

Package ‘simevent’

May 9, 2026

Title Simulation and Analysis of Event History Data

Version 0.1.1

Description Simulate event history data from a framework where treatment decisions and disease progression are represented as counting process. The user can specify number of events and parameters of intensities thereby creating a flexible simulation framework.

License MIT + file LICENSE

Encoding UTF-8

RoxygenNote 7.3.2

URL <https://github.com/miclukacova/simevent>

BugReports <https://github.com/miclukacova/simevent/issues>

Imports stats, data.table, ggplot2, survival, Rcpp

Suggests knitr, rmarkdown, testthat (>= 3.0.0)

Config/testthat/edition 3

LinkingTo Rcpp

VignetteBuilder knitr

NeedsCompilation yes

Author Michaela Lukacova [aut, cre]

Maintainer Michaela Lukacova <michaela.lukacova@sund.ku.dk>

Repository CRAN

Date/Publication 2026-04-24 08:30:02 UTC

Contents

alphaSim	2
intEffectAlpha	3
IntFormatData	5
inverseScHaz	6
inverseScHazTV	7

plotEventData	8
sampleEvents	9
simCRdata	9
simDisease	10
simDropIn	12
simEventCox	16
simEventData	17
simEventObj	20
simEventTV	21
simStatinData	23
simSurvData	24
simTreatment	25

Index	28
--------------	-----------

alphaSim	<i>Simulation and Estimation with Modified Shape Parameter</i>
----------	--

Description

This function simulates event history data from either the Disease, Treatment or Drop In setting. See `simDisease`, `simTreatment` and `simDropIn`. The shape parameter η of respectively the disease process, the Drop In process and the Treatment process is multiplied by `alpha`. The function either

- returns the proportion of individuals who experience death and the proportion of individuals who experience disease/drop in/treatment by a specified time τ (in group $A_0 = a_0$ for drop in and disease).
- returns number of years lost before τ of death and disease/drop in/treatment
- returns simulated data. One can specify all the same parameters as in the functions `simDisease`, `simTreatment` and `simDropIn`.

Usage

```
alphaSim(
  N = 10000,
  eta = rep(0.1, 4),
  nu = rep(1.1, 4),
  alpha = 0.5,
  tau = 5,
  a0 = 1,
  years_lost = FALSE,
  setting = "Disease",
  return_data = FALSE,
  cens = 0,
  ...
)
```

Arguments

N	Integer. Number of individuals to simulate. Default is 10,000.
eta	Numeric vector. Shape parameters for Weibull hazards. Length of the vector should match number of events. For the Disease and Drop In setting this is 4. For the Treatment setting, this is 3. (default <code>rep(0.1, 4)</code>).
nu	Numeric vector. Scale parameters for Weibull hazards. Length of the vector should match number of events. For the Disease and Drop In setting this is 4. For the Treatment setting, this is 3. (default <code>rep(1.1, 4)</code>).
alpha	Numeric scalar. Multiplicative factor applied to the disease process shape parameter η .
tau	Numeric scalar. Time horizon at which proportions are computed.
a0	Binary (0/1). Specifies the group for comparison in setting Drop In and Disease.
years_lost	Logical. If TRUE, computes years lost instead of proportions.
setting	Character string. Must be either "Disease", "Drop In" or "Treatment". Depending on the simulation setting.
return_data	Logical. If TRUE the simulated data is returned.
cens	Binary scalar. Indicates whether individuals are at risk of censoring (default 0).
...	Additional arguments passed to respectively <code>simDisease</code> , <code>simTreatment</code> and <code>simDropIn</code> .

Value

A list with two components:

`effect_L` Proportion (or years lost) of individuals diagnosed with disease by time τ , under intervention.

`effect_death` Proportion (or years lost) of individuals who died by time τ under intervention.

Or the simulated data.

Examples

```
alphaSim(N = 100, eta = rep(0.1,3), nu = rep(1.1,3), alpha = 0.5, setting = "Disease")
alphaSim(N = 100, setting = "Drop In", beta_A0_Z = 1)
```

intEffectAlpha

Estimate Effect of Intervention: Modifying Eta Parameter of Process

Description

This function simulates data from the disease setting in two scenarios. Under intervention on the shape parameter η of the disease process is multiplied by `alpha`, and a baseline (non-intervened) scenario. It computes the proportion of individuals who experience death or disease by a specified time τ in the group $A0 = a0$, optionally returning `years_lost`. The function can also plot a sample of the event data for each scenario for comparison.

Usage

```
intEffectAlpha(
  N = 10000,
  setting = "Disease",
  eta = rep(0.1, 4),
  nu = rep(1.1, 4),
  alpha = 0.5,
  tau = 5,
  a0 = 1,
  years_lost = FALSE,
  plot = TRUE,
  lower = 10^(-30),
  upper = 200,
  cens = 0,
  ...
)
```

Arguments

<code>N</code>	Integer. Number of individuals to simulate. Default is 10,000.
<code>setting</code>	Character string. Must be either "Disease", "Drop In" or "Treatment". Depending on the simulation setting.
<code>eta</code>	Numeric vector of length 3. Shape parameters for Weibull hazards (default <code>rep(0.1, 4)</code>).
<code>nu</code>	Numeric vector of length 3. Scale parameters for Weibull hazards (default <code>rep(1.1, 4)</code>).
<code>alpha</code>	Numeric scalar. Multiplicative factor applied to the disease process shape parameter η .
<code>tau</code>	Numeric scalar. Time horizon at which proportions are computed.
<code>a0</code>	Binary (0/1). Specifies the group for comparison. Only relevant in setting "Drop In" and "Disease".
<code>years_lost</code>	Logical. If TRUE, computes years lost instead of proportions.
<code>plot</code>	Logical. If TRUE, plots timelines for sample of intervention and non intervention data.
<code>lower</code>	Numeric scalar. Lower bound for root-finding in hazard inversion (default <code>1e-30</code>).
<code>upper</code>	Numeric scalar. Upper bound for root-finding in hazard inversion (default 200).
<code>cens</code>	Binary scalar. Indicates whether individuals are at risk of censoring (default 0).
<code>...</code>	Additional arguments passed to respectively <code>simDisease</code> , <code>simTreatment</code> and <code>simDropIn</code> .

Value

A list with two components:

effect_L Proportion (or years lost) of individuals diagnosed with disease by time τ in group $A_0 = a_0$, under intervention.

effect_death Proportion (or years lost) of individuals who died by time τ in group $A_0 = a_0$, under intervention.

Examples

```
intEffectAlpha(N = 1000, alpha = 0.7, tau = 5, years_lost = FALSE, a0 = 1, setting = "Drop In")
```

IntFormatData *Transform Event Data into Interval Format for Classical Inference*

Description

Converts simulated event history data with time-dependent covariates into an interval (start-stop) format, suitable for classical survival analysis functions like `coxph`. Adds interval start and stop times (`tstart`, `tstop`) and a counting variable `k` indexing events. Optionally, the function can split intervals at a specified time point to accommodate estimation of time-varying effects.

Usage

```
IntFormatData(data, N_cols = 6:9, timeVar = FALSE, t_prime = NULL)
```

Arguments

<code>data</code>	A <code>data.frame</code> or <code>data.table</code> containing event data with columns <code>ID</code> , <code>Time</code> , <code>Delta</code> , and counting process columns specified by <code>N_cols</code> .
<code>N_cols</code>	Integer vector. Column indices of data that correspond to counting process variables. Defaults to <code>6:9</code> .
<code>timeVar</code>	Logical. If <code>TRUE</code> , the intervals are split at <code>t_prime</code> to allow time-varying covariate effects. Default is <code>FALSE</code> .
<code>t_prime</code>	Numeric scalar. Time point at which to split intervals if <code>timeVar = TRUE</code> .

Value

A `data.table` with columns `tstart`, `tstop`, `k`, and other original variables, formatted for survival analysis.

Examples

```
data <- simEventData(10)
IntFormatData(data)
```

`inverseSchaz`*Wrapper for inverse cumulative hazard*

Description

A wrapper around the Rcpp function `inverseSchaz`, used to find the inverse by numeric methods in case of no simple analytical solution.

Usage

```
inverseSchaz(  
  p,  
  t,  
  lower,  
  upper,  
  eta,  
  nu,  
  phi,  
  at_risk,  
  tol = 1e-09,  
  max_iter = 100  
)
```

Arguments

<code>p</code>	The random variable (typically $-\log(U)$).
<code>t</code>	Current time.
<code>lower</code>	Lower bound for root finding.
<code>upper</code>	Upper bound for root finding.
<code>eta</code>	Numeric vector of shape parameters.
<code>nu</code>	Numeric vector of scale parameters.
<code>phi</code>	Numeric vector of multiplicative effects.
<code>at_risk</code>	Numeric vector indicating at-risk indicators for each event type.
<code>tol</code>	Numeric tolerance for root-finding. Default is $1e-9$.
<code>max_iter</code>	Maximum iterations. Default is 100.

Value

A numeric scalar, the root u .

Examples

```
eta <- c(0.1, 0.1)
nu <- c(1.1, 1.1)
phi <- c(1, 1)
at_risk <- c(1, 1)
inverseScHaz(0.5, t = 0, lower = 1e-15, upper = 200, eta, nu, phi, at_risk)
```

inverseScHazTV

Wrapper for inverse cumulative hazard

Description

A wrapper around the Rcpp function `inverseScHazCppTV`, used to find the inverse of the summed cumulative hazard.

Usage

```
inverseScHazTV(
  p,
  t,
  lower = 1e-15,
  upper = 200,
  t_prime,
  eta,
  nu,
  phi,
  phi_prime,
  at_risk,
  tol = 1e-09,
  max_iter = 100
)
```

Arguments

<code>p</code>	The random variable (typically $-\log(U)$).
<code>t</code>	The time of the previous event
<code>lower</code>	Lower bound for root finding.
<code>upper</code>	Upper bound for root finding.
<code>t_prime</code>	The time where the time varying effects change
<code>eta</code>	Numeric vector of shape parameters.
<code>nu</code>	Numeric vector of scale parameters.
<code>phi</code>	Numeric vector of multiplicative effect before time <code>t_prime</code>
<code>phi_prime</code>	Numeric vector of multiplicative effects after time <code>t_prime</code>
<code>at_risk</code>	Numeric vector indicating at-risk indicators for each event type.
<code>tol</code>	Numeric tolerance for root-finding. Default is $1e-9$.
<code>max_iter</code>	Maximum iterations. Default is 100.

Value

A numeric scalar, the root u .

Examples

```
eta <- c(0.1, 0.1)
nu <- c(1.1, 1.1)
phi <- c(1, 1)
at_risk <- c(1, 1)
phi_prime <- c(2, 2)
inverseScHazTV(p = 0.5, t = 1, t_prime = 2, eta = eta, nu = nu,
               phi = phi, phi_prime = phi_prime, at_risk = at_risk)
```

plotEventData

Plot Simulated Event History Data

Description

Visualizes event history data by plotting individual event times colored and shaped by event type. Each individual's timeline is displayed horizontally with events marked along it.

Usage

```
plotEventData(data, title = "Event Data")
```

Arguments

data	A data.frame or data.table containing at least the columns ID, Time, and Delta.
title	Character string specifying the plot title. Defaults to "Event Data".

Value

A ggplot object representing the event data visualization.

Examples

```
data <- simEventData(10)
plotEventData(data)
```

sampleEvents	<i>Sample event types from matrix of probabilities</i>
--------------	--

Description

Sample event types from matrix of probabilities

Usage

```
sampleEvents(probs)
```

Arguments

probs A matrix where each column is a probability vector

Value

A vector of sampled event types (0-indexed)

simCRdata	<i>Simulate Competing Risks Data</i>
-----------	--------------------------------------

Description

Simulates competing risks data for N individuals who are at risk of mutually exclusive event types. Three event types are simulated, where one can be interpreted as censoring.

Usage

```
simCRdata(N, beta = NULL, eta = rep(0.1, 3), nu = rep(1.1, 3), cens = 1, ...)
```

Arguments

N	Integer. Number of individuals to simulate.
beta	Numeric matrix of dimension 2x3. Covariate effects of L_0 and A_0 on the three competing processes (columns correspond to processes). Defaults to zero matrix if NULL.
eta	Numeric vector of length 3. Shape parameters for Weibull hazards, parameterized as $\eta\nu t^{\nu-1}$. Defaults to <code>rep(0.1, 3)</code> .
nu	Numeric vector of length 3. Scale parameters for Weibull hazards. Defaults to <code>rep(1.1, 3)</code> .
cens	Binary (0 or 1). Indicates if a censoring process is included. Default is 1.
...	Additional arguments passed to <code>simEventData</code> , including <code>add_cov</code> for extra covariates.

Details

The event intensities follow Weibull hazard models parameterized by shape and scale parameters η and ν . Covariate effects on the hazard are specified by the beta matrix, which models the effects of baseline covariates L_0 and A_0 on each event type.

Value

A data frame with simulated competing risk data including:

- ID - Individual identifier.
- Time - Event time.
- Delta - Event type (0, 1, or 2).
- L_0 - Baseline covariate.
- A_0 - Baseline treatment indicator.

Examples

```
simCRdata(10)
```

simDisease

Simulate Data in a Disease Setting

Description

This function simulates event data representing three event types: Censoring (0), Death (1), and Change in Covariate Process (2). Death and Censoring are terminal events, while Change in Covariate Process can occur only once.

Usage

```
simDisease(  
  N,  
  eta = rep(0.1, 3),  
  nu = rep(1.1, 3),  
  cens = 1,  
  beta_L0_D = 1,  
  beta_L0_L = 1,  
  beta_L_D = 1,  
  beta_A0_D = 0,  
  beta_A0_L = 0,  
  beta_L0_C = 0,  
  beta_A0_C = 0,  
  beta_L_C = 0,  
  followup = Inf,  
  lower = 10(-15),  
  upper = 200,  
)
```

```

    beta_L_D_t_prime = NULL,
    t_prime = NULL,
    gen_A0 = NULL,
    at_risk_cov = NULL,
    ...
)

```

Arguments

N	Numeric scalar. Number of individuals to simulate.
eta	Numeric vector of length 3. Shape parameters for Weibull intensities with parameterization $\eta\nu t^{\nu-1}$. Defaults to rep(0.1, 3).
nu	Numeric vector of length 3. Scale parameters for the Weibull hazards. Defaults to rep(1.1, 3).
cens	Binary scalar. Indicates whether individuals are at risk of censoring (default 1).
beta_L0_D	Numeric scalar. Effect of baseline covariate L0 on death risk (default 1).
beta_L0_L	Numeric scalar. Effect of baseline covariate L0 on covariate change risk (default 1).
beta_L_D	Numeric scalar. Effect of covariate change (L = 1) on death risk (default 1).
beta_A0_D	Numeric scalar. Effect of baseline treatment (A0 = 1) on death risk (default 0).
beta_A0_L	Numeric scalar. Effect of baseline treatment (A0 = 1) on covariate change risk (default 0).
beta_L0_C	Numeric scalar. Effect of baseline covariate L0 on censoring probability (default 0).
beta_A0_C	Numeric scalar. Effect of baseline treatment A0 on censoring probability (default 0).
beta_L_C	Numeric scalar. Effect of covariate change (L = 1) on censoring probability (default 0).
followup	Numeric scalar. Maximum follow-up (censoring) time. Defaults to Inf.
lower	Numeric scalar. Lower bound for root-finding (inverse cumulative hazard) (default 1e-15).
upper	Numeric scalar. Upper bound for root-finding (default 200).
beta_L_D_t_prime	Numeric scalar or NULL. Additional effect of covariate change on death risk after time t_prime (optional).
t_prime	Numeric scalar or NULL. Time point where effects change (optional).
gen_A0	Function. Function to generate the baseline treatment covariate A0. Takes N and L0 as inputs. Default is a Bernoulli(0.5) random variable.
at_risk_cov	Function. Function determining if an individual is at risk for each event type, given their covariates. Takes a numeric vector covariates and returns a binary vector. Default returns 1 for all events.
...	Additional arguments passed to simEventData or simEventTV

Details

Event intensities depend on previous events and predefined parameters ν and η .

The arguments `beta_X_Y` control how the X affects Y. A positive value means that a higher value of X increases the intensity of Y, while a negative value decreases the intensity.

The simulation uses an event history framework with terminal events (death, censoring) and a single recurrent covariate change. The event intensities depend on covariates and previous events according to user-specified parameters. Time-varying effects can be included via `beta_L_D_t_prime` and `t_prime`.

Value

A data frame containing the simulated data with columns:

ID	Individual identifier
Time	Time of the event
Delta	Event type (0 = censoring, 1 = death, 2 = covariate change)
L0	Baseline covariate
L	Covariate indicating change in covariate process

Examples

```
simDisease(10)
```

simDropIn	simDropIn
-----------	-----------

Description

Simulate Event Data from a "Drop In" Setting

Usage

```
simDropIn(
  N,
  eta = c(0.5, 0.5, 0.1, 0.25),
  nu = c(1.1, 1.1, 1.1, 1.1),
  adherence = FALSE,
  followup = Inf,
  cens = 1,
  generate.A0 = function(N, L0) stats::rbinom(N, 1, 0.5),
  lower = 1e-200,
  upper = 1e+10,
  t_prime = NULL,
  at_risk_cov = NULL,
```

```
beta_L_A = 1,  
beta_L_Z = 2,  
beta_L_D = 1.5,  
beta_L_C = 0,  
beta_A_L = -0.5,  
beta_A_Z = -0.5,  
beta_A_D = -1,  
beta_A_C = 0,  
beta_Z_L = -1,  
beta_Z_A = 0,  
beta_Z_D = -1,  
beta_Z_C = 0,  
beta_L0_L = 1,  
beta_L0_A = 1,  
beta_L0_Z = 1,  
beta_L0_D = 1,  
beta_L0_C = 0,  
beta_A0_L = -1.5,  
beta_A0_A = 0,  
beta_A0_Z = 0,  
beta_A0_D = -2,  
beta_A0_C = 0,  
beta_L_A_prime = 0,  
beta_L_Z_prime = 0,  
beta_L_D_prime = 0,  
beta_L_C_prime = 0,  
beta_A_L_prime = 0,  
beta_A_Z_prime = 0,  
beta_A_D_prime = 0,  
beta_A_C_prime = 0,  
beta_Z_L_prime = 0,  
beta_Z_A_prime = 0,  
beta_Z_D_prime = 0,  
beta_Z_C_prime = 0,  
beta_L0_L_prime = 0,  
beta_L0_A_prime = 0,  
beta_L0_Z_prime = 0,  
beta_L0_D_prime = 0,  
beta_L0_C_prime = 0,  
beta_A0_L_prime = 0,  
beta_A0_A_prime = 0,  
beta_A0_Z_prime = 0,  
beta_A0_D_prime = 0,  
beta_A0_C_prime = 0,  
...  
)
```

Arguments

N	Integer. Number of individuals to simulate.
eta	Numeric vector of length 4 (or 5). Shape parameters of the Weibull baseline intensity for each event type. $\eta\nu t^{\nu-1}$
nu	Numeric vector of length 4 (or 5). Scale parameters for the Weibull hazard.
adherence	Logical. Indicator of whether a Treatment process should be simulated.
followup	Numeric. Maximum censoring time. Events occurring after this time are censored. Default is Inf (no censoring).
cens	Logical. Indicator of whether there should be a censoring process.
generate.A0	Function. Function to generate the baseline treatment covariate A0. Takes N and L0 as inputs. Default is a Bernoulli(0.5) random variable.
lower	Numeric. Lower bound for root-finding in inverse cumulative hazard calculations. Default is 10^{-15} .
upper	Numeric. Upper bound for root-finding in inverse cumulative hazard calculations. Default is 200.
t_prime	Numeric scalar or NULL. Time point where effects change (optional).
at_risk_cov	Function. Function determining if an individual is at risk for each event type, given their covariates. Takes a numeric vector covariates and returns a binary vector. Default returns 1 for all events.
beta_L_A	Numeric. Specifies how L affects A.
beta_L_Z	Numeric. Specifies how L affects Z.
beta_L_D	Numeric. Specifies how L affects D.
beta_L_C	Numeric. Specifies how L affects C.
beta_A_L	Numeric. Specifies how L affects A.
beta_A_Z	Numeric. Specifies how L affects Z.
beta_A_D	Numeric. Specifies how L affects D.
beta_A_C	Numeric. Specifies how L affects C.
beta_Z_L	Numeric. Specifies how L affects A.
beta_Z_A	Numeric. Specifies how L affects Z.
beta_Z_D	Numeric. Specifies how L affects D.
beta_Z_C	Numeric. Specifies how L affects C.
beta_L0_L	Numeric. Specifies how L affects A.
beta_L0_A	Numeric. Specifies how L affects Z.
beta_L0_Z	Numeric. Specifies how L affects Z.
beta_L0_D	Numeric. Specifies how L affects D.
beta_L0_C	Numeric. Specifies how L affects C.
beta_A0_L	Numeric. Specifies how L affects A.

beta_A0_A	Numeric. Specifies how L affects Z.
beta_A0_Z	Numeric. Specifies how L affects Z.
beta_A0_D	Numeric. Specifies how L affects D.
beta_A0_C	Numeric. Specifies how L affects C.
beta_L_A_prime	Numeric. Specifies how L additionally affects A after time t_prime.
beta_L_Z_prime	Numeric. Specifies how L additionally affects Z after time t_prime.
beta_L_D_prime	Numeric. Specifies how L additionally affects D after time t_prime.
beta_L_C_prime	Numeric. Specifies how L additionally affects C after time t_prime.
beta_A_L_prime	Numeric. Specifies how L additionally affects A after time t_prime.
beta_A_Z_prime	Numeric. Specifies how L additionally affects Z after time t_prime.
beta_A_D_prime	Numeric. Specifies how L additionally affects D after time t_prime.
beta_A_C_prime	Numeric. Specifies how L additionally affects C after time t_prime.
beta_Z_L_prime	Numeric. Specifies how L additionally affects A after time t_prime.
beta_Z_A_prime	Numeric. Specifies how L additionally affects Z after time t_prime.
beta_Z_D_prime	Numeric. Specifies how L additionally affects D after time t_prime.
beta_Z_C_prime	Numeric. Specifies how L additionally affects C after time t_prime.
beta_L0_L_prime	Numeric. Specifies how L additionally affects after time A.
beta_L0_A_prime	Numeric. Specifies how L additionally affects after time Z.
beta_L0_Z_prime	Numeric. Specifies how L additionally affects after time Z.
beta_L0_D_prime	Numeric. Specifies how L additionally affects after time D.
beta_L0_C_prime	Numeric. Specifies how L additionally affects after time C.
beta_A0_L_prime	Numeric. Specifies how L additionally affects after time A.
beta_A0_A_prime	Numeric. Specifies how L additionally affects after time Z.
beta_A0_Z_prime	Numeric. Specifies how L additionally affects after time Z.
beta_A0_D_prime	Numeric. Specifies how L additionally affects after time D.
beta_A0_C_prime	Numeric. Specifies how L additionally affects after time C.
...	Additional arguments passed to simEventData or simEventTV

Details

simDropIn is a function that simulates data corresponding to N individuals that are at risk for 4 or 5 events. Censoring (C), Death (D), Drop In Initiation (Z), Change in Covariate Process (L) and optionally Treatment (A).

Value

Data frame containing the simulated event history data

Examples

```
simDropIn(10)
```

simEventCox

Simulate Event History Data Based on Cox Models

Description

Simulates recurrent and terminal event data for a cohort of individuals based on a list of fitted Cox proportional hazards models. Each event type is governed by its own model, and simulation proceeds by iteratively sampling event times until a terminal event occurs.

Usage

```
simEventCox(
  N,
  cox_fits,
  list_old_vars = NULL,
  n_event_max = c(1, 1),
  term_events = 1,
  intervention1 = NULL,
  intervention2 = NULL
)
```

Arguments

N	Integer. The number of individuals to simulate.
cox_fits	A named list of fitted Cox proportional hazards models (coxph objects), one for each event type. The names are used as event type labels.
list_old_vars	A named list containing the old covariates. New covariates will be simulated by drawing from the old covariates with replacement.
n_event_max	Integer vector. Maximum number of times each event type can occur per individual.
term_events	Integer or integer vector. Indices of event types that are terminal, i.e., events that stop further simulation for an individual.
intervention1	Optional function. Takes arguments (j, sim_matrix) and returns an updated simulation matrix. Used to modify covariates dynamically at each event iteration.
intervention2	Optional function. Takes arguments (j, H_j) and returns a modified baseline cumulative hazard vector for event type j. Allows dynamic hazard modification. The function intervention2 <- function(j, basehaz) if(j == 2) 1.15 * basehaz else basehaz performs an intervention where the baseline hazard of process 2 is multiplied by 1.15.

Details

The function simulates individual event histories by:

1. Sampling initial baseline covariates by resampling observed values.
2. Extracting baseline cumulative hazard functions from the Cox models.
3. Iteratively sampling event times.
4. Updating covariate histories and event counts.
5. Stopping simulation per individual after a terminal event or maximum events reached.

Value

A data.table with one row per event per individual containing:

- ID — Individual identifier.
- Time — Event time.
- Delta — Event type indicator.
- Baseline covariates
- Columns for each event type indicating cumulative event counts.

Examples

```
# The observed data
data_obs <- simDisease(N = 1000)
data_obs <- IntFormatData(data_obs, N_cols = 6)

# Fit Cox models
cox_death <- survival::coxph(survival::Surv(tstart, tstop, Delta == 1)
~ L0 + A0 + L, data = data_obs)
cox_Disease <- survival::coxph(survival::Surv(tstart, tstop, Delta == 2)
~ L0 + A0, data = data_obs[L == 0])

# Then simulate new data:
cox_fits <- list("D" = cox_death, "L" = cox_Disease)
list_old_vars <- list("L0" = data_obs$L0, "A0" = data_obs$A0)
new_data <- simEventCox(100, cox_fits = cox_fits, list_old_vars = list_old_vars)
```

simEventData

Simulate Event Data with Multiple Event Types and Covariates

Description

Simulate Continuous Time-to-Event Data with Multiple Event Types

Usage

```

simEventData(
  N,
  beta = NULL,
  eta = NULL,
  nu = NULL,
  at_risk = NULL,
  term_deltas = c(0, 1),
  max_cens = Inf,
  add_cov = NULL,
  override_beta = NULL,
  max_events = 10,
  lower = 10-15,
  upper = 200,
  gen_A0 = NULL,
  gen_L0 = NULL,
  at_risk_cov = NULL,
  ...
)

```

Arguments

N	Integer. Number of individuals to simulate.
beta	Numeric matrix. Regression coefficients matrix where columns correspond to event types (N0, N1, ...) and rows correspond to covariates (L0, A0, L1, L2, ...) and event counts (N0, N1, ...). Default is a zero matrix.
eta	Numeric vector. Shape parameters of the Weibull baseline intensity for each event type. Default is 0.1 for all events.
nu	Numeric vector. Scale parameters of the Weibull baseline intensity for each event type. Default is 1.1 for all events.
at_risk	Function. Function determining if an individual is at risk for each event type, given their current event counts. Takes a numeric vector events and returns a binary vector. Default returns 1 for all events.
term_deltas	Integer vector. Event types considered terminal (after which no further events occur). Default is c(0, 1).
max_cens	Numeric. Maximum censoring time. Events occurring after this time are censored. Default is Inf (no maximal censoring).
add_cov	Named list of functions. Functions generating additional baseline covariates. Each function takes integer N and returns a numeric vector of length N. Default is NULL.
override_beta	Named list. Used to specify entries of the beta matrix to override defaults. For example, list("L0" = c("N1" = 2)) sets the effect of L0 on N1 to 2.
max_events	Integer. Maximum number of events to simulate per individual. Default is 10.
lower	Numeric. Lower bound for root-finding in inverse cumulative hazard calculations. Default is 10 ⁻¹⁵ .

upper	Numeric. Upper bound for root-finding in inverse cumulative hazard calculations. Default is 200.
gen_A0	Function. Function to generate the baseline treatment covariate A0. Takes N and L0 as inputs. Default is a Bernoulli(0.5) random variable.
gen_L0	Function. Function to generate the baseline covariate L0. Takes N as inputs. Default is a N(0,1) random variable.
at_risk_cov	Function. Function determining if an individual is at risk for each event type, given their covariates. Takes a numeric vector covariates and returns a binary vector. Default returns 1 for all events.
...	Additional technical arguments

Details

simEventData simulates event times and types for a cohort of individuals in a counting process framework. It supports multiple event types (by default 4), including terminal events, with intensities influenced by baseline covariates and previous event history.

The event intensities for event type x at time t are given by

$$\lambda^x(t) = \lambda_0^x(t) \exp(\beta_x^T L),$$

where the baseline intensity follows a Weibull hazard function:

$$\lambda_0^x(t) = \eta^x \nu^x t^{\nu^x - 1}.$$

Here, L is the vector of covariates and event counts, and β^x is the a vector of coefficients representing the effect of covariates and previous events on the intensity.

Value

A data.table with columns:

ID	Individual identifier
Time	Time of event
Delta	Event type at time
L0	Baseline covariate
A0	Baseline treatment
L1, L2, ...	Additional baseline covariates if specified
N0, N1, ...	Event counts up to the current event

Examples

```
# Simulate data for 10 individuals with default settings
sim_data <- simEventData(N = 10)
head(sim_data)
```

simEventObj	<i>Simulate Survival and Competing Risk Data Based on a General Model</i>
-------------	---

Description

The `simEventObj` function simulates survival or competing risk data for a cohort of individuals based on a general model with a `predict2` method. The function is useful for simulating additional data under the same distribution as an original data set. The procedure consists of fitting a model, such as a random forest or Cox Proportional Hazards model on an original data set. Next the model is equipped with a `predict2` method, and passed as an argument to the `simEventObj` function, which simulates new data using the `predict2` method. The method should output the cumulative hazard array and the jump times of the cumulative hazard. Simulation proceeds by sampling from the uniform distribution and obtaining event times using the inverse of the cumulative hazard function(s).

Usage

```
simEventObj(N, obj, event_names = NULL, list_old_vars = NULL)
```

Arguments

<code>N</code>	Integer. The number of individuals to simulate.
<code>obj</code>	An object of class <code>simevent</code> . The object should have a <code>predict2</code> method. The method should return a list containing <code>chf</code> and <code>time</code> . <code>chf</code> should be an array of dimension <code>Individuals x Times x Events</code> . The array should contain the cumulative hazard values for each individual, at each time for each event. <code>time</code> should be a vector of times where the cumulative hazard function jumps.
<code>event_names</code>	A character vector. Containing the names of the various processes. The argument is optional. By default events will be named <code>N1, N2, ...</code>
<code>list_old_vars</code>	A named list containing the old covariates. New covariates will be simulated by drawing from the old covariates with replacement.

Details

The function simulates individual event histories by:

1. Sampling initial baseline covariates by resampling observed values.
2. Extracting cumulative hazard functions from the object.
3. Iteratively sampling event times.
4. Updating covariate histories and event counts.
5. Stopping simulation per individual after a terminal event or maximum events reached.

Value

A data.table with one row per event per individual containing:

- ID — Individual identifier.
- Time — Event time.
- Delta — Event type indicator.
- Baseline covariates.
- Columns for each event type indicating cumulative event counts.

 simEventTV

simEventTV

Description

Simulate Event Data with Time-Varying Effects

Usage

```
simEventTV(
  N,
  beta = NULL,
  tv_eff = NULL,
  t_prime = Inf,
  eta = NULL,
  nu = NULL,
  at_risk = NULL,
  term_deltas = c(0, 1),
  max_cens = Inf,
  add_cov = NULL,
  override_beta = NULL,
  max_events = 10,
  lower = 10-15,
  upper = 200,
  gen_A0 = NULL,
  at_risk_cov = NULL
)
```

Arguments

N	Integer. Number of individuals to simulate.
beta	Matrix. Coefficients for covariates and processes. Columns correspond to events (N0, N1, ...), rows correspond to covariates (L0, A0, ..., and past event counts).
tv_eff	Matrix. Time-varying changes to beta, applied at time t_prime. Must have same dimensions as beta.
t_prime	Numeric. Time at which tv_eff is added to beta.

eta	Numeric vector. Shape parameters of Weibull intensities for each event.
nu	Numeric vector. Scale parameters of Weibull intensities for each event.
at_risk	Function. Determines which events an individual is at risk for, based on event history.
term_deltas	Integer vector. Event types considered terminal (e.g., death).
max_cens	Numeric. Maximum censoring time. Defaults to Inf.
add_cov	Named list of functions for generating additional baseline covariates. Each function takes one argument N and returns a vector of length N.
override_beta	Named list to override elements of beta. Format: <code>list("covariate" = c("event" = value))</code> .
max_events	Integer. Maximum number of events allowed per individual.
lower	Numeric. Lower bound for the root-finding algorithm used in inverse cumulative hazard computation.
upper	Numeric. Upper bound for the root-finding algorithm used in inverse cumulative hazard computation.
gen_A0	Function. Generates baseline treatment assignment. Takes arguments N and L0.
at_risk_cov	Function. Function determining if an individual is at risk for each event type, given their covariates. Takes a numeric vector covariates and returns a binary vector. Default returns 1 for all events.

Details

simEventTV is a function that simulates event data, with the option of adding time varying effects. The function is build up in the same way as simEventData, with the additional arguments tv_eff and t_prime, which specify the change of the beta matrix at time t_prime.

Value

A data.table with columns:

ID:	Individual identifier
Time:	Time of event
Delta:	Type of event
L0:	Baseline covariate
A0:	Baseline treatment
N0, N1, ...:	Cumulative event counts
L1, L2, ...:	Additional covariates (if specified)

Examples

```
eta <- rep(0.1, 2)
simEventTV(N = 100, t_prime = 1, eta = eta, term_deltas = c(0, 1))
```

simStatinData *Simulate Data in a Statin Setting*

Description

Simulate Data in a Statin Setting

Usage

```
simStatinData(
  N,
  eta = rep(0.1, 6),
  nu = rep(1.1, 6),
  beta = NULL,
  cens = 1,
  followup = 5,
  lower = 10^(-15),
  upper = 200,
  gen_A0 = NULL,
  gen_L0 = NULL,
  ...
)
```

Arguments

N	Numeric scalar. Number of individuals to simulate.
eta	Numeric vector of length 6. Shape parameters for Weibull intensities with parameterization $\eta\nu t^{\nu-1}$. Defaults to rep(0.1, 6).
nu	Numeric vector of length 6. Scale parameters for the Weibull hazards. Defaults to rep(1.1, 6).
beta	Numeric matrix. Of dimension p times 6. Regression coefficients matrix where columns correspond to event types (N0, ..., N5) and rows correspond to covariates (L0, A0, L1, L2, ...) followed by event counts (N0, ..., N5). Default is a zero matrix.
cens	Binary scalar. Indicates whether individuals are at risk of censoring (default 1).
followup	Numeric scalar. Maximum follow-up (censoring) time. Defaults to Inf.
lower	Numeric scalar. Lower bound for root-finding (inverse cumulative hazard) (default 1e-15).
upper	Numeric scalar. Upper bound for root-finding (default 200).
gen_A0	Function. Function to generate the baseline treatment covariate A0. Takes N and L0 as inputs. Default is a Bernoulli(0.5) random variable.
gen_L0	Function. Function to generate the baseline covariate L0. Takes N as input. Default is a N(0,1) random variable.
...	Additional arguments passed to simEventData

Value

A data frame containing the simulated data with columns:

ID	Individual identifier
Time	Time of the event
Delta	Event type (0,...,5)
L0	Baseline covariate
A0	Baseline covariate
L1, ... Lp	Additional baseline covariates

Examples

```
simDisease(10)
```

simSurvData

Simulate Survival Data

Description

Simulate Survival Data with Censoring and Event Times

Usage

```
simSurvData(N, beta = NULL, eta = rep(0.1, 2), nu = rep(1.1, 2), cens = 1, ...)
```

Arguments

N	Numeric scalar. Number of individuals to simulate.
beta	Numeric 2x2 matrix specifying effects of baseline covariates L0 and A0 on censoring and event hazards. <ul style="list-style-type: none"> • Rows correspond to covariates L0 and A0. • Columns correspond to censoring (1st column) and event (2nd column). Defaults to zero matrix if NULL.
eta	Numeric vector of length 2. Shape parameters for Weibull hazard with parameterization $\eta\nu t^{\nu-1}$. Defaults to rep(0.1, 2).
nu	Numeric vector of length 2. Scale parameters for the Weibull hazard. Defaults to rep(1.1, 2).
cens	Numeric binary indicator (0 or 1) specifying if censoring is included (default 1).
...	Additional arguments passed to simEventData, including the argument add_cov to specify extra covariates.

Details

Simulates survival data for N individuals who are at risk for censoring (0) and an event (1). The hazard functions for censoring and event times follow Weibull distributions parameterized by shape parameters η and scale parameters ν . Covariate effects on censoring and event hazards are specified via a matrix beta.

Value

Data frame containing the simulated survival data

Examples

```
simSurvData(10)
```

simTreatment	<i>Simulate Event History Data with Treatment and Time-Dependent Covariate</i>
--------------	--

Description

Simulates event history data with four types of events representing censoring (0), death (1), treatment (2), and covariate change (3). Death and censoring are terminal events; treatment and covariate events can occur only once.

Usage

```
simTreatment(  
  N,  
  eta = rep(0.1, 4),  
  nu = rep(1.1, 4),  
  beta_L_A = 1,  
  beta_L_D = 1,  
  beta_A_D = -1,  
  beta_A_L = -0.5,  
  beta_L0_A = 1,  
  beta_L0_L = 1,  
  beta_L0_D = 1,  
  beta_L0_C = 0,  
  beta_L_C = 0,  
  beta_A_C = 0,  
  beta_L_A_prime = 0,  
  beta_L_D_prime = 0,  
  beta_A_D_prime = 0,  
  beta_A_L_prime = 0,  
  beta_L0_A_prime = 0,  
  beta_L0_L_prime = 0,
```

```

beta_L0_D_prime = 0,
beta_L0_C_prime = 0,
beta_L_C_prime = 0,
beta_A_C_prime = 0,
t_prime = NULL,
at_risk_cov = NULL,
cens = 1,
op = 1,
lower = 10^(-15),
upper = 200,
followup = Inf,
...
)

```

Arguments

N	Integer. Number of individuals to simulate.
eta	Numeric vector of length 4. Shape parameters for Weibull intensities, parameterized as $\eta vt^{\nu-1}$. Default is rep(0.1, 4).
nu	Numeric vector of length 4. Scale parameters for Weibull hazards. Default is rep(1.1, 4).
beta_L_A	Numeric. Effect of covariate L = 1 on treatment hazard. Default 1.
beta_L_D	Numeric. Effect of covariate L = 1 on death hazard. Default 1.
beta_A_D	Numeric. Effect of treatment A = 1 on death hazard. Default -1.
beta_A_L	Numeric. Effect of treatment A = 1 on covariate hazard. Default -0.5.
beta_L0_A	Numeric. Effect of baseline covariate L0 on treatment hazard. Default 1.
beta_L0_L	Numeric. Effect of baseline covariate L0 on covariate hazard. Default 1.
beta_L0_D	Numeric. Effect of baseline covariate L0 on death hazard. Default 1.
beta_L0_C	Numeric. Effect of baseline covariate L0 on censoring hazard. Default 0.
beta_L_C	Numeric. Effect of covariate L = 1 on censoring hazard. Default 0.
beta_A_C	Numeric. Effect of treatment A = 1 on censoring hazard. Default 0.
beta_L_A_prime	Numeric. Additional effect of covariate L = 1 on treatment hazard. Default 0.
beta_L_D_prime	Numeric. Additional effect of covariate L = 1 on death hazard. Default 0.
beta_A_D_prime	Numeric. Effect of treatment A = 1 on death hazard. Default 0.
beta_A_L_prime	Numeric. Effect of treatment A = 1 on covariate hazard. Default 0.
beta_L0_A_prime	Numeric. Effect of baseline covariate L0 on treatment hazard. Default 0.
beta_L0_L_prime	Numeric. Effect of baseline covariate L0 on covariate hazard. Default 0.
beta_L0_D_prime	Numeric. Effect of baseline covariate L0 on death hazard. Default 0.
beta_L0_C_prime	Numeric. Effect of baseline covariate L0 on censoring hazard. Default 0.

beta_L_C_prime	Numeric. Effect of covariate L = 1 on censoring hazard. Default 0.
beta_A_C_prime	Numeric. Effect of treatment A = 1 on censoring hazard. Default 0.
t_prime	Numeric scalar or NULL. Time point where effects change (optional).
at_risk_cov	Function. Function determining if an individual is at risk for each event type, given their covariates. Takes a numeric vector covariates and returns a binary vector. Default returns 1 for all events.
cens	Integer (0 or 1). Indicates if censoring is possible. Default 1.
op	Integer (0 or 1). Indicates if treatment (operation) is possible. Default 1.
lower	Numeric. Lower bound for root finding (inverse cumulative hazard). Default 1e-15.
upper	Numeric. Upper bound for root finding (inverse cumulative hazard). Default 200.
followup	Numeric. Maximum censoring time. Defaults to Inf (no censoring).
...	Additional arguments passed to simEventData or simEventTV

Details

Event intensities are modeled using Weibull hazards with parameters ν (scale) and η (shape), and covariate effects controlled by specified beta parameters. For example, beta_L_A quantifies the effect of covariate L = 1 on the hazard of treatment.

Value

A data.frame with columns:

- ID - Individual identifier.
- Time - Event time.
- Delta - Event type (0=censoring, 1=death, 2=treatment, 3=covariate change).
- L0 - Baseline covariate.
- L - Time-dependent covariate.
- A - Treatment status.

Examples

```
simTreatment(10)
```

Index

[alphaSim](#), [2](#)

[intEffectAlpha](#), [3](#)

[IntFormatData](#), [5](#)

[inverseScHaz](#), [6](#)

[inverseScHazTV](#), [7](#)

[plotEventData](#), [8](#)

[sampleEvents](#), [9](#)

[simCRdata](#), [9](#)

[simDisease](#), [10](#)

[simDropIn](#), [12](#)

[simEventCox](#), [16](#)

[simEventData](#), [9](#), [17](#)

[simEventObj](#), [20](#)

[simEventTV](#), [21](#)

[simStatinData](#), [23](#)

[simSurvData](#), [24](#)

[simTreatment](#), [25](#)