

Package ‘HazReg’

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Type Package

Title Parametric hazard-based regression models

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Description The HazReg R package implements the following parametric hazard-based regression models for survival data, in the overall and relative survival frameworks.

License What license is it under?

Encoding UTF-8

LazyData true

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Imports numDeriv, matrixStats

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chew	<i>Power Exponentiated Weibull (EW) cumulative hazard function.</i> https://rpubs.com/FJRUBIO/EWD
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Description

Power Exponentiated Weibull (EW) cumulative hazard function. <https://rpubs.com/FJRUBIO/EWD>

Usage

```
chew(t, sigma, nu, gamma)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter

Value

the value of the EW cumulative hazard function

chgammma	<i>Gamma (G) cumulative hazard function.</i>
----------	--

Description

Gamma (G) cumulative hazard function.

Usage

```
chgammma(t, shape, scale)
```

Arguments

t	: positive argument
shape	: shape parameter
scale	: scale parameter
log:	log scale (TRUE or FALSE)

Value

the value of the Weibull hazard function

chggamma

Generalised Gamma (GG) cumulative hazard function.
<https://rpubs.com/FJRubio/GG>

Description

Generalised Gamma (GG) cumulative hazard function. <https://rpubs.com/FJRubio/GG>

Usage

```
chggamma(t, sigma, nu, gamma)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter

Value

the value of the GG cumulative hazard function

chllogis

Log-logistic (LL) cumulative hazard function.

Description

Log-logistic (LL) cumulative hazard function.

Usage

```
chllogis(t, mu, sigma)
```

Arguments

t	: positive argument
mu	: mean parameter in the log scale
sigma	: scale parameter in the log scale

Value

the value of the LL cumulative hazard function

chlnorm

Lognormal (LN) cumulative hazard function.

Description

Lognormal (LN) cumulative hazard function.

Usage

```
chlnorm(t, mu, sigma)
```

Arguments

- | | |
|-------|------------------------------------|
| t | : positive argument |
| mu | : mean parameter in the log scale |
| sigma | : scale parameter in the log scale |

Value

the value of the LN cumulative hazard function

chpgw

Power Generalised Weibull (PGW) cumulative hazard function.
<http://rpubs.com/FJRubio/PGW>

Description

Power Generalised Weibull (PGW) cumulative hazard function. <http://rpubs.com/FJRubio/PGW>

Usage

```
chpgw(t, sigma, nu, gamma)
```

Arguments

- | | |
|-------|---------------------|
| t | : positive argument |
| sigma | : scale parameter |
| nu | : shape parameter |
| gamma | : shape parameter |

Value

the value of the PGW cumulative hazard function

chweibull	<i>Weibull (W) cumulative hazard function.</i>
-----------	--

Description

Weibull (W) cumulative hazard function.

Usage

```
chweibull(t, sigma, nu)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter

Value

the value of the Weibull cumulative hazard function

CH_TVC	<i>Compute the Cumulative Hazard for a Proportional Hazards or Accelerated Failure Model (2- and 3-parameter baseline)</i>
--------	--

Description

Computes the cumulative hazard $H(t | x(t))$ at multiple time points for each individual under a proportional hazards (PH) model or Accelerated Failure Time (AFT) model with a two-parameter or three-parameter parametric baseline hazard.

Usage

```
CH_TVC(df, beta, npar, ae0, be0, ce0 = NULL, chfun, hstr)
```

Arguments

df	A data frame containing:
	<ul style="list-style-type: none"> • ‘time’: numeric vector of time points. • Covariate columns named with prefix “des” (e.g., ‘des1’, ‘des2’, ...), representing $x(t)$.
beta	Numeric vector of regression coefficients.
npar	Number of parameters in the baseline hazard (2 or 3).
ae0, be0, ce0	Numeric baseline parameters of the cumulative hazard.
chfun	A function computing the baseline cumulative hazard: ‘chfun(time, ae0, be0, ce0)’.
hstr	Hazard structure ("PH" or "AFT")

Details

The PH model assumes

$$H(t \mid x(t)) = H_0(t; a_0, b_0, c_0) \exp(x(t)^\top \beta).$$

In the AFT model, event time is rescaled as

$$H(t \mid x(t)) = H_0(t \exp(x(t)^\top \beta); a_0, b_0, c_0).$$

Value

A numeric vector with the cumulative hazard evaluated at each time point in ‘df’.

Conf_Int

Function to calculate the normal confidence intervals. The parameters indicated with "index" are transformed to the real line using log().

Description

Function to calculate the normal confidence intervals. The parameters indicated with "index" are transformed to the real line using log().

Usage

```
Conf_Int(FUN, MLE, level = 0.95, index = NULL)
```

Arguments

- FUN : minus log-likelihood function to be used to calculate the confidence intervals
- MLE : maximum likelihood estimator of the parameters of interest
- level : confidence level
- index : position of the positive parameters under the original parameterisation

Value

a list containing the upper and lower conf.int limits, the transformed MLE, and std errors

dggamma *Generalised Gamma (GG) probability density function.*
<https://rpubs.com/FJRubio/GG>

Description

Generalised Gamma (GG) probability density function. <https://rpubs.com/FJRubio/GG>

Usage

```
dggamma(t, sigma, nu, gamma, log = FALSE)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter
log:	log scale (TRUE or FALSE)

Value

the value of the GG probability density function

dpgw *Power Generalised Weibull (PGW) probability density function.*
<http://rpubs.com/FJRubio/PGW>

Description

Power Generalised Weibull (PGW) probability density function. <http://rpubs.com/FJRubio/PGW>

Usage

```
dpgw(t, sigma, nu, gamma, log = FALSE)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter
log:	log scale (TRUE or FALSE)

Value

the value of the PGW probability density function

GEHMLE

Relative (Net) Survival models Log likelihood and MLE for the GH excess hazards model. Baseline hazards: Lognormal, Log-logistic, Gamma, PGW, EW, GG

Description

Relative (Net) Survival models Log likelihood and MLE for the GH excess hazards model. Baseline hazards: Lognormal, Log-logistic, Gamma, PGW, EW, GG

Usage

```
GEHMLE(
  init,
  times,
  status,
  hp,
  hstr = NULL,
  dist = NULL,
  des = NULL,
  des_t = NULL,
  method = "Nelder-Mead",
  maxit = 100
)
```

Arguments

init	: initial point for optimisation step under the parameterisation (log(scale), log(shape1), log(shape2), alpha, beta) for scale-shape1-shape2 models or (mu, log(scale), alpha, beta) for log-location scale models.
times	: times to event
status	: vital status indicators (TRUE or 1 = observed, FALSE or 0 = censored)
hp	: population hazard (for all individuals)
hstr	: hazard structure: No covariates ("baseline"), AFT model with PGW baseline hazard ("AFT"), PH model with PGW baseline hazard ("PH"), AH model with PGW baseline hazard ("AH"), GH model with PGW baseline hazard ("GH") *GH is not available with Weibull dist
dist	: distribution for the baseline hazard: Power Generalised Weibull ("PGW") Generalised Gamma ("GenGamma")) Exponentiated Weibull ("EW") Weibull ("Weibull") Gamma ("Gamma") LogNormal ("LogNormal") LogLogistic ("LogLogistic")
des	: design matrix for hazard-level effects
des_t	: design matrix for time-level effects (it is recommended not to use splines here)
method	: "nlminb" or optimisation method to be used in optim (see ?optim)
maxit	: maximum number of iterations in optim or nlminb

Value

It returns the output from optim or nlminb for the selected model and the negative log likelihood function

GHMLE	<i>GHMLE function: Hazard Regression Models with a parametric baseline hazard</i>
-------	---

Description

GHMLE function: Hazard Regression Models with a parametric baseline hazard

Usage

```
GHMLE(
  init,
  times,
  status,
  hstr = NULL,
  dist = NULL,
  des = NULL,
  des_t = NULL,
  method = "Nelder-Mead",
  maxit = 100
)
```

Arguments

init	: initial point for optimisation step under the parameterisation (log(scale), log(shape1), log(shape2), alpha, beta) for scale-shape1-shape2 models or (mu, log(scale), alpha, beta) for log-location scale models.
times	: times to event
status	: vital status indicators (TRUE or 1 = observed, FALSE or 0 = censored)
hstr	: hazard structure: No covariates ("baseline"), AFT model with PGW baseline hazard ("AFT"), PH model with PGW baseline hazard ("PH"), AH model with PGW baseline hazard ("AH"), GH model with PGW baseline hazard ("GH") *GH is not available with Weibull dist
dist	: distribution for the baseline hazard: Power Generalised Weibull ("PGW") Generalised Gamma ("GenGamma") Exponentiated Weibull ("EW") Weibull ("Weibull") Gamma ("Gamma") LogNormal ("LogNormal") LogLogistic ("LogLogistic")
des	: design matrix for hazard-level effects
des_t	: design matrix for time-level effects (it is recommended not to use splines here)
method	: "nlminb" or optimisation method to be used in optim (see ?optim)
maxit	: maximum number of iterations in optim or nlminb

Value

It returns the output from optim or nlminb for the selected model and the negative log likelihood function

hew *Power Exponentiated Weibull (EW) hazard function.*
<https://rpubs.com/FJRubio/EWD>

Description

Power Exponentiated Weibull (EW) hazard function. <https://rpubs.com/FJRubio/EWD>

Usage

```
hew(t, sigma, nu, gamma, log = FALSE)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter
log:	log scale (TRUE or FALSE)

Value

the value of the EW hazard function

hgamma *Gamma (G) hazard function.*

Description

Gamma (G) hazard function.

Usage

```
hgmma(t, shape, scale, log = FALSE)
```

Arguments

t	: positive argument
shape	: shape parameter
scale	: scale parameter
log:	log scale (TRUE or FALSE)

Value

the value of the Gamma hazard function

hggamma	<i>Generalised Gamma (GG) hazard function.</i>
	https://rpubs.com/FJRubio/GG

Description

Generalised Gamma (GG) hazard function. <https://rpubs.com/FJRubio/GG>

Usage

```
hggamma(t, sigma, nu, gamma, log = FALSE)
```

Arguments

- t : positive argument
- sigma : scale parameter
- nu : shape parameter
- gamma : shape parameter
- log: log scale (TRUE or FALSE)

Value

the value of the GG hazard function

hllogis	<i>Log-logistic (LL) hazard function.</i>
---------	---

Description

Log-logistic (LL) hazard function.

Usage

```
hllogis(t, mu, sigma, log = FALSE)
```

Arguments

- t : positive argument
- mu : mean parameter in the log scale
- sigma : scale parameter in the log scale
- log: log scale (TRUE or FALSE)

Value

the value of the LL hazard function

hlnorm	<i>Lognormal (LN) hazard function.</i>
--------	--

Description

Lognormal (LN) hazard function.

Usage

```
hlnorm(t, mu, sigma, log = FALSE)
```

Arguments

t	: positive argument
mu	: mean parameter in the log scale
sigma	: scale parameter in the log scale
log:	log scale (TRUE or FALSE)

Value

the value of the LN hazard function

HMLE_TVC	<i>Maximum Likelihood Estimation for Parametric Hazard Models with Time-Varying Covariates</i>
----------	--

Description

'HMLE_TVC()' fits parametric survival models in the presence of **time-varying covariates**, using maximum likelihood estimation.

The function supports:

- * **Proportional Hazards (PH)** models with time-varying covariates
- * Fully parametric baseline hazards (2-parameter or 3-parameter)

The likelihood is constructed from the cumulative hazard differences across observation intervals for each individual, using a counting-process representation.

For each individual, the data must contain several rows: one per time-varying covariate measurement, along with the corresponding time.

Usage

```
HMLE_TVC(
  init,
  df,
  status,
  hstr = NULL,
  dist = NULL,
  des = NULL,
  method = "Nelder-Mead",
  maxit = 100
)
```

Arguments

df	A data frame in **long format**, containing one row per individual per covariate-measurement time. Required columns: * ‘ID’ — individual identifier * ‘time’ — time at which the covariates are measured * ‘status’ — event indicator (1 = event at the final time; 0 = censored) * ‘des*’ — covariate columns used in the model (e.g., ‘des1’, ‘des2’, …) The last row for each ID represents the individual’s event/censoring time, even if the event time does not coincide with a measurement time.
method	Optimisation method for the likelihood. Either ““nlminb”“ or a valid ‘optim()‘ method.
maxit	Maximum number of optimisation iterations.
beta	Numeric vector of regression coefficients associated with the time-varying covariate design matrix (‘des*’ columns).
ae0	be0, ce0 Baseline hazard parameters. * For **2-parameter** baselines, only ‘ae0’ and ‘be0’ are used. * For **3-parameter** baselines, all three are used. These parameters are passed directly to the user-supplied baseline cumulative hazard function ‘chfun()‘.
chfun	A function computing the **baseline cumulative hazard**: * 2-parameter case: ‘chfun(time, ae0, be0)‘ * 3-parameter case: ‘chfun(time, ae0, be0, ce0)‘ The function must return a vector of values of equal length to ‘time‘.

Details

Likelihood formulation

For each individual $\backslash(i\backslash)$, let $\backslash(t_i1 < t_i2 < \dots < t_iK_i\backslash)$ denote the *observation / measurement times*.

The cumulative hazard contribution over interval $\backslash((t_ij-1, t_ij)\backslash)$ is:

$$\Delta H_{ij} = [H_0(t_{ij}) - H_0(t_{ij-1})] \exp(x_{ij}^\top \beta),$$

where $\backslash(x_ij\backslash)$ is the vector of covariates measured at time $\backslash(t_ij\backslash)$.

The full log-likelihood is:

$$\ell = \sum_i \left(- \sum_j \Delta H_{ij} + \delta_i \log [h_0(T_i) \exp(x_{iK}^\top \beta)] \right),$$

where:

* $\backslash(\Delta H_{ij}\backslash)$ comes from cumulative hazard increments * $\backslash(T_i = t_iK\backslash)$ is the final event or censoring time * $\backslash(\delta_i\backslash)$ is the event indicator * hazard and cumulative hazard are computed from ‘chfun()‘

The function internally: 1. Splits the data by ID 2. Computes cumulative hazard at all measurement times 3. Computes increments $\backslash(\Delta H_{ij}\backslash)$ for each ID 4. Constructs the likelihood 5. Optimises over $\backslash(\beta\backslash)$ and baseline parameters

Value

A list containing:

- * The full output from ‘optim()‘ or ‘nlminb()‘
 - * The **negative log-likelihood function** used for optimisation
 - * A vector giving, for each ID, the cumulative hazard increments used in the likelihood
- Returned invisibly where appropriate.

Data structure

The input data frame must contain:

- * varying number of rows per ID
- * strictly increasing ‘time‘ within each ID
- * last row containing the event/censoring time

Covariates must be named as ‘des1‘, ‘des2‘, etc.

hpgw	<i>Power Generalised Weibull (PGW) hazard function.</i>
	<i>http://rpubs.com/FJRubio/PGW</i>

Description

Power Generalised Weibull (PGW) hazard function. <http://rpubs.com/FJRubio/PGW>

Usage

```
hpgw(t, sigma, nu, gamma, log = FALSE)
```

Arguments

- | | |
|--------------------|---------------------------|
| <code>t</code> | : positive argument |
| <code>sigma</code> | : scale parameter |
| <code>nu</code> | : shape parameter |
| <code>gamma</code> | : shape parameter |
| <code>log:</code> | log scale (TRUE or FALSE) |

Value

the value of the PGW hazard function

hweibull

*Weibull (W) hazard function.***Description**

Weibull (W) hazard function.

Usage

```
hweibull(t, sigma, nu, log = FALSE)
```

Arguments

- t : positive argument
- sigma : scale parameter
- nu : shape parameter
- log: log scale (TRUE or FALSE)

Value

the value of the Weibull hazard function

pgamma

Generalised Gamma (GG) cumulative distribution function.
<https://rpubs.com/FJRubi/GG>

Description

Generalised Gamma (GG) cumulative distribution function. <https://rpubs.com/FJRubi/GG>

Usage

```
pgamma(t, sigma, nu, gamma, log.p = FALSE)
```

Arguments

- t : positive argument
- sigma : scale parameter
- nu : shape parameter
- gamma : shape parameter
- log.p: log scale (TRUE or FALSE)

Value

the value of the GG cumulative distribution function

qew

Power Exponentiated Weibull (EW) quantile function.
<https://rpubs.com/FJRubio/EWD>

Description

Power Exponentiated Weibull (EW) quantile function. <https://rpubs.com/FJRubio/EWD>

Usage

```
qew(p, sigma, nu, gamma)
```

Arguments

- p : probability. A value in (0,1)
- sigma : scale parameter
- nu : shape parameter
- gamma : shape parameter

Value

the value of the EW quantile function

qggamma

Generalised Gamma (GG) quantile function.
<https://rpubs.com/FJRubio/GG>

Description

Generalised Gamma (GG) quantile function. <https://rpubs.com/FJRubio/GG>

Usage

```
qggamma(p, sigma, nu, gamma)
```

Arguments

- p : probability. A value in (0,1)
- sigma : scale parameter
- nu : shape parameter
- gamma : shape parameter

Value

the value of the GG quantile function

qlllogis *Log-logistic (LL) quantile function.*

Description

Log-logistic (LL) quantile function.

Usage

```
qlllogis(p, mu, sigma)
```

Arguments

- p : probability. A value in (0,1)
- mu : mean parameter in the log scale
- sigma : scale parameter in the log scale

Value

the value of the LL quantile function

qpgw *Power Generalised Weibull (PGW) quantile function.*
<http://rpubs.com/FJRubi/PGW>

Description

Power Generalised Weibull (PGW) quantile function. <http://rpubs.com/FJRubi/PGW>

Usage

```
qpgw(p, sigma, nu, gamma)
```

Arguments

- p : probability. A value in (0,1)
- sigma : scale parameter
- nu : shape parameter
- gamma : shape parameter

Value

the value of the PGW quantile function

rggamma*Generalised Gamma (GG) random number generation.*
<https://rpubs.com/FJRubio/GG>

Description

Generalised Gamma (GG) random number generation. <https://rpubs.com/FJRubio/GG>

Usage

```
rggamma(n, sigma, nu, gamma)
```

Arguments

n	: number of observations
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter

Value

generates random deviates

rpgw*Power Generalised Weibull (PGW) random number generation.*
<http://rpubs.com/FJRubio/PGW>

Description

Power Generalised Weibull (PGW) random number generation. <http://rpubs.com/FJRubio/PGW>

Usage

```
rpgw(n, sigma, nu, gamma)
```

Arguments

n	: number of observations
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter

Value

generates random deviates

sggamma	<i>Generalised Gamma (GG) survival function.</i>
	https://rpubs.com/FJRubio/GG

Description

Generalised Gamma (GG) survival function. <https://rpubs.com/FJRubio/GG>

Usage

```
sggamma(t, sigma, nu, gamma, log.p = FALSE)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter
log.p:	log scale (TRUE or FALSE)

Value

the value of the GG survival function

simGH	<i>simGH function: Function to simulate times to event from a model with a GH structure for different parametric baseline hazards. Distributions: LogNormal, LogLogistic, GenGamma, Gamma, Weibull, PGW, EW. See: https://github.com/FJRubio67/HazReg</i>
-------	---

Description

simGH function: Function to simulate times to event from a model with a GH structure for different parametric baseline hazards. Distributions: LogNormal, LogLogistic, GenGamma, Gamma, Weibull, PGW, EW. See: <https://github.com/FJRubio67/HazReg>

Usage

```
simGH(
  seed,
  n,
  des = NULL,
  des_h = NULL,
  des_t = NULL,
  theta,
  beta_h = NULL,
  beta_t = NULL,
  beta = NULL,
  hstr,
  baseline
)
```

Arguments

seed	: seed for simulation
n	: sample size (number of individuals)
des	: Design matrix for AFT, PH, and AH models
des_h	: Design matrix for GH model (hazard scale)
des_t	: Design matrix for GH model (time scale)
theta	: parameters of the baseline hazard
beta_h	: regression parameters multiplying the hazard for GH model
beta_t	: regression parameters multiplying the time scale for GH model
beta	: regression parameters for AFT, PH, and AH models
hstr	: hazard structure (AH, AFT, PH, GH)
baseline	: baseline hazard distribution

Value

a vector containing the simulated times to event

spgw *Power Generalised Weibull (PGW) survival function.*
<http://rpubs.com/FJRubio/PGW>

Description

Power Generalised Weibull (PGW) survival function. <http://rpubs.com/FJRubio/PGW>

Usage

```
spgw(t, sigma, nu, gamma, log.p = FALSE)
```

Arguments

t	: positive argument
sigma	: scale parameter
nu	: shape parameter
gamma	: shape parameter
log.p:	log scale (TRUE or FALSE)

Value

the value of the PGW survival function

SPred_TVC	<i>Compute the Survival Function for a Proportional Hazards or Accelerated Failure Model (2- and 3-parameter baseline)</i>
-----------	--

Description

Computes the survival function $\exp(-H(t | x(t)))$ at the last time point for each individual under a proportional hazards (PH) model or Accelerated Failure Time (AFT) model with a two-parameter or three-parameter parametric baseline hazard.

Usage

```
SPred_TVC(df, beta, npar, ae0, be0, ce0 = NULL, chfun, hstr)
```

Arguments

df	A data frame containing: <ul style="list-style-type: none"> • ‘time’: numeric vector of time points. • Covariate columns named with prefix “des” (e.g., ‘des1’, ‘des2’, ...), representing $x(t)$.
beta	Numeric vector of regression coefficients.
npar	Number of parameters in the baseline hazard (2 or 3).
ae0, be0, ce0	Numeric baseline parameters of the cumulative hazard.
chfun	A function computing the baseline cumulative hazard: ‘chfun(time, ae0, be0, ce0)’.
hstr	Hazard structure ("PH" or "AFT")

Details

The PH model assumes

$$H(t | x(t)) = H_0(t; a_0, b_0, c_0) \exp(x(t)^\top \beta).$$

In the AFT model, event time is rescaled as

$$H(t | x(t)) = H_0(t \exp(x(t)^\top \beta); a_0, b_0, c_0).$$

Value

A numeric vector with the survival function evaluated at the last time point in ‘df’.

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