Package 'NPCox'

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Type Package
Title Nonparametric and Semiparametric Proportional Hazards Model
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Description An estimation procedure for the analysis of nonparametric proportional haz- ards model (e.g. $h(t) = h0(t)exp(b(t)'Z)$), providing estimation of $b(t)$ and its pointwise standard er- rors, and semiparametric proportional hazards model (e.g. $h(t) = h0(t)exp(b(t)'Z1 + c*Z2))$, pro- viding estimation of $b(t)$, c and their standard errors. More de- tails can be found in Lu Tian et al. (2005) <doi:10.1198 01621450400000845="">.</doi:10.1198>
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npcox

Description

Estimation of proportional hazards (PH) model with time-varying coefficients.

Usage

npcox(cva, delta, obstime, SE = FALSE, bandwidth = FALSE, resamp = 100)

Arguments

сvа	Covariate Z in $h(t) = hO(t)exp(b(t)'Z)$
delta	Right censoring indicator for the model
obstime	The observed time = min(censoring time, observed failure time)
SE	Whether or not the estimation of standard error through resampling method will be done. The default value is FALSE.
bandwidth	Bandwidth for kernel function, which can be specified. The default value is FALSE and can be selected through least prediction error over all subjects.
resamp	Number of resampling for estimation of pointwise standard error. The default value is 100.

Details

This is some description of this function.

'npcox' function is designed for PH model with time-varying coefficients, h(t) = h0(t)exp(b(t)'Z), providing estimation of b(t) and its pointwise standard errors on [bandwidth, max(obstime)-badwidth].

Value

a list that contain the estimation result of temporal coefficients, standard error estimation, selected or predesigned bandwidth, dataset, unconverged time points.

Examples

```
data(pbc)
pbc = pbc[(pbc$time < 3000) & (pbc$time > 800), ]
Z = pbc[,c("age","edema")]
colnames(Z) = c("age","edema")
del = pbc$status
tim = pbc$time
res = npcox(cva = Z,delta = del, obstime = tim, bandwidth = 500)
```

npplot

Description

Plot of time-varying coefficient estimated through function 'npcox' or 'spcox'.

Usage

npplot(temp, xrange = NULL, CIlevel = 0.95)

Arguments

temp	Estimation result from function 'npcox' or 'spcox'.
xrange	Illustration range for x-axis.
CIlevel	The default confidence level stands at 0.95.

Details

This is some description of this function.

This is the plot function for

Value

The plot of nonparametric coefficients function 'npcox' or 'spcox'.

Examples

```
data(pbc)
pbc = pbc[(pbc$time < 3000) & (pbc$time > 800), ]
Z = pbc[,c("age","edema")]
colnames(Z) = c("age","edema")
del = pbc$status
tim = pbc$time
res = npcox(cva = Z,delta = del, obstime = tim, bandwidth = 500)
op = par(mfrow = c(1,2))
npplot(res)
par(op)
```

npsimu

Description

An example of data generation for nonparametric PH model.

Usage

npsimu(n, cenpara = NULL)

Arguments

n	The number of sample size, which can be self-determined.
cenpara	Censoring parameter, which is supposed to be positive, for adjustment of cen- soring rate.

Details

This is some description of this function.

'np_simulata' is designed for PH model with time-varying coefficients, h(t) = h0(t)exp(b(t)'Z), generating the covariates, observed time and censoring indicator.

Value

a list that contain covariates, observed time and censoring indicator.

Examples

data = npsimu(200)

pbc

Primary Biliary Cirrhosis data set.

Description

Primary sclerosing cholangitis is an autoimmune disease leading to destruction of the small bile ducts in the liver. Progression is slow but inexhortable, eventually leading to cirrhosis and liver decompensation. The condition has been recognised since at least 1851 and was named "primary biliary cirrhosis" in 1949. Because cirrhosis is a feature only of advanced disease, a change of its name to "primary biliary cholangitis" was proposed by patient advocacy groups in 2014.

This data is from the Mayo Clinic trial in PBC conducted between 1974 and 1984. A total of 424 PBC patients, referred to Mayo Clinic during that ten-year interval, met eligibility criteria for the randomized placebo controlled trial of the drug D-penicillamine. The first 312 cases in the data set

spcox

participated in the randomized trial and contain largely complete data. The additional 112 cases did not participate in the clinical trial, but consented to have basic measurements recorded and to be followed for survival. Six of those cases were lost to follow-up shortly after diagnosis, so the data here are on an additional 106 cases as well as the 312 randomized participants.

Usage

data(pbc)

Format

age: in years albumin: serum albumin (g/dl) alk.phos: alkaline phosphotase (U/liter) ascites: presence of ascites ast: aspartate aminotransferase, once called SGOT (U/ml) bili: serum bilirunbin (mg/dl) chol: serum cholesterol (mg/dl) copper: urine copper (ug/day) edema: 0 no edema, 0.5 untreated or successfully treated 1 edema despite diuretic therapy hepato: presence of hepatomegaly or enlarged liver id: case number platelet: platelet count protime: standardised blood clotting time sex: m/f spiders: blood vessel malformations in the skin stage: histologic stage of disease (needs biopsy) status: status at endpoint, 0/1/2 for censored, transplant, dead time: number of days between registration and the earlier of death, transplantion, or study analysis in July, 1986 trt: 1/2/NA for D-penicillmain, placebo, not randomised trig: triglycerides (mg/dl)

spcox

Nonparametric and semiparametric Cox regression model.

Description

Estimation of proportional hazards (PH) model with time-varying coefficients and constant coefficients.

Usage

```
spcox(
  cva_cons,
  cva_time,
 delta.
  obstime,
 SE = FALSE,
 bandwidth = FALSE,
  resamp = 100
```

Arguments

)

cva_cons	Covariate Z1 with constant coefficient c in $h(t) = h0(t)exp(c'Z1 + b(t)'Z2)$
cva_time	Covariate Z2 with time-varying coefficient $b(t)$ in $h(t) = h0(t)exp(c'Z1 + b(t)'Z2)$
delta	Right censoring indicator for the model
obstime	The observed time = min(censoring time, observed failure time)

SE	Whether or not the estimation of standard error through resampling method will be done. The default value is FALSE.
bandwidth	Bandwidth for kernel function, which can be specified. The default value is FALSE and can be selected through least prediction error over all subjects.
resamp	Number of resampling for estimation of pointwise standard error. The default value is 100.

Details

This is some description of this function.

'spcox' is designed for PH model with both time-varying and constant coefficients, h(t) = h0(t)exp(b(t)'Z1 + c*Z2), providing estimation of b(t), c and their standard errors.

Value

a list that contain the estimation result of both temporal and constant coefficients, standard error estimation, selected or predesigned bandwidth, dataset, unconverged time points.

Examples

```
data(pbc)
pbc = pbc[(pbc$time < 3000) & (pbc$time > 800), ]
Z1 = as.matrix(pbc[,5])
Z2 = as.matrix(pbc[,c('albumin')])
colnames(Z1) = c('age')
colnames(Z2) = c('albumin')
del = pbc$status
tim = pbc$time
res1 = spcox(cva_cons = Z1, cva_time = Z2, delta = del, obstime = tim, bandwidth = 500)
```

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