

# Package ‘fastlogranktest’

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**Type** Package

**Title** A Fast Way to Calculate the p-Value of One or Multiple  
Log-Rank-Tests

**Version** 0.2.1

**Author** Andreas Stelzer, Manuela Lautizi, Tim Kacprowski, Research group of Computational Sys-  
tems Medicine, Chair of Experimental Bioinformatics, TU Munich

**Maintainer** Tim Kacprowski <tim.kacprowski@wzw.tum.de>

**Description** A very fast Log-Rank-Test implementation that is several orders of magni-  
tude faster than the implementation in the 'survival' package.  
Log-Rank-Tests can be computed individually or concurrently using threading.

**License** GPL-3

**URL** <https://github.com/compsysmed/fastlogranktest.git>

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.0.2

**LinkingTo** Rcpp, BH

**Imports** Rcpp

**Suggests** testthat (>= 2.1.0), survival (>= 3.1)

**NeedsCompilation** yes

**Repository** CRAN

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logrank_test	<i>Calculate the Log-Rank-Test very fast</i>
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**Description**

Calculate the Log-Rank-Test very fast

**Usage**

```
logrank_test(groupa, groupb, groupacensored, groupbcensored, onlyz = FALSE)
```

**Arguments**

groupa	vector of group a's survival times
groupb	vector of group b's survival times
groupacensored	vector of censored information of group a's survival times
groupbcensored	vector of censored information of group b's survival times
onlyz	(optional) calculate only z-statistic

**Value**

chi2 statistic, z-statistic, p-value

**Examples**

```
T1 <- c(6, 6, 6, 6, 7, 9, 10, 10, 11, 13, 16, 17, 19, 20, 22, 23, 25, 32, 32, 34, 35)
E1 <- c(1, 1, 1, 0, 1, 0, 1, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 0, 0, 0)
T2 <- c(1, 1, 2, 2, 3, 4, 4, 5, 5, 8, 8, 8, 8, 11, 11, 12, 12, 15, 17, 22, 23)
E2 <- c(1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1)
logrank_test(T1, T2, E1, E2)
#1.679294e+01 -4.097919e+00, 4.168809e-05
```

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multi_logrank_test	<i>Calculate multiple Log-Rank-Tests very fast</i>
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**Description**

Calculate multiple Log-Rank-Tests very fast

**Usage**

```
multi_logrank_test(
  groupas,
  groupbs,
  groupacensoreds,
  groupbcensoreds,
  threadnumber = NULL,
  onlyz = FALSE
)
```

**Arguments**

groupas	list of vectors of groupa's survival times
groupbs	list of vectors of groupb's survival times
groupacensoreds	list of vectors of censored information of groupa's survival times
groupbcensoreds	list of vectors of censored information of groupb's survival times
threadnumber	(optional) set the number of threads used for this function
onlyz	(optional) calculate only z-statistic

**Value**

vector of chi2 statistic, z-statistic, p-value (same order as input)

**Examples**

```
T1 <- c(6, 6, 6, 6, 7, 9, 10, 10, 11, 13, 16, 17, 19, 20, 22, 23, 25, 32, 32, 34, 35)
E1 <- c(1, 1, 1, 0, 1, 0, 1, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0)
T2 <- c(1, 1, 2, 2, 3, 4, 4, 5, 5, 8, 8, 8, 8, 11, 11, 12, 12, 15, 17, 22, 23)
E2 <- c(1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1)
t1s<-list(T1, T1, T1)
e1s<-list(E1, E1, E1)
t2s<-list(T2, T2, T2)
e2s<-list(E2, E2, E2)
multi_logrank_test(t1s, t2s, e1s, e2s)
#1.679294e+01 -4.097919e+00 4.168809e-05 1.679294e+01 -4.097919e+00 4.168809e-05
#1.679294e+01 -4.097919e+00 4.168809e-05
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