Package 'coxsei'

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Type Package

Title Fitting a CoxSEI Model

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Description
Fit a CoxSEI (Cox type Self-Exciting Intensity) model to right-censored counting process data.

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coxsei-package | *Fit a Cox-type self-exciting intensity model (CoxSEI) to right-censored counting process data*

Description

Fit the CoxSEI model using the partial likelihood method.

Details

To use the package, the data needs to be prepared into a data frame containing a column named Y for observed event times in ascending order of each individual process, a column named delta indicating if the event is 'death' (1) or 'censoring' (0), a column named id indicating the process id of each event time, and one or more columns giving the value of any covariate variable at the observed event times of each process. Then call the coxseiest function or the identical but much faster coxseiest2 function to estimate the parametric part of the model and then the coxseiInt function to estimate the cumulative baseline intensity function.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

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References

Feng Chen and Kani Chen. (2014). Modeling Event Clustering Using the m-Memory Cox-Type Self-Exciting Intensity Model. International Journal of Statistics and Probability. 3(3): 126-137. doi:10.5539/ijsp.v3n3p126 URL: http://dx.doi.org/10.5539/ijsp.v3n3p126

Feng Chen and Kani Chen. (2014). Case-cohort analysis of clusters of recurrent events. 20(1): 1-15. doi: 10.1007/s10985-013-9275-3

See Also

survival

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coxsei | *CoxSEI model*

Description

Fit a CoxSEI model to counting process data

Usage

```
coxsei(x,...)
## Default S3 method:
coxsei(x,y,delta,id,par.init,m=2,mit=1000,tr=TRUE,
       method="L-BFGS-B",lower=c(rep(-Inf,ncol(x)),-Inf,0),
       upper=rep(Inf,ncol(x) + 2),...)
## S3 method for class 'coxsei'
print(x,...)
## S3 method for class 'coxsei'
plot(x,...)
## S3 method for class 'coxsei'
summary(object,...)
```

Arguments

x	a covariate matrix, or an object of class `coxsei`
y	a vector of observed times
delta	a vector of event indicators: 1=event, 0=censoring
id	the individual/group id to which the event/censoring time correspond
par.init	initial parameter guess to start the iteration
m	lag parameter as in m-dependence
mit	max number of iteration
tr	whether to trace the optimization or not
method	method used in optimization
lower	the lower bound of the parameter space if the L-BFGS-B method of optimization is used.
upper	the upper bound of the paramter space if the L-BFGS-B methodof optimaization is used.
...	further arguments to plot.stepfun
object	an object of the class coxsei

Value

an object of class `coxsei`, basically a list of the following components

coefficients	a numeric vector of coefficients
vcov	the variance-covariance matrix
zval	the vector of z-value of the Wald test statistic
pval	the vector of p-values
details.par	a list returned by theoptim routine
cintfn	a step function as the estimated cumulative baseline intensity function
cintvar	a step function as the variance of the cumulative baseline intensity function estimator
details.cint	a list containing more details about the `cint`

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

References

Feng Chen and Kani Chen. (2014). Modeling Event Clustering Using the m-Memory Cox-Type Self-Exciting Intensity Model. International Journal of Statistics and Probability. 3(3): 126-137. doi:10.5539/ijsp.v3n3p126 URL: http://dx.doi.org/10.5539/ijsp.v3n3p126

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See Also

coxseifit.ex

Examples

```
data(dat,package="coxsei")
acoxsei <- coxsei(dat[,3:5],dat[,1],dat[,2],dat[,6],
                  c(0.2*1:3,log(0.07),log(10)))
summary(acoxsei)
plot(acoxsei,do.points=FALSE)
```

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coxseiest	*Function to estimate the parametric part of the Cox (proportional intensity) self-exciting point process (CoxSEI) model*

Description

Estimate the parametric part of the CoxSEI model using (conditionally) right-censored counting process data.

Usage

```
coxseiest(dat, par.init, m = 2, mit = 1000, tr = TRUE,
          method = "L-BFGS-B", lower=c(rep(-Inf,ncol(dat)-3),-Inf,0),
          upper=rep(Inf,ncol(dat)-3 + 2),
          gfun = function(x, pa) {
           ifelse(x <= 0, rep(0, length(x)), pa[1] * exp(-pa[2] * x))
          })
coxseiest2(dat, par.init, m = 2, mit = 1000, tr = TRUE,
          method = "L-BFGS-B", lower=c(rep(-Inf,ncol(dat)-3),-Inf,0),
          upper=rep(Inf,ncol(dat)-3 + 2),
          gfun = function(x, pa) {
             ifelse(x <= 0, rep(0, length(x)), pa[1] * exp(-pa[2] * x))
          })
coxseiest3(dat, par.init, m = 2, mit = 1000, tr = TRUE,
          method = "L-BFGS-B", lower=c(rep(-Inf,ncol(dat)-3),-Inf,0),
          upper=rep(Inf,ncol(dat)-3 + 2))
```

Arguments

dat	a data frame with columns Y containing the censored event times of each individual process arranged in ascending order with the last time always being the the censoring time, delta containing the event time indicator with value indicator an event time and 0 a censoring time, id specifying the id (process number) of each event time recorded, and the others giving the value of the associated covariate process at the corresponding event times.
par.init	init guess of the value of the parameters to start the optimization iteration with.
m	order of "autoregression" of the excitation term.
mit	maximum number of iteration in the optimization routine
tr	if set to TRUE, print some summary information while the optimization routine is running.
method	method of optimization
lower	vector of lower boundary values of the parameter space
upper	vector of upper boundary of the parameter space
gfun	the excitation function. Defaults to the exponential decay function

$$g(t; \gamma) = \gamma_1 \gamma_2 e^{-\gamma_2 t}$$

Details

coxseiest uses only R code; coxseiest2 uses external C code, and is expected to be 3~4 times fasters than the former; coxseiest3 assumes the excitation function is the exponential function as defaulted by the former two, and hardwares it in the C side of the code, and therefore is much faster than the former two when the exponential excitation function is desired.

Value

A list as that returned by the call to the optimizer routine. For instance,

par	gives the estimate of the parameters
hessian	gives the inverse of the estimate of the variance-covariance matrix

Note

the excitation function has to contain exactly two parameters; a feature that does not seem desiable and might change later.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

References

Feng Chen and Kani Chen. (2014). Modeling Event Clustering Using the m-Memory Cox-Type Self-Exciting Intensity Model. International Journal of Statistics and Probability. 3(3): 126-137. doi:10.5539/ijsp.v3n3p126 URL: http://dx.doi.org/10.5539/ijsp.v3n3p126

Feng Chen and Kani Chen. (2014). Case-cohort analysis of clusters of recurrent events. 20(1): 1-15. doi: 10.1007/s10985-013-9275-3

See Also

See optim for the components of the returned value

Examples

```
data("dat")
## this takes over 15 minutes
##est0 <- coxseiest(dat,par.init=c(0.2,0.4,0.6,0.6,5))
## this one takes about 4 minutes
##est1 <- coxseiest2(dat,par.init=c(0.2,0.4,0.6,0.6,5))
## this one takes about 10 seconds
est2 <- coxseiest3(dat,par.init=c(0.2,0.4,0.6,0.6,5))
```

---

coxseiexp   *CoxSEI model with exponential function*

Description

fit CoxSEI model using an exponential excitation function

Usage

```
coxseiexp(Y, delta, id, Z, par.init, m = 2, mit = 1000, tr = TRUE,
          method = "L-BFGS-B",lower=c(rep(-Inf,ncol(Z)),-Inf,0),
          upper=rep(Inf,ncol(Z) + 2),...)
```

Arguments

Y	the observed times (including censoring times)
delta	indicator of event: 1=event, 0=censoring
id	the id of the individual/group the event/censoring corresponds to
Z	covariate matrix
par.init	initial parameter value to start the iteration
m	the lag parameter as in M-dependence
mit	maximum number of iteration allowed in maximizing the loag partial likelihood
tr	should the optimization process be 'tr'aced
method	method of optimization; defaults to "L-BFGS-B"
lower	vector of lower boundary values of the parameter space
upper	vector of upper boundary of the parameter space
...	other arguments to be passed to the optimization routine

Value

an object of class "coxsei", basically a list with components

coefficients	a named vector of coefficients
vcov	a symmetric matrix which is supposed to be positive definite when m>0, or with the (np-2)x(np-2) major submatrix positive definite when m=0

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

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coxseifit.ex	*CoxSEI model with exponential function*

Description

Fit a CoxSEI model with exponential function to right censored counting process data

Usage

```
coxseifit.ex(dat, par.init, m = 2, mit = 1000, tr = TRUE,
             method = "L-BFGS-B",lower=c(rep(-Inf,ncol(dat)-3),-Inf,0),
             upper=rep(Inf,ncol(dat)-3 + 2),...)
```

Arguments

dat	The data
par.init	initial value of the regression coefficients and coefficients in the excitation function
m	the lag parameter (the m-dependence parameter)
mit	maximum number of iterations allowed in the optimizer
tr	whether to trace the optimization or not
method	the method of optimization used by the optim routine
lower	vector of lower boundary values of the parameter space
upper	vector of upper boundary of the parameter space
...	other arguments to be passed to the optimization routine

Value

A list of some components with kind of self-evident meanings by their name

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

See Also

[coxseiest](#), [coxseiInt](#)

Examples

```
data("dat")
csfit <- coxseifit.ex(dat,c(1:3*0.2,0.7,10))
coef(csfit)
plot(csfit$cintfn,do.points=FALSE)
```

---

coxseiInt	*Calculate the estimator of the cumulative baseline intensity function in the CoxSEI model.*

Description

It takes the paramter of the parametric part (or its theorized value) and calculate the values of the estimator at the jump times; it also gives the values of the estimator for the variance of the intensity estimator.

Usage

```
coxseiInt(dat, parest, hessian=NULL, vcovmat=solve(hessian), m = 2,
        gfun = function(x, pa) {
          ifelse(x <= 0, 0, pa[1] * pa[2] * exp(-pa[2] * x))
        },
        gfungrd = function(x, pa){
          if(length(x)==0)return(matrix(0,2,0));
          rbind(pa[2]*exp(-pa[2]*x),
                pa[1]*exp(-pa[2]*x)*(1-pa[2]*x)
              )
        })
```

Arguments

dat	a data frame containing the right-censored counting process data
parest	the estimate of parameter of the parametric part of the CoxSEI model
hessian	the hessian matrix returned by the optimization procedure in the estimation of the parametric part based on partial likelihood
vcovmat	the variance-covariance matrix of the estimator of the the parametric components; defaulted to the inverse of the hessian matrix
m	autoregressive order in the excitation part of the intensity
gfun	the excitation function; defaults to the exponential decay function
gfungrd	derivative/gradient function of the excitation function

Value

a list giving the jump times and values at these of the estimator of the cumulative baseline intensity function.

x	the ordered death/event times
y	the value of the estimator of the intensity function at the observed death/event times
varest	the value of the estimator of the variance of the estimator of the intensity function, at the jump times

The step function can be obtained using `stepfun`, and plotted by setting `type="s"` in the `plot` function.

Note

Currently doesn't compute the standard error or variance estimator of the baseline cumulative intensity estimator.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
data("dat")
est <- coxseiest3(dat,c(0.2,0.4,0.6,log(0.06),log(5)))
pe <- est$par; pe[4:5] <- exp(pe[4:5]);
ve <- diag(pe) %*% solve(est$hessian, diag(pe));
cintest <- coxseiInt(dat,pe,vcovmat=ve)
plot(cintest,type="s")
```

---

coxseisim	*A function to simulate a CoxSEI process conditional on specified covariate values*

Description

simulate the sample path of the CoxSEI model with given covariate process values, and excitation function and order of autodependence in the excitation term.

Usage

```
coxseisim(parreg, parg, lmd0 = function(tt) (1 + 0.5 * cos(2 * pi *
tt)),
          g = function(x, parg) {
                ifelse(x <= 0, 0, parg[1] * parg[2] * exp(-parg[2] * x))
            },
          censor = 1, m = 2, trace=TRUE,
          Z = function(x) matrix(0, length(x), length(parreg))
          )
```

Arguments

`parreg`	the regression parameter
`parg`	parameters of the excitation function
`lmd0`	the baseline intensity function
`g`	the excitation function
`censor`	the censoring time
`m`	order of autoregression in the excitation component of the intensity process
`trace`	whether to trace the data generation process; defaults to `TRUE`
`Z`	a function to calculate the covariate values at a specified event time

Value

A data frame with provided covariate values and the censoring time, and the generated event times.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
n.smp <- 100;
z <- matrix(NA,n.smp,3)
for(i in 1:n.smp)
z[i,] <- round(c(runif(1,0.5,1.5),runif(1,1.5,2.5),rbinom(1,1,0.5)),2)
dat <- coxseisim(1:3*0.2,c(0.07,10),censor=rlnorm(1,0,0.1),m=2,
Z=function(x)matrix(z[1,],length(x),3,byrow=TRUE))
dat$id <- 1;
for(i in 2:n.smp){
  dattmp <- coxseisim(1:3*0.2,c(0.07,10),censor=rlnorm(1,0,0.1),m=2,
  Z=function(x)matrix(z[i,],length(x),3,byrow=TRUE))
  dattmp$id <- i;
  dat <- rbind(dat,dattmp)
}
```

CumInt	*Cumulative intensity function*

Description

Calculate the cumulative/integrated hazard/intensity function

Usage

```
CumInt(x, int, ...)
```

Arguments

x	the value at which to calculate the cumulative function value
int	the intensity/hazard rate function. Has to be vectorized.
...	the arguments to be passed in to control the behavior of the underlying `integrate` function.

Details

Relies on the numerical integration routine of R.

Value

The value(s) of the cumulative hazard function at the specified x value(s).

Warning

The validity of the user supplied intensity function is not checked.

Note

Not intended to be called by the user directly.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
curve(CumInt(x,int=function(y)1*( y>=0 & y<2)+3*(y>=2 & y<3)+1*(y>=3)),
      0,5,xlab="t",ylab="H(t) of a piece-wise constant hazard fun h(t)")
```

---

dat   *A simulated data set from a CoxSEI model*

Description

Simulated from a CoxSEI model with an exponential excitation function and an AR order 2 for
the self-excitation effects. Generated using the following code: set.seed(1); n.smp <- 50; z <-
matrix(NA,n.smp,3); for(i in 1:n.smp) z[i,] <- round(c(runif(1,0.5,1.5),runif(1,1.5,2.5),rbinom(1,1,0.
dat <- coxseisim(1:3*0.2,c(0.07,10),censor=rlnorm(1,0,0.1),m=2, Z=function(x)matrix(z[1,],length(x),3
dat$id <- 1; for(i in 2:n.smp){ dattmp <- coxseisim(1:3*0.2,c(0.07,10),censor=rlnorm(1,0,0.1),m=2,
Z=function(x)matrix(z[i,],length(x),3,byrow=T)) dattmp$id <- i; dat <- rbind(dat,dattmp)
}

Usage

```
data(dat)
```

Format

A data frame with 307 observations on the following 6 variables.

Y a numeric vector

delta a numeric vector

Z.1 a numeric vector

Z.2 a numeric vector

Z.3 a numeric vector

id a numeric vector

Examples

```
data(dat)
## maybe str(dat) ; plot(dat) ...
```

Dens   *Density function*

Description

Evaluate the density function corresponding to the specified intensity/hazard function int.

Usage

```
Dens(x, int, ...)
```

Arguments

x	the value at which to evaluate the density function
int	the intensity/hazard function. Has to be vectorized.
...	other arguments to be passed to the underlying integrator

Value

A numerical value or vector giving the value(s) of the density function

Note

Relies on R's integrate function

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
set.seed(1); dat <- RND(1000,int=function(x)3*x^2)
hist(dat,freq=FALSE); curve(Dens(x,int=function(x)3*x^2),add=TRUE)
```

Dist   *Distribution function*

Description

Calculate the value at `x` of the distribution function associated with the intensity/hazard function probived through `int`.

Usage

```
Dist(x, int, ...)
```

Arguments

x	the value to evaluate the distribution function at.
int	vectorized function specifying the intensity/hazard function
...	arguments to be passed to the `integrate` function

Value

A number between 0 and 1 inclusive, that gives the value of the distribution function at the specified x value.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
curve(Dist(x,int=function(x)3*x^2),0,5)
curve(pweibull(x,shape=3),0,5,add=TRUE,col=3,lty=3)
```

Quant   *Quantile function*

Description

calculates the value of the quantile function (inverse of the distribution function) of the survival variable with given intensity/hazard function.

Usage

```
Quant(p, int, tolerance = .Machine$double.eps, ...)
```

Arguments

p	the (probability) values to calculate the quantiles at
int	the intensity/hazard function. Has to be vectorized.
tolerance	tolerated numerical error in inverting the distribution function.
...	arguments to be passed to CumInt (eventually to `integrate`)

Value

a numerical value or vector giving the values of the quantile function at x

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
curve(Quant(x,int=function(x)3*x^2),from=1e-3,to=1 - 1e-3)
curve(qweibull(x,shape=3),col=3,lty=3,add=TRUE)
```

RND   *Random number generator*

Description

RND takes a vectorized positive R function defined on positive reals and returns a vector of n values of the random variable (survival time) with the specifed function as its hazard/intensity rate function.

Usage

```
RND(n, int, tol = .Machine$double.eps^0.5, epsabs = 1e-10, epsrel =
1e-10, limit = 1000)
```

Arguments

n	number of observations.
int	hazard rate function of the survival variable, or the intensity function of the one-event point process counting the number (0 or 1) of deaths by following a sample of the surviving suject.
tol	tolerance of the numerical error in calculating the inverse of the cumulative distribution function of the survival variable. Defaults to the square root of the machine epsilon.
epsabs	maximum absolute error to be tolerated by the integrator.
epsrel	maximum relative error to be tolerated by the integrator.
limit	maximum number of iterations permitted by the integrator.

Value

a vector of n observations of the survival variable with the supplied intensity/hazard function.

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
set.seed(1)
dat <- RND(100,int=function(x)x^2)
ks.test(dat,pweibull,shape=3) # p-value = 0.6058
qqplot(dat,rweibull(100,shape=3))
```

Surv *Survival function*

Description

Evaluate the survival function corresponding to the given intensity/hazard function.

Usage

```
Surv(x, int, ...)
```

Arguments

x	value to calculate the value of the survival function for
int	the intensity/hazard function
...	further arguments to be passed to CumInt

Value

a numerical value or vector giving the value(s) of the survival function at x

Author(s)

Feng Chen <feng.chen@unsw.edu.au>

Examples

```
curve(Surv(x, int=function(x)3*x^2), from=0, to=5)
curve(pweibull(x,shape=3,lower=FALSE), add=TRUE, col=2, lty=3)
```

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