

# Package: controlTest (via r-universe)

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**Type** Package

**Title** Quantile Comparison for Two-Sample Right-Censored Survival Data

**Version** 1.1.0

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**Description** Nonparametric two-sample procedure for comparing survival quantiles.

**Encoding** UTF-8

**Imports** survival (>= 2.41), graphics (>= 3.4.0), stats (>= 3.4.0)

**LazyData** true

**License** GPL-3

**RoxygenNote** 6.0.1

**Repository** <https://erickawaguchi.r-universe.dev>

**RemoteUrl** <https://github.com/erickawaguchi/controltest>

**RemoteRef** HEAD

**RemoteSha** 9ee6b25300edf000430a876e8c909043f20542c7

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quantileControlTest *Find standard error for survival quantile*

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### Description

Find standard error for survival quantile

### Usage

```
quantileControlTest(timevar1, censor1, timevar2, censor2, q = 0.5, B = 1000,  
  seed = 1234, plots = FALSE)
```

### Arguments

timevar1	Vector of observed survival times for sample 1 (control).
censor1	Vector of censoring indicators for sample 1 (1 = uncensored, 0 = censored).
timevar2	Vector of observed survival times for sample 2 (treatment).
censor2	Vector of censoring indicators for sample 2 (1 = uncensored, 0 = censored).
q	Quantile of interest (in terms of CDF). Default is median.
B	Number of bootstrap samples.
seed	Seed number (for reproducibility).
plots	Logical. TRUE to show plot of cumulative distribution functions.

### Details

It is important to note the possibility that the estimated quantile may not be estimable in our bootstrap samples. In such cases the largest observed survival time will be considered as an estimate for the quantile.

### Value

Returns quantile estimate, bootstrapped standard error, test statistic, and two-sided p-value.

### References

- Li, G., Tiwari, R.C., and Wells, M. (1996). "Quantile Comparison Functions in Two-Sample Problems: With Applications to Comparisons of Diagnostic Markers." *Journal of the American Statistical Association*, 91, 689-698.
- Chakraborti, S., and Mukerjee, R. (1989), "A Confidence Interval for a Measure Associated With the Comparison of a Treatment With a Control," *South African Statistical Journal*, 23, 219-230.
- Gastwirth, J. L., and Wang, J. L. (1988), "Control Percentile Test for Censored Data," *Journal of Statistical Planning and Inference*, 18, 267-276.

**Examples**

```
#Reference: Survival Analysis Techniques for Censored and Truncated Data.
#Klein and Moeschberger (1997) Springer.
#Data: Chapter 7.6 Example 7.9 (p. 211)
library(controlTest)
t1 <- c(1, 63, 105, 129, 182, 216, 250, 262, 301, 301,
        342, 354, 356, 358, 380, 383, 383, 338, 394, 408, 460, 489,
        499, 523, 524, 535, 562, 569, 675, 676, 748, 778, 786, 797,
        955, 968, 1000, 1245, 1271, 1420, 1551, 1694, 2363, 2754, 2950)
t2 <- c(17, 42, 44, 48, 60, 72, 74, 95, 103, 108, 122, 144, 167, 170,
        183, 185, 193, 195, 197, 208, 234, 235, 254, 307, 315, 401, 445,
        464, 484, 528, 542, 547, 577, 580, 795, 855, 1366, 1577, 2060,
        2412, 2486, 2796, 2802, 2934, 2988)
c1 <- c(rep(1, 43), 0, 0)
c2 <- c(rep(1, 39), rep(0, 6))
quantileControlTest(t1, c1, t2, c2, q = 0.5, B = 500)
```

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 quantileSE

*Find standard error for survival quantile*


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**Description**

Find standard error for survival quantile

**Usage**

```
quantileSE(timevar, censor, q = 0.5, B = 1000, alpha = 0.05,
  seed = 1991, plots = FALSE)
```

**Arguments**

timevar	Vector of observed survival times.
censor	Vector of censoring indicators (1 = uncensored, 0 = censored).
q	Quantile of interest (Default is median).
B	Number of bootstrap samples.
alpha	Significance level for confidence interval of quantile.
seed	Seed number (for reproducibility).
plots	Logical. TRUE to show Kaplan-Meier plot

**Value**

Returns quantile estimate, bootstrapped standard error, and  $(1 - \alpha / 2) * 100$

**Examples**

```
#Reference: Survival Analysis Techniques for Censored and Truncated Data.
#Klein and Moeschberger (1997) Springer.
#Data: Chapter 7.6 Example 7.9 (p. 211)
library(controlTest)
t1 <- c(1, 63, 105, 129, 182, 216, 250, 262, 301, 301,
        342, 354, 356, 358, 380, 383, 383, 338, 394, 408, 460, 489,
        499, 523, 524, 535, 562, 569, 675, 676, 748, 778, 786, 797,
        955, 968, 1000, 1245, 1271, 1420, 1551, 1694, 2363, 2754, 2950)
c1 <- c(rep(1, 43), 0, 0)
quantileSE(timevar = t1, censor = c1, q = 0.5, B = 500)
```

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<code>supControlTest</code>	<i>Supremum-type test for two-sample comparison of survival quantiles.</i>
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**Description**

Supremum-type test for two-sample comparison of survival quantiles.

**Usage**

```
supControlTest(timevar1, censor1, timevar2, censor2, q.min = NULL,
               q.max = 0.5, gridpts = 50, B = 500, seed = 1234, plots = FALSE)
```

**Arguments**

<code>timevar1</code>	Vector of observed survival times for sample 1 (control).
<code>censor1</code>	Vector of censoring indicators for sample 1 (1 = uncensored, 0 = censored).
<code>timevar2</code>	Vector of observed survival times for sample 2 (treatment).
<code>censor2</code>	Vector of censoring indicators for sample 2 (1 = uncensored, 0 = censored).
<code>q.min</code>	Smallest quantile (in terms of CDF) to test. Default is the time to earliest event for sample 1.
<code>q.max</code>	Largest quantile (in terms of CDF) to test.
<code>gridpts</code>	Number of grid points between <code>q.min</code> and <code>q.max</code> to test.
<code>B</code>	Number of bootstrap samples.
<code>seed</code>	Seed number (for reproducibility).
<code>plots</code>	Logical. TRUE to show plot of cumulative distribution functions.

**Details**

It is important to note the possibility that the estimated quantile may not be estimable in our bootstrap samples. In such cases the largest observed survival time will be considered as an estimate for the quantile.

**Value**

Returns quantile estimate, bootstrapped standard error, test statistic, and two-sided p-value.

**References**

Li, G., Tiwari, R.C., and Wells, M. (1996). "Quantile Comparison Functions in Two-Sample Problems: With Applications to Comparisons of Diagnostic Markers." *Journal of the American Statistical Association*, 91, 689-698.

Chakraborti, S., and Mukerjee, R. (1989), "A Confidence Interval for a Measure Associated With the Comparison of a Treatment With a Control," *South African Statistical Journal*, 23, 219-230.

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**Examples**

```
#Reference: Survival Analysis Techniques for Censored and Truncated Data.
#Klein and Moeschberger (1997) Springer.
#Data: Chapter 7.6 Example 7.9 (p. 211)
library(controlTest)
t1 <- c(1, 63, 105, 129, 182, 216, 250, 262, 301, 301,
        342, 354, 356, 358, 380, 383, 383, 338, 394, 408, 460, 489,
        499, 523, 524, 535, 562, 569, 675, 676, 748, 778, 786, 797,
        955, 968, 1000, 1245, 1271, 1420, 1551, 1694, 2363, 2754, 2950)
t2 <- c(17, 42, 44, 48, 60, 72, 74, 95, 103, 108, 122, 144, 167, 170,
        183, 185, 193, 195, 197, 208, 234, 235, 254, 307, 315, 401, 445,
        464, 484, 528, 542, 547, 577, 580, 795, 855, 1366, 1577, 2060,
        2412, 2486, 2796, 2802, 2934, 2988)
c1 <- c(rep(1, 43), 0, 0)
c2 <- c(rep(1, 39), rep(0, 6))
supControlTest(t1, c1, t2, c2, q.max = 0.5, B = 500)
```

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