

# Package ‘cpsurvsim’

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**Type** Package

**Title** Simulating Survival Data from Change-Point Hazard Distributions

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**Description** Simulates time-to-event data

with type I right censoring using two methods: the inverse CDF method and our proposed memoryless method. The latter method takes advantage of the memoryless property of survival and simulates a separate distribution between change-points. We include two parametric distributions: exponential and Weibull. Inverse CDF method draws on the work of Rainer Walke (2010),  
[<https://www.demogr.mpg.de/papers/technicalreports/tr-2010-003.pdf>](https://www.demogr.mpg.de/papers/technicalreports/tr-2010-003.pdf).

**Depends** R (>= 3.6.0)

**License** GPL (>= 3)

**Encoding** UTF-8

**Imports** plyr (>= 1.8.5), stats, Hmisc (>= 4.3.0), knitr (>= 1.27)

**Suggests** rmarkdown, testthat

**RoxygenNote** 7.2.3

**VignetteBuilder** knitr

**URL** <https://github.com/camillejo/cpsurvsim>

**BugReports** <https://github.com/camillejo/cpsurvsim/issues>

**NeedsCompilation** no

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## R topics documented:

cpsurvsim . . . . .	2
exp_cdfsims . . . . .	2
exp_icdf . . . . .	4
exp_memsim . . . . .	5
weib_cdfsims . . . . .	6
weib_icdf . . . . .	7
weib_memsim . . . . .	8

<b>Index</b>	<b>10</b>
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<b>cpsurvsim</b>	<i>cpsurvsim: Simulating Survival Data from Change-Point Hazard Distributions</i>
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### Description

The `cpsurvsim` package simulates time-to-event data with type I right censoring using two methods: the inverse CDF method and a memoryless method (for more information on simulation methods, see the vignette). We include two parametric distributions: exponential and Weibull.

### `cpsurvsim` functions

For the exponential distribution, the `exp_icdf` function simulates values from the inverse exponential distribution. `exp_cdfsims` and `exp_memsim` return time-to-event datasets simulated using the inverse CDF and memoryless methods respectively.

For the Weibull distribution, the `weib_icdf` function simulates values from the inverse Weibull distribution. `weib_cdfsims` and `weib_memsim` return time-to-event datasets simulated using the inverse CDF and memoryless methods respectively.

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<b>exp_cdfsims</b>	<i>Inverse CDF simulation for the exponential change-point hazard distribution</i>
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### Description

`exp_cdfsims` simulates time-to-event data from the exponential change-point hazard distribution by implementing the inverse CDF method.

### Usage

```
exp_cdfsims(n, endtime, theta, tau = NA)
```

## Arguments

n	Sample size
endtime	Maximum study time, point at which all participants are censored
theta	Scale parameter $\theta$
tau	Change-point(s) $\tau$

## Details

This function simulates data for the exponential change-point hazard distribution with  $K$  change-points by simulating values of the exponential distribution and substituting them into the inverse hazard function. This method applies Type I right censoring at the endtime specified by the user. This function allows for up to four change-points.

## Value

Dataset with n participants including a survival time and censoring indicator (0 = censored, 1 = event).

## Examples

```

nochangepoint <- exp_cdfsim(n = 10, endtime = 20, theta = 0.05)
onechangepoint <- exp_cdfsim(n = 10, endtime = 20,
                             theta = c(0.05, 0.01), tau = 10)
twochangepoints <- exp_cdfsim(n = 10, endtime = 20,
                               theta = c(0.05, 0.01, 0.05), tau = c(8, 12))

# Pay attention to how you parameterize your model!
# This simulates a decreasing hazard
set.seed(7830)
decreasingHazard <- exp_cdfsim(n = 10, endtime = 20,
                                theta = c(0.5, 0.2, 0.01), tau = c(8, 12))
# This tries to fit an increasing hazard, resulting in biased estimates
cp2.nll <- function(par, tau = tau, dta = dta){
  theta1 <- par[1]
  theta2 <- par[2]
  theta3 <- par[3]
  ll <- log(theta1) * sum(dta$time < tau[1]) +
    log(theta2) * sum((tau[1] <= dta$time) * (dta$time < tau[2])) +
    log(theta3) * sum((dta$time >= tau[2]) * dta$censor) -
    theta1 * sum(dta$time * (dta$time < tau[1])) +
    tau[1] * (dta$time >= tau[1]) -
    theta2 * sum((dta$time - tau[1]) * (dta$time >= tau[1]) *
      (dta$time < tau[2]) + (tau[2] - tau[1]) * (dta$time >= tau[2])) -
    theta3 * sum((dta$time - tau[2]) * (dta$time >= tau[2]))
  return(-ll)
}
optim(par = c(0.001, 0.1, 0.5), fn = cp2.nll,
      tau = c(8, 12), dta = decreasingHazard)

```

**exp\_icdf***Inverse CDF for the exponential distribution***Description**

`exp_icdf` simulates values from the inverse CDF of the exponential distribution.

**Usage**

```
exp_icdf(n, theta)
```

**Arguments**

<code>n</code>	Number of output exponential values
<code>theta</code>	Scale parameter $\theta$

**Details**

This function uses the exponential distribution of the form

$$f(t) = \theta \exp(-\theta t)$$

to get the inverse CDF

$$F^{-1}(u) = (-\log(1-u))/\theta$$

where  $u$  is a uniform random variable. It can be implemented directly and is also called by the function [exp\\_memsim](#).

**Value**

Output is a value or a vector of values from the exponential distribution.

**Examples**

```
simdta <- exp_icdf(n = 10, theta = 0.05)
```

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exp\_memsim*Memoryless simulation for the exponential change-point hazard distribution*

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## Description

`exp_memsim` simulates time-to-event data from the exponential change-point hazard distribution by implementing the memoryless method.

## Usage

```
exp_memsim(n, endtime, theta, tau = NA)
```

## Arguments

<code>n</code>	Sample size
<code>endtime</code>	Maximum study time, point at which all participants are censored
<code>theta</code>	Scale parameter $\theta$
<code>tau</code>	Change-point(s) $\tau$

## Details

This function simulates time-to-event data between  $K$  change-points from independent exponential distributions using the inverse CDF implemented in `exp_icdf`. This method applies Type I right censoring at the `endtime` specified by the user.

## Value

Dataset with `n` participants including a survival time and censoring indicator (0 = censored, 1 = event).

## Examples

```
nochangepoint <- exp_memsim( n = 10, endtime = 20, theta = 0.05)
onechangepoint <- exp_memsim(n = 10, endtime = 20,
                             theta = c(0.05, 0.01), tau = 10)
twochangepoints <- exp_memsim(n = 10, endtime = 20,
                               theta = c(0.05, 0.01, 0.05), tau = c(8, 12))

# Pay attention to how you parameterize your model!
# This simulates a decreasing hazard
set.seed(1245)
decreasingHazard <- exp_memsim(n = 10, endtime = 20,
                                theta = c(0.05, 0.02, 0.01), tau = c(8, 12))
# This tries to fit an increasing hazard, resulting in biased estimates
cp2.nll <- function(par, tau = tau, dta = dta){
  theta1 <- par[1]
  theta2 <- par[2]
```

```

theta3 <- par[3]
l1 <- log(theta1) * sum(dta$time < tau[1]) +
  log(theta2) * sum((tau[1] <= dta$time) * (dta$time < tau[2])) +
  log(theta3) * sum((dta$time >= tau[2]) * dta$censor) -
  theta1 * sum(dta$time * (dta$time < tau[1])) +
  tau[1] * (dta$time >= tau[1])) -
  theta2 * sum((dta$time - tau[1]) * (dta$time >= tau[1]) *
    (dta$time < tau[2]) + (tau[2] - tau[1]) * (dta$time >= tau[2])) -
  theta3 * sum((dta$time - tau[2]) * (dta$time >= tau[2]))
return(-l1)
}
optim(par = c(0.001, 0.1, 0.5), fn = cp2.nll,
      tau = c(8, 12), dta = decreasingHazard)

```

**weib\_cdfsim**

*Inverse CDF simulation for the Weibull change-point hazard distribution*

**Description**

`weib_cdfsim` simulates time-to-event data from the Weibull change-point hazard distribution by implementing the inverse CDF method.

**Usage**

```
weib_cdfsim(n, endtime, gamma, theta, tau = NA)
```

**Arguments**

<code>n</code>	Sample size
<code>endtime</code>	Maximum study time, point at which all participants are censored
<code>gamma</code>	Shape parameter $\gamma$
<code>theta</code>	Scale parameter $\theta$
<code>tau</code>	Change-point(s) $\tau$

**Details**

This function simulates data from the Weibull change-point hazard distribution with  $K$  change-points by simulating values of the exponential distribution and substituting them into the inverse hazard function. This method applies Type I right censoring at the endtime specified by the user. This function allows for up to four change-points and  $\gamma$  is held constant.

**Value**

Dataset with `n` participants including a survival time and censoring indicator (0 = censored, 1 = event).

## Examples

```

nochangepoint <- weib_cdfs(n = 10, endtime = 20, gamma = 2,
                             theta = 0.5)
onechangepoint <- weib_cdfs(n = 10, endtime = 20, gamma = 2,
                             theta = c(0.05, 0.01), tau = 10)
twochangepts <- weib_cdfs(n = 10, endtime = 20, gamma = 2,
                           theta = c(0.05, 0.01, 0.05), tau = c(8, 12))

#' # Pay attention to how you parameterize your model!
# This simulates an increasing hazard
set.seed(9945)
increasingHazard <- weib_cdfs(n = 100, endtime = 20, gamma = 2,
                               theta = c(0.001, 0.005, 0.02), tau = c(8, 12))
# This tries to fit a decreasing hazard, resulting in biased estimates
cp2.nll <- function(par, tau = tau, gamma = gamma, dta = dta){
  theta1 <- par[1]
  theta2 <- par[2]
  theta3 <- par[3]
  ll <- (gamma - 1) * sum(dta$censor * log(dta$time)) +
    log(theta1) * sum((dta$time < tau[1])) +
    log(theta2) * sum((tau[1] <= dta$time) * (dta$time < tau[2])) +
    log(theta3) * sum((dta$time >= tau[2]) * dta$censor) -
    (theta1/gamma) * sum((dta$time^gamma) * (dta$time < tau[1]) +
      (tau[1]^gamma) * (dta$time >= tau[1])) -
    (theta2/gamma) * sum((dta$time^gamma - tau[1]^gamma) *
      (dta$time >= tau[1]) * (dta$time < tau[2])) +
    (tau[2]^gamma - tau[1]^gamma) * (dta$time >= tau[2])) -
    (theta3/gamma) * sum((dta$time^gamma - tau[2]^gamma) *
      (dta$time >= tau[2])))
  return(-ll)
}
optim(par = c(0.2, 0.02, 0.01), fn = cp2.nll,
      tau = c(8, 12), gamma = 2,
      dta = increasingHazard)

```

## Description

weib\_icdf returns a value from the Weibull distribution by using the inverse CDF.

## Usage

```
weib_icdf(n, gamma, theta)
```

### Arguments

<code>n</code>	Number of output Weibull values
<code>gamma</code>	Shape parameter $\gamma$
<code>theta</code>	Scale parameter $\theta$

### Details

This function uses the Weibull density of the form

$$f(t) = \theta t^{(\gamma - 1)} \exp(-\theta/\gamma t^{(\gamma)})$$

to get the inverse CDF

$$F^{(-1)}(u) = (-\gamma/\theta \log(1-u))^{(1/\gamma)}$$

where  $u$  is a uniform random variable. It can be implemented directly and is also called by the function `weib_memsim`.

### Value

Output is a value or vector of values from the Weibull distribution.

### Examples

```
simdta <- weib_icdf(n = 10, theta = 0.05, gamma = 2)
```

---

`weib_memsim`

*Memoryless simulation for the Weibull change-point hazard distribution*

---

### Description

`weib_memsim` simulates time-to-event data from the Weibull change-point hazard distribution by implementing the memoryless method.

### Usage

```
weib_memsim(n, endtime, gamma, theta, tau = NA)
```

### Arguments

<code>n</code>	Sample size
<code>endtime</code>	Maximum study time, point at which all participants are censored
<code>gamma</code>	Shape parameter $\gamma$
<code>theta</code>	Scale parameter $\theta$
<code>tau</code>	Change-point(s) $\tau$

## Details

This function simulates time-to-event data between  $K$  change-points  $\tau$  from independent Weibull distributions using the inverse Weibull CDF implemented in [weib\\_icdf](#). This method applies Type I right censoring at the endtime specified by the user.  $\gamma$  is held constant.

## Value

Dataset with  $n$  participants including a survival time and censoring indicator (0 = censored, 1 = event).

## Examples

```

nochangept <- weib_memsim(n = 10, endtime = 20, gamma = 2,
    theta = 0.05)
onechangepoint <- weib_memsim(n = 10, endtime = 20, gamma = 2,
    theta = c(0.05, 0.01), tau = 10)
twochangepts <- weib_memsim(n = 10, endtime = 20, gamma = 2,
    theta = c(0.05, 0.01, 0.05), tau = c(8, 12))

# Pay attention to how you parameterize your model!
# This simulates an increasing hazard
set.seed(5738)
increasingHazard <- weib_memsim(n = 100, endtime = 20, gamma = 2,
    theta = c(0.001, 0.005, 0.02), tau = c(8, 12))
# This tries to fit a decreasing hazard, resulting in biased estimates
cp2.nll <- function(par, tau = tau, gamma = gamma, dta = dta){
    theta1 <- par[1]
    theta2 <- par[2]
    theta3 <- par[3]
    ll <- (gamma - 1) * sum(dta$censor * log(dta$time)) +
        log(theta1) * sum((dta$time < tau[1])) +
        log(theta2) * sum((tau[1] <= dta$time) * (dta$time < tau[2])) +
        log(theta3) * sum((dta$time >= tau[2]) * dta$censor) -
        (theta1/gamma) * sum((dta$time^gamma) * (dta$time < tau[1])) +
        (tau[1]^gamma) * (dta$time >= tau[1])) -
        (theta2/gamma) * sum((dta$time^gamma - tau[1]^gamma) *
            (dta$time >= tau[1]) * (dta$time < tau[2])) +
        (tau[2]^gamma - tau[1]^gamma) * (dta$time >= tau[2])) -
        (theta3/gamma) * sum((dta$time^gamma - tau[2]^gamma) *
            (dta$time >= tau[2])))
    return(-ll)
}
optim(par = c(0.2, 0.02, 0.01), fn = cp2.nll,
    tau = c(8, 12), gamma = 2,
    dta = increasingHazard)

```

# Index

`cpsurvsim`, 2

`cpsurvsim`-package (`cpsurvsim`), 2

`exp_cdfsim`, 2, 2

`exp_icdf`, 2, 4

`exp_memsim`, 2, 4, 5

`weib_cdfsim`, 2, 6

`weib_icdf`, 2, 7, 9

`weib_memsim`, 2, 8, 8