

# Package ‘isoSurv’

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**Type** Package

**Title** Isotonic Regression on Survival Analysis

**Version** 0.2.0

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**Author** Yunro Chung [aut, cre]

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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>.

**Depends** R (>= 3.6.0), Iso, survival

**License** GPL (>= 2)

**Encoding** UTF-8

**LazyData** true

**NeedsCompilation** no

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## R topics documented:

isoSurv-package . . . . .	2
disoph . . . . .	2
iso . . . . .	4
isoph . . . . .	4

**Index**

7

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)**

Yunro Chung [aut,cre] Maintainer: Yunro Chung <yunro.chung@asu.edu>

**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**

<b>formula</b>	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.
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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  $\lambda_0$  and  $\psi$  based on the full likelihood under the isotonic proportional hazards model, defined as

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

where  $\lambda_0$  is an isotonic baseline hazard function and  $\psi$  is an isotonic function. One point has to be fixed with  $\psi(K) = 0$ , where  $K$  is an anchor point. A direction of  $\lambda_0$  is defined as monotone increasing or monotone decreasing in  $t$ . A direction of  $\psi$  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 $\lambda_0$ .
iso.cov	data.frame with <i>z</i> and estimated $\psi$ .
conv	algorithm convergence status.
iter	total number of iterations.
Zk	anchor satisfying estimated $\psi(Zk)=0$ .
shape.bh	order restriction on $\lambda_0$ .
shape.cov	order restriction on $\psi$ .

## 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, 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

<code>z</code>	a univariate covariate.
<code>shape</code>	a direction of <code>z</code> ( <code>shape="increasing"</code> or <code>"decreasing"</code> ) (default is <code>"increasing"</code> ).
<code>K</code>	an anchor point (default is <code>NA</code> 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  $\psi$  and  $\beta$  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  $\lambda_0$  is a baseline hazard function,  $\psi$  is an isotonic function,  $z$  is a univariate variable,  $w$  is a q by 1 dimensional covariate vector and  $\beta$  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  $\psi$  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 $\psi$ .
beta	estimated $\beta$ .
conv	algorithm convergence status.
iter	total number of iterations.
Zk	anchor satisfying estimated $\psi(Zk)=0$ . See the formular argument above.
shape	order restriction on $\psi$ . 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, 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, 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, 0, 1, 0, 1, 1, 0, 0)  
)  
  
isoph.fit2 = isoph(Surv(time, status)~iso(z,shape="inc")+trt, data=test2)  
plot(isoph.fit2)
```

# Index

- \* **Isotonic regression**
  - isoSurv-package, [2](#)
- \* **Nonparametric shape restricted estimation**
  - isoSurv-package, [2](#)
- \* **Survival analysis**
  - isoSurv-package, [2](#)
  - disoph, [2](#)
  - iso, [4](#)
  - isoph, [4](#)
  - isoSurv-package, [2](#)