

Package ‘surrosurvROC’

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Type Package

Title Surrogate Survival ROC

Version 0.1.0

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Description Nonparametric and semiparametric estimations of the time-dependent ROC curve for an incomplete failure time data with surrogate failure time endpoints.

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License GPL (>= 2)

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surrosurvROC-package *Surrogate Survival ROC*

Description

Nonparametric and semiparametric estimations of the time-dependent ROC curve for an incomplete failure time data with surrogate failure time endpoints.

Details

Package: isoph
 Type: Package
 Version: 0.1.0
 Date: 2018-08-17
 License: GPL (>= 2)

Author(s)

Yunro Chung [cre]

Maintainer: Yunro Chung <yunro.chung@asu.edu>

References

Yunro Chung and Yingye Zheng, Improving efficiency of evaluating prognostic accuracy of biomarkers for incomplete failure-time data with surrogate outcome (in progress)

surrosurvROC

Surrogate Survival ROC

Description

Nonparametric and semiparametric estimations of the time-dependent ROC curve for an incomplete failure time data with surrogate failure time endpoints

Usage

```
surrosurvROC(DATA, method, pred.time, wt=NULL, span=NULL, b.rep=200)
```

Arguments

DATA	data frame, consisting of Marker: Predictor or marker value; Survival time; Status: Event indicator (1: event; 0: censoring); STime: Surrogate survival Time; SStatus: Surrogate event indicator (1: event; 0: censoring)
method	"KNN" for nonparametric model using nearest neighborhood kernel; "COX" for semiparametric proportional hazard model
pred.time	Prediction time of the ROC curve
wt	Weight, such as inverse probability weighting
span	Smoothing bandwidth parameter for KNN
b.rep	Number of bootstrap

Details

It provides a more efficient time-dependent ROC curve for an incomplete failure time data, when surrogate failure time endpoints are additionally observed for all subjects.

Author(s)

Yunro Chung [cre]

References

Yunro Chung and Yingye Zheng, Evaluating Prognostic Accuracy of Biomarkers for Incomplete and Right-Censored Data with Surrogate Outcome (in progress)

Examples

```
DATA=data.frame(
  Time= c(1,2,5,3,9,NA,8,9,10,NA,NA,NA,6,4,NA,NA,NA,NA,NA,NA,NA,NA,NA),
  Status= c(1,1,0,0,1,NA,1,1,0, NA,NA,NA,0,0,NA,NA,NA,NA,NA,NA,NA,NA,NA),
  STime= c(3,2,4,2,8,5,8,7,11,1,8,9,3,5,2,5,10,3,5,8,5,2,4,6,7),
  SStatus=c(1,0,1,0,1,1,1,0,0,1,1,1,1,0,1,1,0,0,1,0,1,0,1,0,0),
  Marker= c(1,5,1,2,3,1,2,3,4,5,9,8,5,7,3,4,2,5,3,4,7,5,9,3,8)
)

#COX at year 3
RES1=surrosurvROC(DATA, method="COX", pred.time=3)
print(RES1)

#KNN at year 3
nobs=sum(!is.na(DATA$Time))
span=0.25*nobs^(-0.20)
RES2=surrosurvROC(DATA, method="KNN",pred.time=3,span=span)
print(RES2)
```

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* **biomarkers, inverse probability weighing,**

bootstrap

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