

# Package: blendR (via r-universe)

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**Title** Blended Survival Curves

**Version** 1.0.0

**Description** Create blended survival curves, see 'Che et al.' (2022)

<doi:10.1177/0272989X221134545>.

**License** GPL (>= 3)

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

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**Imports** INLA, dplyr, flexsurv, ggplot2, manipulate, sn, survHE, tibble

**Depends** R (>= 4.4.0)

**Suggests** knitr, rlang, rmarkdown, remotes, survival, testthat (>= 3.0.0)

**VignetteBuilder** knitr

**Additional\_repositories** <https://inla.r-inla-download.org/R/stable/>

**URL** <https://github.com/StatisticsHealthEconomics/blendR/>,

<https://StatisticsHealthEconomics.github.io/blendR/>,

<https://github.com/StatisticsHealthEconomics/blendR/>

**BugReports** <https://github.com/StatisticsHealthEconomics/blendR/issues/>

**Config/testthat.edition** 3

**Repository** <https://n8thangreen.r-universe.dev>

**RemoteUrl** <https://github.com/StatisticsHealthEconomics/blendR>

**RemoteRef** HEAD

**RemoteSha** c98a9b78cc95303bd38da1c8472c3c830abaa2e

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|           |                                |
|-----------|--------------------------------|
| blendsurv | <i>Blended survival object</i> |
|-----------|--------------------------------|

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### Description

This is the main function in the **blendR** package. Two survival curves are supplied and blended according to the blending distribution characterised by the blending interval and the beta distribution parameters.

### Usage

```
blendsurv(
  obs_Surv,
  ext_Surv,
  blend_interv,
  beta_params = list(alpha = 3, beta = 3),
  times = NULL,
  nsim = 100
)
```

### Arguments

|                    |  |
|--------------------|--|
| obs_Surv, ext_Surv | Observed and external data survival curves. These can come from <b>survHE</b> , <b>INLA</b> or <b>flexsurv</b> fits. |
| blend_interv       | Maximum and minimum values for the blending interval.  |
| beta_params        | coefficients of a beta distribution  |
| times              | A vector of times for which the survival curves are to be computed; optional   |
| nsim               | The number of simulations from the distribution of the survival curves; default 100                                  |

### Value

List of S for observed, external and blended curves.

## Examples

```

library(survHE)

## trial data
data("TA174_FCR", package = "blendR")

## externally estimated data
data_sim <- ext_surv_sim(t_info = 144,
                         S_info = 0.05,
                         T_max = 180)

obs_Surv <- fit.models(formula = Surv(death_t, death) ~ 1,
                       data = dat_FCR,
                       distr = "exponential",
                       method = "hmc")

ext_Surv <- fit.models(formula = Surv(time, event) ~ 1,
                       data = data_sim,
                       distr = "exponential",
                       method = "hmc")

blend_interv <- list(min = 48, max = 150)
beta_params <- list(alpha = 3, beta = 3)

ble_Surv <- blendsurv(obs_Surv, ext_Surv, blend_interv, beta_params)

plot(ble_Surv)

```

dat\_FCR

*Survival data*

## Description

Survival data

ext\_surv\_sim

*Create an external survival data based on expert opinion*

## Description

Generally, the sampling is done in two steps

$$p(T) = p(T|interval_i)p(interval_i)$$

**Usage**

```
ext_surv_sim(t_info, S_info, T_max, n = 100)
```

**Arguments**

|                     |  |
|---------------------|--|
| <code>t_info</code> | A vector of times for which expert opinion is elicited   |
| <code>S_info</code> | A vector of mean survival probabilities estimated by experts corresponding to time points in <code>t_info</code> |
| <code>T_max</code>  | The maximum survival time to be used   |
| <code>n</code>      | The number of patients to construct the artificial external data set; default 100                                |

**Details**

In particular  $T \sim U(x_i, x_{i+1})$   $i \sim \text{multinomial}(\hat{\pi})$

**Value**

Dataframe of times and censoring status

**Examples**

```
dat <- ext_surv_sim(t_info = c(10, 20, 50),
                     S_info = c(0.9, 0.8, 0.2),
                     T_max = 100, n = 100)
if (require(survival)) {
  km_fit <- survfit(Surv(time, event) ~ 1, data = dat)
  plot(km_fit)
}
```

|                          |  |
|--------------------------|--|
| <code>fit_inla_pw</code> | <i>Generate survival estimates with a piecewise exponential Cox model (using INLA)</i> |
|--------------------------|--|

**Description**

Generate survival estimates with a piecewise exponential Cox model (using INLA)

**Usage**

```
fit_inla_pw(
  inla.formula = inla.surv(death_t, death) ~ -1,
  data,
  cutpoints,
  nsim = 100,
  ...
)
```

### Arguments

|              |   |
|--------------|---|
| inla.formula | The formula for PEM which must be an <code>inla.surv</code> object                                |
| data         | A dataframe for survival data with time ( <code>death_t</code> ) and event ( <code>death</code> ) |
| cutpoints    | A sequence of cut points for intervals in the baseline hazard                                     |
| nsim         | The number of simulations from posteriors; default 100  |
| ...          | Additional arguments  |

### Value

INLA object

### Examples

```
## Not run:
data("TA174_FCR", package = "blendR")
head(dat_FCR)

obs_Surv <- fit_inla_pw(data = dat_FCR, cutpoints = seq(0, 180, by = 5))

## End(Not run)
```

make\_surv\_methods      *Create survival probabilities*

### Description

These function are version of the `survHE::make.surv()` function from **survHE**. These are needed prior to blending.

### Usage

```
make_surv(Surv, ...)

## S3 method for class 'survHE'
make_surv(Surv, t, nsim = 100, ...)

## S3 method for class 'flexsurvreg'
make_surv(Surv, t = NULL, nsim = 100, ...)

## S3 method for class 'inla'
make_surv(Surv, t = NULL, nsim = 100, ...)

## Default S3 method:
make_surv(Surv, t = NULL, nsim = 100, ...)
```

**Arguments**

|                   |                                |
|-------------------|--------------------------------|
| <code>Surv</code> | survival analysis object       |
| <code>...</code>  | Additional arguments           |
| <code>t</code>    | Time points; vector            |
| <code>nsim</code> | Number of simulations; integer |

**Value**

matrix of survival probabilities

**Examples**

```
library(survHE)

## trial data
data("TA174_FCR", package = "blendR")

## externally estimated data
data_sim <- ext_surv_sim(t_info = 144,
                           S_info = 0.05,
                           T_max = 180)

ext_Surv <- fit.models(formula = Surv(time, event) ~ 1,
                        data = data_sim,
                        distr = "exponential",
                        method = "hmc")

S_ext <- make_surv(ext_Surv, t = 1:100, nsim = 100)
```

`manip_plot`

*Blended survival plot with manipulate*

**Description**

RStudio bug need to run base R first `manipulate(plot(1:x), x = slider(5, 10))`

**Usage**

```
manip_plot(obs_Surv, ext_Surv, blend_interv)
```

**Arguments**

|                           |                   |
|---------------------------|-------------------|
| <code>obs_Surv</code>     | Observed survival |
| <code>ext_Surv</code>     | External survival |
| <code>blend_interv</code> | Blending interval |

**Value**

Blended survival plot

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|              |   |
|--------------|---|
| plot.blended | <i>Blended survival curve based on short-term data and external information</i> |
|--------------|---|

---

**Description**

Blended survival curve based on short-term data and external information

**Usage**

```
## S3 method for class 'blended'  
plot(x, alpha = c(0.1, 0.05), ...)
```

**Arguments**

|       |  |
|-------|--|
| x     | A blended survival curve object obtain from <a href="#">blendsurv()</a>          |
| alpha | A vector specifying the opacity of ribbon for the blended curve and other curves |
| ...   | Additional arguments   |

**Value**

**ggplot2** object

**See Also**

[blendsurv\(\)](#)

**Examples**

```
library(survHE)  
  
## trial data  
data("TA174_FCR", package = "blendR")  
  
## externally estimated data  
data_sim <- ext_surv_sim(t_info = 144,  
                           S_info = 0.05,  
                           T_max = 180)  
  
obs_Surv <- fit.models(formula = Surv(death_t, death) ~ 1,  
                        data = dat_FCR,  
                        distr = "exponential",  
                        method = "hmc")  
  
ext_Surv <- fit.models(formula = Surv(time, event) ~ 1,
```

```
data = data_sim,  
distr = "exponential",  
method = "hmc")  
  
blend_interv <- list(min = 48, max = 150)  
beta_params <- list(alpha = 3, beta = 3)  
  
ble_Surv <- blendsurv(obs_Surv, ext_Surv, blend_interv, beta_params)  
  
plot(ble_Surv)
```

---

**weightplot**

*Plots the weights for the blending procedure*

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**Description**

Plots the weights for the blending procedure

**Usage**

```
weightplot(x, ...)
```

**Arguments**

|     |   |
|-----|---|
| x   | A blended survival curve object obtained from <a href="#">blendsurv()</a> |
| ... | Additional arguments  |

**Value**

[ggplot2](#) object

**See Also**

[blendsurv\(\)](#)

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