

Package ‘isoSurv’

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Type Package

Title Isotonic Regression on Survival Analysis

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Description Nonparametric estimation on survival analysis under order restrictions. It estimates monotone increasing or decreasing covariate effects in the proportional hazards model. Yunro Chung et al. (2018) <[doi:10.1093/biomet/asx064](https://doi.org/10.1093/biomet/asx064)>.

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License GPL (>= 2)

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 isoSurv-package

Isotonic Regression on Survival Analysis

Description

Nonparametric estimation on survival analysis under order restrictions

Details

Package: isoph
 Type: Package
 Version: 0.2.0
 Date: 2021-3-22
 License: GPL (>= 2)

Author(s)

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References

Yunro Chung, Anastasia Ivanova, Michael G. Hudgens, Jason P. Fine (2018), Partial likelihood estimation of isotonic proportional hazards models, *Biometrika*, 105(1), 133-148. doi:10.1093/biomet/asx064

 disoph

Fit Double Isotonic Proportional Hazards Model

Description

Nonparametric full likelihood estimation of monotone baseline hazard and covariate effect functions in the proportional hazards model.

Usage

```
disoph(formula, bshape, data, maxiter, eps)
```

Arguments

`formula` a formula object: `response ~ iso(covariate1)`. The response must be survival outcome using the `Surv` function in the survival package. The `iso(covariate1, shape="increasing", K="median")` is for isotonic estimation of `covariate1` with `shape="increasing"` (or "decreasing") direction and `K="median"` anchor.

bshape	direction of the baseline hazard function (bshape="increasing" or "decreasing").
data	data.frame includes variables named in the formula argument.
maxiter	maximum number of iteration (default is 10^4).
eps	stopping convergence criteria (default is 10^{-3}).

Details

The disoph function estimates λ_0 and ψ based on the full likelihood under the isotonic proportional hazards model, defined as

$$\lambda(t|z) = \lambda_0(t) \exp(\psi(z)),$$

where λ_0 is an isotonic baseline hazard function and ψ is an isotonic function. One point has to be fixed with $\psi(K) = 0$, where K is an anchor point. A direction of λ_0 is defined as monotone increasing or monotone decreasing in t . A direction of ψ is defined as monotone increasing or monotone decreasing in z . Back-and-forth iterative pool adjacent violators algorithm is used to maximize the full likelihood.

Value

A list of class fisoph:

iso.bh	data.frame with <i>time</i> and estimated λ_0 .
iso.cov	data.frame with <i>z</i> and estimated ψ .
conv	algorithm convergence status.
iter	total number of iterations.
Zk	anchor satisfying estimated $\psi(Zk)=0$.
shape.bh	order restriction on λ_0 .
shape.cov	order restriction on ψ .

Author(s)

Yunro Chung [auth, cre]

References

Yunro Chung, Double Isotonic Proportional Hazards Models with Monotone Baseline Hazard and Covariate Effect Functions. In preparation.

Examples

```
test1=data.frame(
  time= c(2, 5, 1, 7, 9, 5, 3, 6, 8, 9, 7, 4, 5, 2, 8),
  status=c(0, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 0, 1),
  z=    c(2, 1, 1, 3, 5, 6, 7, 9, 3, 0, 2, 7, 3, 9, 4)
)

disoph.fit1=disoph(Surv(time, status)~iso(z,shape="inc"),bshape="inc",data=test1)
plot(disoph.fit1)
```

 iso

Attributions of isotonic covariate effect

Description

InternalIt attributes the covariate with respect to the name, direction and anchor constrain.

Usage

```
iso(z, shape, K)
```

Arguments

z a univariate covariate.
 shape a direction of z (shape="increasing" or "decreasing") (default is "increasing").
 K an anchor point (default is *NA* for the median anchor).

Details

Internal function. The iso function attributes the covariate z for its name, shape direction and anchor.

Value

The value *z* with attribution of its name, shape and *K*.

Author(s)

Yunro Chung [cre]

 isoph

Fit Isotonic Proportional Hazards Model

Description

Nonparametric partial likelihood estimation of a monotone covariate effect in the proportional hazards model.

Usage

```
isoph(formula, data, maxiter, eps)
```

Arguments

formula	a formula object: response ~ iso(covariate1)+covariate2+.... The response must be survival outcome using the Surv function in the survival package. The iso(covariate1,shape="increasing" is for isotonic estimation of covariate1 with shape="increasing" (or "decreasing") direction and K="median" anchor.
data	data.frame includes variables named in the formula argument.
maxiter	maximum number of iteration (default is 10 ⁴).
eps	stopping convergence criteria (default is 10 ⁻³).

Details

The isoph function estimates ψ and β based on the partial likelihood under the isotonic proportional hazards model, defined as

$$\lambda(t|z, w) = \lambda_0(t) \exp(\psi(z) + \beta w),$$

where λ_0 is a baseline hazard function, ψ is an isotonic function, z is a univariate variable, w is a q by 1 dimensional covariate vector and β is a 1 by q dimensional regression parameter. One point has to be fixed with $\psi(K) = 0$, where K is an anchor point. A direction of ψ is defined as monotone increasing or monotone decreasing in z . Pseudo iterative convex minorant algorithm is used to maximize the partial likelihood.

Value

A list of class isoph:

iso.cov	data.frame with z and estimated ψ .
beta	estimated β .
conv	algorithm convergence status.
iter	total number of iterations.
Zk	anchor satisfying estimated $\psi(Z_k)=0$. See the formular argument above.
shape	order restriction on ψ . See the formular argument above.

Author(s)

Yunro Chung [aut, cre]

References

Yunro Chung, Anastasia Ivanova, Michael G. Hudgens, Jason P. Fine, Partial likelihood estimation of isotonic proportional hazards models, *Biometrika*. 2018, 105 (1), 133-148. doi:10.1093/biomet/asx064

Examples

```
# test1
test1=data.frame(
  time= c(2, 5, 1, 7, 9, 5, 3, 6, 8, 9, 7, 4, 5, 2, 8),
  status=c(0, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 0, 1),
  z=    c(2, 1, 1, 3, 5, 6, 7, 9, 3, 0, 2, 7, 3, 9, 4)
```

```
)  
  
isoph.fit1 = isoph(Surv(time, status)~iso(z,shape="inc"),data=test1)  
plot(isoph.fit1)  
  
# test2  
test2=data.frame(  
  time= c(2, 5, 1, 7, 9, 5, 3, 6, 8, 9, 7, 4, 5, 2, 8),  
  status=c(0, 1, 0, 1, 1, 0, 1, 0, 1, 1, 1, 0, 1, 1, 1),  
  z=     c(2, 1, 1, 3, 5, 6, 7, 9, 3, 0, 2, 7, 3, 9, 4),  
  trt=   c(1, 1, 1, 0, 1, 0, 1, 0, 0, 1, 0, 1, 1, 0, 0)  
)  
  
isoph.fit2 = isoph(Surv(time, status)~iso(z,shape="inc")+trt, data=test2)  
plot(isoph.fit2)
```

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