

# Package ‘compareC’

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**Type** Package

**Title** Compare Two Correlated C Indices with Right-Censored Survival Outcome

**Version** 1.3.3

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**Description** Proposed by Harrell, the C index or concordance C, is considered an overall measure of discrimination in survival analysis between a survival outcome that is possibly right censored and a predictive-score variable, which can represent a measured biomarker or a composite-score output from an algorithm that combines multiple biomarkers. This package aims to statistically compare two C indices with right-censored survival outcome, which commonly arise from a paired design and thus resulting two correlated C indices.

**License** GPL ( $\geq 2.0$ )

**NeedsCompilation** yes

**Repository** CRAN

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 compareC

*Testing the difference between two correlated overall C indices*


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**Description**

This is a function used to test if the difference in two correlated overall  $C$  indices is statistically significant

**Usage**

```
compareC(timeX, statusX, scoreY, scoreZ)
```

**Arguments**

timeX	The vector of actual survival time $X$ , one survival time for each observation
statusX	The matching vector of event indicator for time $X$ , 1 if occurred and 0 otherwise
scoreY	The vector of the first measured biomarker or score $Y$ , one for each of the same observations
scoreZ	The vector of the second measured biomarker or score $Z$ , one for each of the same observations

**Value**

est.c	The estimated two $C$ indices
est.diff_c	The estimated difference of the two $C$ indices, i.e., $C_{XY} - C_{XZ}$
est.vardiff_c	The estimated variance of the difference of two $C$ indices
est.varCxy	The estimated variance of the $C$ index for <i>scoreY</i>
est.varCxz	The estimated variance of the $C$ index for <i>scoreZ</i>
est.cov	The estimated covariance between the two $C$ indices for <i>scoreY</i> and that for <i>scoreZ</i>
zscore	Z score (test statistic) for hypothesis testing
pval	P value for the comparison of two $C$ indices

**Note**

Under non-random censoring, the returned  $p$ -value is statistically invalid as the assumption of random censoring is violated.

**Author(s)**

Le Kang, Weijie Chen

## References

Harrell FE, Califf RM, Pryor DB, Lee KL, and Rosati RA. (1982) Evaluating the yield of medical tests. *The Journal of the American Medical Association*, 247(18), 2543–2546

Pencina MJ and D'Agostino RB. (2004) Overall  $C$  as a measure of discrimination in survival analysis: model specific population value and confidence interval estimation. *Statistics in Medicine*, 23(13), 2109–2123

Kang L, Chen W, Petrick NA, and Gallas BD. (2014) Comparing two correlated  $C$  indices with right-censored survival outcome: a one-shot nonparametric approach. *Statistics in Medicine*, 34(4), 685–703, doi: 10.1002/sim.6370

## See Also

[estC, vardiffC](#)

## Examples

```
demo(testC)
```

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estC

*Estimation of a single overall C index*

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## Description

This is a function used to estimate a single overall  $C$  index

## Usage

```
estC(timeX, statusX, scoreY)
```

## Arguments

timeX	The vector of actual survival time $X$ , one survival time for each observation
statusX	The matching vector of event indicator for time $X$ , 1 if occurred and 0 otherwise
scoreY	The vector of the measured biomarker $Y$ , one for each of the same observations, based on which the $C$ index is to be estimated

## Value

The estimated  $C$  index

## Author(s)

Le Kang, Weijie Chen

## References

Harrell FE, Califf RM, Pryor DB, Lee KL, and Rosati RA. (1982) Evaluating the yield of medical tests. *The Journal of the American Medical Association*, 247(18), 2543–2546

Pencina MJ and D’Agostino RB. (2004) Overall  $C$  as a measure of discrimination in survival analysis: model specific population value and confidence interval estimation. *Statistics in Medicine*, 23(13), 2109–2123

Kang L, Chen W, Petrick NA, and Gallas BD. (2014) Comparing two correlated  $C$  indices with right-censored survival outcome: a one-shot nonparametric approach. *Statistics in Medicine*, 34(4), 685–703, doi: 10.1002/sim.6370

## See Also

[vardiffC,compareC](#)

## Examples

```
demo(testC)
```

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vardiffC

*Variance estimation of the difference between two correlated overall  $C$  indices*

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## Description

This is a function used to estimate the variance of the difference between two correlated overall  $C$  indices

## Usage

```
vardiffC(timeX, statusX, scoreY, scoreZ)
```

## Arguments

timeX	The vector of actual survival time $X$ , one survival time for each observation
statusX	The matching vector of event indicator for time $X$ , 1 if occurred and 0 otherwise
scoreY	The vector of the first measured biomarker $Y$ , one for each of the same observations, based on which the variance of the difference in overall $C$ indices is to be estimated
scoreZ	The vector of the second measured biomarker $Z$ , one for each of the same observations, based on which the variance of the difference in overall $C$ indices is to be estimated

**Value**

<code>est.vardiff_c</code>	The estimated variance of the difference of two <i>C</i> indices
<code>est.varCxy</code>	The estimated variance of the <i>C</i> index for <i>scoreY</i>
<code>est.varCxz</code>	The estimated variance of the <i>C</i> index for <i>scoreZ</i>
<code>est.cov</code>	The estimated covariance between the two <i>C</i> indices for <i>scoreY</i> and that for <i>scoreZ</i>

**Author(s)**

Le Kang, Weijie Chen

**References**

Harrell FE, Califf RM, Pryor DB, Lee KL, and Rosati RA. (1982) Evaluating the yield of medical tests. *The Journal of the American Medical Association*, 247(18), 2543–2546

Pencina MJ and D’Agostino RB. (2004) Overall *C* as a measure of discrimination in survival analysis: model specific population value and confidence interval estimation. *Statistics in Medicine*, 23(13), 2109–2123

Kang L, Chen W, Petrick NA, and Gallas BD. (2014) Comparing two correlated *C* indices with right-censored survival outcome: a one-shot nonparametric approach. *Statistics in Medicine*, 34(4), 685–703, doi: 10.1002/sim.6370

**See Also**

[estC,compareC](#)

**Examples**

```
demo(testC)
```

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