

# Package ‘CPE’

September 26, 2018

**Title** Concordance Probability Estimates in Survival Analysis

**Version** 1.5.1

**Depends** R (>= 2.10.0),survival,rms

**Author** Qianxing Mo, Mithat Gonen and Glenn Heller

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**Description** Functions to calculate concordance probability estimates in survival analysis.

**LazyData** no

**License** GPL (>= 2)

**NeedsCompilation** yes

**Repository** CRAN

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## R topics documented:

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| phcpe | <i>Gonen \&amp; Heller Concordance Probability Estimate for the Cox Proportional Hazards model</i> |
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### Description

A function to calculate Gonen \& Heller concordance probability estimate (CPE) for the Cox proportional hazards model.

### Usage

```
phcpe(coxfit, CPE.SE=FALSE,out.ties=FALSE)
```

**Arguments**

|          |  |
|----------|--|
| coxfit   | A coxph or cph object  |
| CPE.SE   | A logical value indicating whether the standard error of the CPE should be calculated  |
| out.ties | If out.ties is set to FALSE, pairs of observations tied on covariates will be used to calculate the CPE. Otherwise, they will not be used. |

**Value**

|        |  |
|--------|--|
| CPE    | Concordance Probability Estimate                           |
| CPE.SE | the Standard Error of the Concordance Probability Estimate |

**Author(s)**

Qianxing Mo, Mithat Gonen and Glenn Heller; <qianxing.mo@moffitt.org>

**References**

Mithat Gonen and Glenn Heller. (2005). Concordance probability and discriminatory power in proportional hazards regression. *Biometrika*, 92, 4, pp.965-970  
 Glenn Heller and Qianxing Mo. (2016). Estimating the concordance probability in a survival analysis with a discrete number of risk groups. *Lifetime Data Analysis*, 22(2):263-79.

**See Also**

[phcpe2](#)

**Examples**

```
### create a simple data set for testing
set.seed(199)
nn <- 1000
time <- rexp(nn)
status <- sample(0:1, nn, replace=TRUE)
covar <- matrix(rnorm(3*nn), ncol=3)
survd <- data.frame(time, status, covar)
names(survd) <- c("time", "status", "x1", "x2", "x3")

coxph.fit <- coxph(Surv(time, status)~x1+x2+x3, data=survd)

### Calculate CPE only (needs much less time).
phcpe(coxph.fit)
phcpe(coxph.fit, out.ties=TRUE)
#result is identical because the covariates are not tied #

### Calculate CPE and CPE.SE
phcpe(coxph.fit, CPE.SE=TRUE)
phcpe(coxph.fit, CPE.SE=TRUE, out.ties=TRUE)
```

```

**** For unknown reason, 'coxph.fit' may need to be removed before running cph()****
rm(coxph.fit)

cph.fit <- cph(Surv(time, status)~x1+x2+x3, data=survd,method="breslow")

### Calculate CPE only (needs much less time).
phcpe(cph.fit)
phcpe(cph.fit,out.ties=TRUE)

### Calculate CPE and CPE.SE
phcpe(cph.fit, CPE.SE=TRUE)
phcpe(cph.fit, CPE.SE=TRUE,out.ties=TRUE)

```

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|        |  |
|--------|--|
| phcpe2 | <i>Gonen \&amp; Heller Concordance Probability Estimate for the Cox Proportional Hazards model</i> |
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### Description

A function to calculate Gonen \& Heller concordance probability estimate (CPE) for the Cox proportional hazards model.

### Usage

```
phcpe2(coef,coef.var,design, CPE.SE=FALSE,out.ties=FALSE)
```

### Arguments

|          |   |
|----------|---|
| coef     | The coefficients of the Cox model.  |
| coef.var | The covariance matrix of the coefficients of the Cox model.   |
| design   | A design matrix for covariates. The rows correspond to subjects, and the columns correspond to covariates.                                |
| CPE.SE   | A logical value indicating whether the standard error of the CPE should be calculated   |
| out.ties | If out.ties is set to FALSE,pairs of observations tied on covariates will be used to calculate the CPE. Otherwise, they will not be used. |

### Value

|        |  |
|--------|--|
| CPE    | Concordance Probability Estimate                           |
| CPE.SE | the Standard Error of the Concordance Probability Estimate |

### Author(s)

Qianxing Mo, Mithat Gonen and Glenn Heller; <qianxing.mo@moffitt.org>

## References

Mithat Gonen and Glenn Heller. (2005). Concordance probability and discriminatory power in proportional hazards regression. *Biometrika*, 92, 4, pp.965-970  
Glenn Heller and Qianxing Mo. (2016). Estimating the concordance probability in a survival analysis with a discrete number of risk groups. *Lifetime Data Analysis*, 22(2):263-79.

## See Also

[phcpe](#)

## Examples

```
### create a simple data set for testing
set.seed(199)
nn <- 1000
time <- rexp(nn)
status <- sample(0:1, nn, replace=TRUE)
covar <- matrix(rnorm(3*nn), ncol=3)
survd <- data.frame(time, status, covar)
names(survd) <- c("time", "status", "x1", "x2", "x3")

coxph.fit <- coxph(Surv(time, status)~x1+x2+x3, data=survd)

phcpe(coxph.fit, CPE.SE=TRUE)
phcpe2(coef=coxph.fit$coefficients, coef.var=coxph.fit$var, design=model.matrix(coxph.fit))

*** For unknown reason, 'coxph.fit' may need to be removed before running cph()***
rm(coxph.fit)

cph.fit <- cph(Surv(time, status)~x1+x2+x3, data=survd, method="breslow")

### Calculate CPE only (needs much less time).
phcpe2(cph.fit$coefficients, coef.var=cph.fit$var, design=model.matrix(cph.fit), CPE.SE=TRUE)
```

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\*Topic **survival**

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