

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.

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Imports numDeriv, matrixStats

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

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

Description

Gamma (G) cumulative hazard function.

Usage

```
chgamma(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	<i>Generalised Gamma (GG) cumulative hazard function.</i> <i>https://rpubs.com/FJRubio/GG</i>
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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	<i>Log-logistic (LL) cumulative hazard function.</i>
----------	--

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

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	<i>Power Generalised Weibull (PGW) cumulative hazard function.</i> <i>http://rpubs.com/FJRubio/PGW</i>
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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 $H(t \mid x(t))$ 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	<i>Function to calculate the normal confidence intervals. The parameters indicated with "index" are transformed to the real line using log().</i>
----------	---

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

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	<i>Power Generalised Weibull (PGW) probability density function.</i> <i>http://rpubs.com/FJRubio/PGW</i>
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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	<i>Relative (Net) Survival models Log likelihood and MLE for the GH excess hazards model. Baseline hazards: Lognormal, Log-logistic, Gamma, PGW, EW, GG</i>
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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

<code>init</code>	: 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.
<code>times</code>	: times to event
<code>status</code>	: vital status indicators (TRUE or 1 = observed, FALSE or 0 = censored)
<code>hp</code>	: population hazard (for all individuals)
<code>hstr</code>	: 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
<code>dist</code>	: distribution for the baseline hazard: Power Generalised Weibull ("PGW") Generalised Gamma ("GenGamma") Exponentiated Weibull ("EW") Weibull ("Weibull") Gamma ("Gamma") LogNormal ("LogNormal") LogLogistic ("LogLogistic")
<code>des</code>	: design matrix for hazard-level effects
<code>des_t</code>	: design matrix for time-level effects (it is recommended not to use splines here)
<code>method</code>	: "nlminb" or optimisation method to be used in optim (see ?optim)
<code>maxit</code>	: 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

*GHMLE function: Hazard Regression Models with a parametric baseline hazard***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

<code>init</code>	: 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.
<code>times</code>	: times to event
<code>status</code>	: vital status indicators (TRUE or 1 = observed, FALSE or 0 = censored)
<code>hstr</code>	: 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
<code>dist</code>	: distribution for the baseline hazard: Power Generalised Weibull ("PGW") Generalised Gamma ("GenGamma") Exponentiated Weibull ("EW") Weibull ("Weibull") Gamma ("Gamma") LogNormal ("LogNormal") LogLogistic ("LogLogistic")
<code>des</code>	: design matrix for hazard-level effects
<code>des_t</code>	: design matrix for time-level effects (it is recommended not to use splines here)
<code>method</code>	: "nlminb" or optimisation method to be used in optim (see ?optim)
<code>maxit</code>	: 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	<i>Power Exponentiated Weibull (EW) hazard function.</i> <i>https://rpubs.com/FJRubio/EWD</i>
-----	--

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	<i>Gamma (G) hazard function.</i>
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Description

Gamma (G) hazard function.

Usage

```
hgamma(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	<p>A data frame in long format, containing one row per individual per covariate-measurement time. Required columns:</p> <ul style="list-style-type: none"> * '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', ...) <p>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.</p>
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	<p>be0, ce0 Baseline hazard parameters.</p> <ul style="list-style-type: none"> * For 2-parameter baselines, only 'ae0' and 'be0' are used. * For 3-parameter baselines, all three are used. <p>These parameters are passed directly to the user-supplied baseline cumulative hazard function 'chfun()'.</p>
chfun	<p>A function computing the baseline cumulative hazard:</p> <ul style="list-style-type: none"> * 2-parameter case: 'chfun(time, ae0, be0)' * 3-parameter case: 'chfun(time, ae0, be0, ce0)' <p>The function must return a vector of values of equal length to 'time'.</p>

Details

Likelihood formulation

For each individual i , let $t_{i1} < t_{i2} < \dots < t_{iK_i}$ denote the *observation / measurement times*.

The cumulative hazard contribution over interval (t_{ij-1}, t_{ij}) is:

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

where x_{ij} is the vector of covariates measured at time t_{ij} .

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:

- * ΔH_{ij} comes from cumulative hazard increments
- * $(T_i = t_{iK})$ is the final event or censoring time
- * δ_i 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 ΔH_{ij} for each ID 4. Constructs the likelihood 5. Optimises over β 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

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

Value

the value of the PGW hazard function

hweibull	<i>Weibull (W) hazard function.</i>
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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	<i>Generalised Gamma (GG) cumulative distribution function.</i> <i>https://rpubs.com/FJRubio/GG</i>
--------	---

Description

Generalised Gamma (GG) cumulative distribution function. <https://rpubs.com/FJRubio/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	<i>Power Exponentiated Weibull (EW) quantile function.</i> <i>https://rpubs.com/FJRubio/EWD</i>
-----	--

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

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

qllogis	<i>Log-logistic (LL) quantile function.</i>
---------	---

Description

Log-logistic (LL) quantile function.

Usage

```
qllogis(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	<i>Power Generalised Weibull (PGW) quantile function.</i> <i>http://rpubs.com/FJRubio/PGW</i>
------	---

Description

Power Generalised Weibull (PGW) quantile function. <http://rpubs.com/FJRubio/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	<i>Generalised Gamma (GG) random number generation.</i> <i>https://rpubs.com/FJRubio/GG</i>
---------	---

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

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

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

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

Description

Computes the survival function $\exp(-H(t \mid x(t)))$ 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

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

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 survival function evaluated at the last time point in ‘df’.

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