying the method for the variance estimation. See Details.
- **boot**: nonparametric bootstrap approach
- **sand**: resampling-based sandwich estimator
- **control**: a list of control parameters. See `reReg.control` for default values.

Details

Model specification:

Suppose the recurrent event process and the failure events are observed in the time interval \( t \in [0, \tau] \), for some constant \( \tau \). We formulate the recurrent event rate function, \( \lambda(t) \), and the terminal event hazard function, \( h(t) \), in the form of

\[
\lambda(t) = Z \lambda_0(te^{X^T \alpha})e^{X^T \beta},
\]

\[
h(t) = Z h_0(te^{X^T \eta})e^{X^T \theta},
\]

where \( \lambda_0(t) \) is the baseline rate function, \( h_0(t) \) is the baseline hazard function, \( X \) is a \( n \) by \( p \) covariate matrix and \( \alpha \), \( Z \) is an unobserved shared frailty variable, and \( (\alpha, \eta) \) and \( (\beta, \theta) \) correspond to the shape and size parameters, respectively. The model includes several popular semiparametric models as special cases, which can be specified via the `model` argument with the rate function and the hazard function separated by "|". For examples, Wang, Qin and Chiang (2001) \( (\alpha = \eta = \theta = 0) \) can be called with `model = "cox"`; Huang and Wang (2004) \( (\alpha = \eta = 0) \) can be called with `model = "cox|cox"`; Xu et al. (2017) \( (\alpha = \beta \text{ and } \eta = \theta = 0) \) can be called with `model = "am|am"`; Xu et al. (2019) \( (\eta = \theta = 0) \) can be called with `model = "gsc"`. Users can mix the models depending on the application. For example, `model = "cox|ar"` postulate a Cox proportional model for the recurrent event rate function and an accelerated rate model for the terminal event hazard function \( (\alpha = \theta = 0) \). If only one model is specified without an "|", it is used for both the rate function and the hazard function. For example, specifying `model = "cox"` is equivalent to `model = "cox|cox"`. Some models that assumes \( Z = 1 \) and requires independent censoring are also implemented in `reReg`; these includes `model = "cox.LWYY"` for Lin et al. (2000), `model = "cox.GL"` for Ghosh and Lin (2002), and `model = "am.GL"` for Ghosh and Lin (2003). Additionally, an improved estimation of the proportional rate model (Huang and Huang 2022) can be called by `model = "cox.HH"` with additional control options to specify the underlying procedure. See online vignette for a detailed discussion of the implemented regression models.
**Variance estimation:**
The available methods for variance estimation are:

- **boot** performs nonparametric bootstrap.
- **sand** performs the efficient resampling-based variance estimation.

**Improving proportional rate model:** A common semiparametric regression model for recurrent event process under the noninformative censoring assumption is the Cox-type proportional rate model (available in `reReg()` via `model = "cox.LWYY"`). However, the construction of the pseudo-partial score function ignores the dependency among recurrent events and thus could be inefficient. To improve upon this popular method, Huang and Huang (2022) proposed to combine a system of weighted pseudo-partial score equations via the generalized method of moments (GMM) and empirical likelihood (EL) estimation. The proposed GMM and EL procedures are available in `reReg` via `model = "cox.HH"` with additional control specifications. See online vignette for an illustration of this feature.

**Control options:**
The control list consists of the following parameters:

- **tol** absolute error tolerance.
- **init** a list contains initial guesses used for root search.
- **solver** the equation solver used for root search. The available options are `BB::BBsolve`, `BB::dfsane`, `BB::BBoptim`, `optimx::optimr`, `dfoptim::hjk`, `dfoptim::mads`, `optim`, and `nleqslv::nleqslv`.
- **eqType** a character string indicating whether the log-rank type estimating equation or the Gehan-type estimating equation (when available) will be used.
- **boot.parallel** a logical value indicating whether parallel computation will be applied when `se = "boot"` is called.
- **boot.parCl** an integer value specifying the number of CPU cores to be used when `parallel = TRUE`. The default value is half the CPU cores on the current host.
- **cppl** A character string indicating either to improve the proportional rate model via the generalized method of moments (`cppl = "GMM"`) or empirical likelihood estimation (`cppl = "EL"`). This option is only used when `model = "cox.HH"`.
- **cppl.wfun** A list of (up to two) weight functions to be combined with the weighted pseudo-partial likelihood scores. Available options are "Gehan" and "cumbase", which correspond to the Gehan’s weight and the cumulative baseline hazard function, respectively. Alternatively, the weight functions can be specified with function formulas. This option is only used when `model = "cox.HH"`.
- **trace** A logical variable denoting whether some of the intermediate results of iterations should be displayed to the user. Default is `FALSE`.

**References**


Huang, M.-Y. and Huang, C.-Y. (2022). Improved semiparametric estimation of the proportional rate model with recurrent event data. *In revision*.

See Also

`Recur`, `simGSC`

Examples

data(simDat)

```r
## Nonparametric estimate
plot(reReg(Recur(t.start %to% t.stop, id, event, status) ~ 1, data = simDat, B = 50))

fm <- Recur(t.start %to% t.stop, id, event, status) ~ x1 + x2
## Fit the Cox rate model
summary(reReg(fm, data = simDat, model = "cox", B = 50))
## Fit the joint Cox/Cox model
summary(reReg(fm, data = simDat, model = "cox|cox", B = 50))
## Fit the scale-change rate model
summary(reReg(fm, data = simDat, model = "gsc", B = 50, se = "sand"))
```

---

**reReg.control**

Package options for reReg

### Description

This function provides the fitting options for the `reReg()` function.

### Usage

```r
reReg.control(
  eqType = c("logrank", "gehan", "gehan_s"),
  solver = c("BB::dfsane", "BB::BBsolve", "BB::BBoptim", "optimx::optimr",
            "dfoptim::hjk", "dfoptim::mads", "optim", "nleqslv::nleqslv"),
```
tol = 1e-07,
cpl = NULL,
cpl.wfun = list(NULL, NULL),
init = list(alpha = 0, beta = 0, eta = 0, theta = 0),
boot.parallel = FALSE,
boot.parCl = NULL,
maxit1 = 100,
maxit2 = 10,
trace = FALSE,
umAdj = 0.001
)

Arguments

- **eqType**: A character string indicating whether the log-rank type estimating equation or the Gehan-type estimating equation (when available) will be used.
- **solver**: A character string specifying the equation solver to be used for root search.
- **tol**: A character string indicating either to improve the proportional rate model via the generalized method of moments (cpl = "GMM") or empirical likelihood estimation (cpl = "EL"). This option is only used when model = "cox.HH".
- **cpl.wfun**: A list of (up to two) weight functions to be combined with the weighted pseudo-partial likelihood scores. Available options are "Gehan" and "cumbase", which correspond to the Gehan’s weight and the cumulative baseline hazard function, respectively. Alternatively, the weight functions can be specified with function formulas. This option is only used when model = "cox.HH".
- **init**: A list contains the initial guesses used for root search.
- **boot.parallel**: An logical value indicating whether parallel computation will be applied when se = "boot" is specified in reReg().
- **boot.parCl**: An integer value specifying the number of CPU cores to be used when parallel = TRUE. The default value is half the CPU cores on the current host.
- **maxit1, maxit2**: Max number of iteration used when model = "cox.HH".
- **trace**: A logical variable denoting whether some of the intermediate results of iterations should be displayed to the user. Default is FALSE.
- **numAdj**: A positive numerical value specifying the small constant used in heuristic adjustment of the borrow strength method.

See Also

reReg
residuals.reReg

Calculate Residuals for a ‘reReg’ Fit

Description

Calculates residuals for a joint frailty scale-change model fitted by ‘reReg’. Under the recurrent
event model, at each observation time, \( t \), the residual is calculated as

\[
\text{observed number of recurrent events at } t - \text{expected number of recurrent events at } t.
\]

The expected number of recurrent events at \( t \) is calculated by the cumulative rate function at \( t \).
Under the failure time model, the residual is calculated as

\[
\Delta - H(t),
\]

where \( \Delta \) is the terminal event indicator and \( H(t) \) is the cumulative hazard function at \( t \).

Usage

## S3 method for class 'reReg'
residuals(object, model = c("recurrent", "failure"), ...)

Arguments

- **object**: an object of class reReg returned by the reReg() function.
- **model**: a character string specifying whether the residuals will be calculated under the
  recurrent event model or the failure time model.
- ... additional parameters for future development.

---

reSurv

Create an reSurv Object

Description

Create a recurrent event survival object, used as a response variable in reReg. This function is
deprecated in Version 1.1.6. A recurrent event object is now being created with Recur(). See
'!?Recur()' for details.

Usage

reSurv(time1, time2, id, event, status, origin = 0)
Arguments

time1 when "time2" is provided, this vector is treated as the starting time for the gap time between two successive recurrent events. In the absence of "time2", this is the observation time of recurrence on calendar time scale, in which, the time corresponds to the time since entry/inclusion in the study.

time2 an optional vector for ending time for the gap time between two successive recurrent events.

id subject’s id.

event a binary vector used as the recurrent event indicator. event = 1 for recurrent times.

status a binary vector used as the status indicator for the terminal event. status = 0 for censored times.

origin a numerical vector indicating the time origin of subjects. When origin is a scalar, reSurv assumes all subjects have the same origin. Otherwise, origin needs to be a numerical vector, with length equals to the number of subjects. In this case, each element corresponds to different origins for different subjects. This argument is only needed when "time2" is missing.

Examples

```r
## Not run:
data(simDat)
## being deprecated in Version 1.1.7
with(dat, reSurv(Time, id, event, status))
## Use Recur() instead
with(dat, Recur(Time, id, event, status))
## End(Not run)
```

---

**simDat**

*Simulated dataset for demonstration*

Description

A simulated data frame with the following variables

- **id** subjects identification
- **t.start** start of the interval
- **t.stop** endpoint of the interval; when time origin is 0 this variable also marks the recurrence or terminal/censoring time
- **status** terminal event indicator; 1 if a terminal event is recorded
- **event** recurrent event indicator; 1 if a recurrent event is recorded
- **x1** baseline covariate generated from a standard uniform distribution
- **x2** baseline covariate generated from a standard uniform distribution (independent from z1
Usage

data(simDat)

Format

A data frame with 874 rows and 7 variables.

Details

The sample dataset simDat is generated by set.seed(0); dat <- simGSC(200). See simGSC for instruction on simulating recurrent event data from scale-change models.

Description

The function simGSC() generates simulated recurrent event data from either a Cox-type model, an accelerated mean model, an accelerated rate model, or a generalized scale-change model.

Usage

simGSC(
  n, 
  summary = FALSE, 
  para, 
  xmat, 
  censoring, 
  frailty, 
  tau, 
  origin, 
  Lam0, 
  Haz0
)

Arguments

n number of observation.
summary a logical value indicating whether a brief data summary will be printed.
para a list of numerical vectors for the regression coefficients in the joint scale-change model. The names of the list elements are alpha, beta, eta, and theta, correspond to \( \alpha, \beta, \eta, \) and \( \theta \) in the joint scale-change model, respectively. See Details for reReg.
xmat an optional matrix specifying the design matrix.
censoring a numeric variable specifying the censoring times for each of the \( n \) observation.
**frailty**  
a numeric variable specifying the frailty variable.

**tau**  
a numeric value specifying the maximum observation time.

**origin**  
a numeric value specifying the time origin.

**Lam0**  
is an optional function that specifies the baseline cumulative rate function. When left-unspecified, the recurrent events are generated using the baseline rate function of

\[ \lambda_0(t) = \frac{2}{1 + t}, \]

or equivalently, the cumulative rate function of

\[ \Lambda_0(t) = 2 \log(1 + t). \]

**Haz0**  
is an optional function that specifies the baseline hazard function. When left-unspecified, the recurrent events are generated using the baseline hazard function

\[ h_0(t) = \frac{1}{5(1 + t)}, \]

or equivalently, the cumulative hazard function of

\[ H_0(t) = \log(1 + t)/5. \]

### Details

The function `simGSC()` generates simulated recurrent event data over the interval \((0, \tau)\) based on the specification of the recurrent process and the terminal events. Specifically, the rate function, \(\lambda(t)\), of the recurrent process can be specified as one of the following model:

\[
\lambda(t) = Z \lambda_0(te^{X^T \alpha})e^{X^T \beta}, \quad h(t) = Z h_0(te^{X^T \eta})e^{X^T \theta},
\]

where \(\lambda_0(t)\) is the baseline rate function, \(h_0(t)\) is the baseline hazard function, \(X\) is a \(n\) by \(p\) covariate matrix and \(\alpha, Z\) is an unobserved shared frailty variable, and \((\alpha, \eta)\) and \((\beta, \theta)\) correspond to the shape and size parameters of the rate function and the hazard function, respectively.

Under the default settings, the `simGSC()` function assumes \(p = 2\) and the regression parameters to be \(\alpha = \eta = (0, 0)^T\), and \(\beta = \theta = (1, 1)^T\). When the `xmat` argument is not specified, the `simGSC()` function assumes \(X_i\) is a two-dimensional vector \(X_i = (X_{i1}, X_{i2})\), \(i = 1, \ldots, n\), where \(X_{i1}\) is a Bernoulli variable with rate 0.5 and \(X_{i2}\) is a standard normal variable. With the default `xmat`, the censoring time \(C\$\) is generated from an independent uniform distribution in \([0, 2\tau X_{i1} + 2Z^2\tau(1 - X_{i1})]\). Thus, the censoring distribution is covariate dependent and is informative when \(Z\) is not a constant. When the `frailty` argument is not specified, the frailty variable \(Z\) is generated from a gamma distribution with a unit mean and a variance of 0.25. The default values for `tau` and `origin` are 60 and 0, respectively. When arguments `Lam0` and `Haz0` are left unspecified, the `simGSC()` function uses \(\Lambda_0(t) = 2 \log(1 + t)\) and \(H_0(t) = \log(1 + t)/5\), respectively. This is equivalent to setting \(\text{Lam0} = \text{function}(x) \times 2 \times \log(1 + x)\) and \(\text{Haz0} = \text{function}(x) \times \log(1 + x) / 5\). Overall, the default specifications generate the recurrent events and the terminal events from the model:

\[
\lambda(t) = Z \frac{2}{1 + te^{-X_{i1} - X_{i2}}}, \quad h(t) = Z \frac{1}{5(1 + te^{X_{i1} + X_{i2}})}, \quad t \in [0, 60].
\]

See online vignette for more examples.
See Also

reReg

Examples

set.seed(123)
simGSC(100, summary = TRUE)
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