

# Package ‘HazReg’

December 16, 2025

**Type** Package

**Title** Parametric hazard-based regression models

**Version** 0.1.0

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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

**RoxygenNote** 7.3.3

**Imports** numDeriv, matrixStats

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chew	<i>Power Exponentiated Weibull (EW) cumulative hazard function.</i> <a href="https://rpubs.com/FJRUBIO/EWD">https://rpubs.com/FJRUBIO/EWD</a>
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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>
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**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

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

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CH_TVC	<i>Compute the Cumulative Hazard for a Proportional Hazards or Accelerated Failure Model (2- and 3-parameter baseline)</i>
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### 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"> <li>• ‘time’: numeric vector of time points.</li> <li>• Covariate columns named with prefix “des” (e.g., ‘des1’, ‘des2’, ...), representing <math>x(t)</math>.</li> </ul>
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 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

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dggamma

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

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### 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

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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

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GHMLE	<i>GHMLE function: Hazard Regression Models with a parametric baseline hazard</i>
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**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")<br>*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

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

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**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

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

<b>df</b>	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.
<b>method</b>	Optimisation method for the likelihood. Either ““nlminb”“ or a valid ‘optim()‘ method.
<b>maxit</b>	Maximum number of optimisation iterations.
<b>beta</b>	Numeric vector of regression coefficients associated with the time-varying covariate design matrix (‘des*’ columns).
<b>ae0</b>	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()‘.
<b>chfun</b>	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><a href="http://rpubs.com/FJRubio/PGW">http://rpubs.com/FJRubio/PGW</a></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>

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**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>
	<a href="https://rpubs.com/FJRubio/GG">https://rpubs.com/FJRubio/GG</a>

---

**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: <a href="https://github.com/FJRubio67/HazReg">https://github.com/FJRubio67/HazReg</a></i>
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**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>
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---

**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"> <li>• ‘time’: numeric vector of time points.</li> <li>• Covariate columns named with prefix “des” (e.g., ‘des1’, ‘des2’, ...), representing <math>x(t)</math>.</li> </ul>
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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