

Package ‘CoxRMSTSimpson’

October 22, 2026

Title Restricted Mean Survival Time from Cox Models

Version 0.1.0

Description Calculates restricted mean survival time from predicted Cox proportional hazards survival curves using composite Simpson's rule and provides plots with shaded areas under the survival curves. Includes simulated survival data for reproducible examples.

License MIT + file LICENSE

Encoding UTF-8

Depends R (>= 4.1.0)

Imports survival, graphics, grDevices, stats

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

Config/roxygen2/version 8.1.0

NeedsCompilation no

Author Ian Forde [aut, cre]

Maintainer Ian Forde <fordeia@appstate.edu>

Repository CRAN

Date/Publication 2026-10-22 16:20:14 UTC

Contents

cox_rmst	2
plot_cox_rmst	3
simpson_rmst	4
simulate_survival_data	4
Index	6

`cox_rmst`*Calculate RMST from a Cox Model*

Description

Predicts survival curves for supplied covariate profiles and calculates their RMST using Simpson's rule.

Usage

```
cox_rmst(fit, newdata, tau, n_grid = 101)
```

Arguments

<code>fit</code>	A fitted <code>survival::coxph</code> model.
<code>newdata</code>	Data frame of covariate profiles. Each row defines one survival curve.
<code>tau</code>	Positive restriction time.
<code>n_grid</code>	Odd number of equally spaced grid points.

Value

An object of class `"cox_rmst"` containing survival curves, RMST estimates, the fitted model, and the time grid.

Examples

```
library(survival)
data(lung, package = "survival")

fit <- coxph(Surv(time, status == 2) ~ age + sex, data = lung)

profiles <- data.frame(
  age = c(50, 70),
  sex = c(1, 2),
  row.names = c("Younger", "Older")
)

result <- cox_rmst(fit, newdata = profiles, tau = 365)
print(result)
```

plot_cox_rmst*Plot Cox Survival Curves with Shaded RMST Areas*

Description

Plot Cox Survival Curves with Shaded RMST Areas

Usage

```
plot_cox_rmst(x, colors = NULL, alpha = 0.2, legend_position = "topright", ...)
```

Arguments

x	An object returned by <code>cox_rmst()</code> .
colors	Colors for the survival curves.
alpha	Transparency for shaded areas, between 0 and 1.
legend_position	Position passed to <code>graphics::legend()</code> .
...	Additional arguments reserved for future methods.

Value

Invisibly returns x.

Examples

```
library(survival)
data(lung, package = "survival")

fit <- coxph(Surv(time, status == 2) ~ age + sex, data = lung)

profiles <- data.frame(
  age = c(50, 70),
  sex = c(1, 2),
  row.names = c("Younger", "Older")
)

result <- cox_rmst(fit, newdata = profiles, tau = 365)
plot_cox_rmst(result)
```

simpson_rmst

Integrate a Survival Curve Using Simpson's Rule

Description

Calculates the area under a survival curve using composite Simpson's 1/3 rule.

Usage

```
simpson_rmst(t, S)
```

Arguments

t	Equally spaced, increasing time points.
S	Survival probabilities corresponding to t.

Value

A numeric scalar representing the integrated area.

simulate_survival_data

Simulate Survival Data Under a Proportional Hazards Model

Description

Simulate Survival Data Under a Proportional Hazards Model

Usage

```
simulate_survival_data(
  n = 200,
  beta = 0.7,
  lambda = 0.03,
  censor_rate = 0.005,
  seed = NULL
)
```

Arguments

n	Number of subjects.
beta	True regression coefficient for X.
lambda	Baseline exponential hazard rate.
censor_rate	Independent exponential censoring rate.
seed	Optional random seed.

simulate_survival_data

5

Value

A data frame with time, status, and X.

Index

`cox_rmst`, [2](#)

`plot_cox_rmst`, [3](#)

`simpson_rmst`, [4](#)

`simulate_survival_data`, [4](#)