

# Package ‘SurvSparse’

May 7, 2026

**Type** Package

**Title** Survival Analysis with Sparse Longitudinal Covariates

**Version** 0.1

**Description** Survival analysis with sparse longitudinal covariates under right censoring scheme. Different hazards models are involved. Please cite the manuscripts corresponding to this package: Sun, Z. et al. (2022) <[doi:10.1007/s10985-022-09548-6](https://doi.org/10.1007/s10985-022-09548-6)>, Sun, Z. and Cao, H. (2023) <[doi:10.48550/arXiv.2310.15877](https://doi.org/10.48550/arXiv.2310.15877)> and Sun, D. et al. (2023) <[doi:10.48550/arXiv.2308.15546](https://doi.org/10.48550/arXiv.2308.15546)>

**License** GPL-3

**Encoding** UTF-8

**Imports** splines, stats, dplyr, MASS, nloptr, nleqslv, tibble,foreach, gaussquad, tidyr, purrr

**RoxygenNote** 7.2.3

**NeedsCompilation** no

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**Repository** CRAN

**Date/Publication** 2023-10-31 17:30:02 UTC

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add.haz *Additive hazards model with sparse longitudinal covariates*

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### Description

Regression analysis of additive hazards model with sparse longitudinal covariates. Three different weighting schemes are provided to impute the missing values.

### Usage

```
add.haz(data, n, tau, h, method)
```

### Arguments

data	An object of class tibble. The structure of the tibble must be: tibble(id = id, X = failure time, covariates = observation for covariates, obs_times = observation times, delta = censoring indicator).
n	An object of class integer. The sample size.
tau	An object of class numeric. The pre-specified time endpoint.
h	An object of class vector. If use auto bandwidth selection, the structure of the vector must be: h = c(the maximum bandwidth, the minimum bandwidth, the number of bandwidth divided). If use fixed bandwidth, h is the chosen bandwidth.
method	An object of class integer. If use weighted LVCF, method = 1. If use half kernel, method = 2. If use full kernel, method = 3.

### Value

a list with the following elements:

est	The estimation for the corresponding parameters.
se	The estimation for the standard error of the estimated parameters.

### References

Sun, Z. et al. (2022) <doi:10.1007/s10985-022-09548-6>

### Examples

```
library(gaussquad)
library(dplyr)
library(nleqslv)
library(MASS)
n=500
lqrule64 <- legendre.quadrature.rules(64)[[64]]
simdata <- function(alpha,beta ) {
  cen=1
```

```

nstep=20
Sigmat_z <- exp(-abs(outer(1:nstep, 1:nstep, "-")) / nstep)
z <- c(mvnorm( 1, c(1: nstep)/2, Sigmat_z ))
left_time_points <- (0:(nstep - 1)) / nstep
z_fun <- stepfun(left_time_points, c(0,z ))
lam_fun <- function(tt) { alpha(tt)+beta*z_fun(tt)}
u <- runif(1)
fail_time <- nleqslv( 0 , function(ttt)
  legendre.quadrature(lam_fun,
    lower = 0,
    upper = ttt,
    lqrule64) + log(u))$x

X <- min(fail_time, cen)
obs=rpois(1,5)+1
tt= sort(runif(obs, min = 0, max = 1))
obs_times <- tt[which(tt<=cen)]
if (length(obs_times) == 0)
  obs_times <- cen
covariates_obscov <- z_fun(obs_times)
return( tibble(X = X,delta = fail_time < cen,
  covariates = covariates_obscov,obs_times = obs_times, censoring = cen ) ) }
data <- replicate(n, simdata(alpha = function(tt) tt, 1 ),
  simplify = FALSE ) %>% bind_rows(.id = "id")

add.haz(data,n,1,n^(-0.5),3)

```

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trans.haz

*Transformed hazards model with sparse longitudinal covariates*


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## Description

Statistical inference on transformed hazards model with sparse longitudinal covariates. Kernel-weighted log-likelihood and sieve maximum log-likelihood estimation are combined to conduct statistical inference on the hazards function.

## Usage

```
trans.haz(data, n, nknots, norder, tau, s, h)
```

## Arguments

data	An object of class tibble. The structure of the tibble must be: tibble(id = id, X = failure time, covariates = observation for covariates, obs_times = observation times, delta = censoring indicator).
n	An object of class integer. The sample size.
nknots	An object of class integer. The number of knots for B-spline.
norder	An object of class integer. The order of B-spline.

tau            An object of class numeric. The maximum follow-up time.  
 s              An object of class numeric. The parameter for Box-Cox transformation.  
 h              An object of class vector. If use auto bandwidth selection, the structure of the vector must be:  $h = c(\text{the maximum bandwidth, the minimum bandwidth, the number of bandwidth divided})$ . If use fixed bandwidth, h is the chosen bandwidth.

### Value

a list with the following elements:

est            The estimation for the corresponding parameters.  
 se             The estimation for the standard error of the estimated parameters.

### References

Sun, D. et al. (2023) <arXiv:2308.15549>

### Examples

```

library(dplyr)
library(gaussquad)
library(nleqslv)
library(MASS)
n=200
lqrule64 <- legendre.quadrature.rules(64)[[64]]
simdata <- function( beta ) {
  cen=1
  nstep=20
  Sigmat_z <- exp(-abs(outer(1:nstep, 1:nstep, "-")) / nstep)
  z <- 2*(pnorm(c(mvrnorm( 1, rep(0,20), Sigmat_z ))) - 0.5)
  left_time_points <- (0:(nstep - 1)) / nstep
  z_fun <- stepfun(left_time_points, c(0,z ))
  h_fun <- function(x) { beta * z_fun(x) }
  lam_fun <- function(tt) 2 * exp(h_fun(tt))
  u <- runif(1)
  fail_time <- nleqslv(0, function(ttt)
  legendre.quadrature(lam_fun, lower = 0, upper = ttt, lqrule64) + log(u))$x
  X <- min(fail_time, cen)
  obs=rpois(1, 5)+1
  tt= sort(runif(obs, min = 0, max = 1))
  obs_times <- tt[which(tt<=cen)]
  if (length(obs_times) == 0)
    obs_times <- cen
  covariates_obs cov <- z_fun(obs_times)
  return( tibble(X = X, delta = fail_time < cen,
  covariates = covariates_obs cov, obs_times = obs_times, censoring = cen ) )
}
beta=1
data <- replicate( n, simdata( beta ), simplify = FALSE ) %>% bind_rows(.id = "id")
trans.haz(data,n,3,3,1,s=0,n^(-0.35))

```

**Description**

Regression analysis of multiplicative hazards model with sparse longitudinal covariates. The kernel weighting approach is employed to impute the missing value and localize the estimating equation. A wild bootstrap-based simultaneous confidence band for the nonparametric function is also provided.

**Usage**

```
tv.co.Cox(data, n, l, times, bd, scb)
```

**Arguments**

data	An object of class tibble. The structure of the tibble must be: tibble(id = id, X = failure time, covariates = observation for covariates, obs_times = observation times, delta = censoring indicator).
n	An object of class integer. The sample size.
l	An object of class vector. The selection vector. For example, for the p dimensional regression coefficient function, if we want to construct simultaneous confidence band for the first regression coefficient function, we can take $l=c(1,0,\dots,0)$ .
times	An object of class vector. The interest time.
bd	An object of class vector. If use auto bandwidth selection, the structure of the vector must be: $bd=c(\text{the maximum bandwidth, the minimum bandwidth, the number of bandwidth divided})$ . If use fixed bandwidth, bd is the chosen bandwidth.
scb	An object of class vector. If need to construct the simultaneous confidence band, the structure of the vector must be: $c(\text{desirable confidence level, repeat times})$ . Otherwise, $scb=0$ .

**Value**

a list with the following elements:

est	The estimation for the corresponding parameters.
se	The estimation for the standard error of the estimated parameters.
scb	The quantile used to construct simultaneous confidence band.

**References**

Sun, Z. and Cao, H. (2023) <arXiv:2310.15877>

**Examples**

```

library(dplyr)
library(gaussquad)
library(MASS)
library(nleqslv)
n=500
beta<-function(t){
  0.5*(t+0.5)^2
}
lqrule64 <- legendre.quadrature.rules(64)[[64]]
simdata <- function( beta ) {
  cen=1
  nstep=20
  Sigmat_z <- exp(-abs(outer(1:nstep, 1:nstep, "-")) / nstep)
  z <-c(mvnorm( 1, rep(0,20), Sigmat_z ))
  left_time_points <- (0:(nstep - 1)) / nstep
  z_fun <- stepfun(left_time_points, c(0,z ))
  h_fun <- function(x) { beta(x) * z_fun(x) }
  lam_fun <- function(tt) 2 * exp(h_fun(tt))
  u <- runif(1)
  fail_time <- nleqslv(0, function(ttt)
  legendre.quadrature(lam_fun, lower = 0,upper = ttt, lqrule64) + log(u))$x
  X <- min(fail_time, cen)
  obs=rpois(1, 5)+1
  tt= sort(runif(obs, min = 0, max = 1))
  obs_times <- tt[which(tt<=cen)]
  if (length(obs_times) == 0)
    obs_times <- cen
  covariates_obscov <-z_fun(obs_times)
  return( tibble(X = X,delta = fail_time < cen,
    covariates = covariates_obscov,obs_times = obs_times  ) ) }

data <- replicate( n, simdata( beta ), simplify = FALSE ) %>% bind_rows(.id = "id")
tv.co.Cox(data,n,1,0.2,bd=c(n^(-0.4),n^(-0.4)),scb=0)

```

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