

Package ‘Temporal’

October 7, 2018

Title Parametric Time to Event Analysis

Version 0.1.1

Description

Performs likelihood-based estimation and inference on time to event data, possibly subject to non-informative right censoring. `fitParaSurv()` provides maximum likelihood estimates of model parameters and distributional characteristics. `compParaSurv()` compares the mean and median survival experiences of two treatment arms. Candidate distributions currently include the exponential, gamma, generalized gamma, log-logistic, log-normal, and Weibull.

Depends R (>= 3.4.0)

License GPL-3

Encoding UTF-8

LazyData true

LinkingTo Rcpp, RcppEigen

Imports expint, methods, numDeriv, plyr, Rcpp, stats

RoxygenNote 6.1.0

Suggests knitr, rmarkdown

VignetteBuilder knitr

NeedsCompilation yes

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Repository CRAN

Date/Publication 2018-10-07 18:32:20 UTC

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compParaSurv	<i>Compare Parametric Survival Distribution</i>
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Description

Compares the means and medians of parametric survival distributions fit to each of two treatment arms. Available distributions include: exponential, gamma, generalized gamma, log-logistic, log-normal, and Weibull.

Usage

```
compParaSurv(time, status, arm, dist1 = "weibull", dist0, init1 = NULL,
             init0 = NULL, sig = 0.05, eps = 1e-06, maxit = 10, report = F)
```

Arguments

time	Observation time.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
arm	Treatment indicator, coded as 1 for the target group, 0 for the reference group.
dist1	Distribution to fit for the target group. Selected from among: exp, gamma, gengamma, log-logistic, log-normal, and weibull.
dist0	Distribution to fit for the reference group. Same choices as for the target group. If omitted, defaults to the distribution specified for the target group.
init1	List of initial parameter values for the target group. See individual distributions for naming convention.
init0	List of initial parameter values for the reference group. See individual distributions for naming convention.
sig	Significance level, for CIs.
eps	Tolerance for Newton-Raphson iterations.
maxit	Maximum number of NR iterations.
report	Report fitting progress?

Value

An object of class `contrast` containing the following:

Model1 The fitted model for the target group.

Model0 The fitted model for the reference group.

Contrast Contrasts of means and medians.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
set.seed(100);
## Weibull and Weibull, different means and medians
# Target group
D1 = rWeibull(n=1e3, a=1, l=1, p=0.2);
D1$arm = 1;
# Reference group
D0 = rWeibull(n=1e3, a=1, l=2, p=0.2);
D0$arm = 0;
# Overall data set
D = rbind(D1, D0);
# Comparison
E = compParaSurv(time=D$time, status=D$status, arm=D$arm, dist1="weibull");

## Gamma and Weibull, different means and medians
# Target group
D1 = rGamma(n=1e3, a=2, l=2, p=0.2);
```

```

D1$arm = 1;
# Reference group
D0 = rWeibull(n=1e3,a=2,l=2/sqrt(pi),p=0.2);
D0$arm = 0;
# Overall data set
D = rbind(D1,D0);
# Comparison
E = compParaSurv(time=D$time,status=D$status,arm=D$arm,dist1="gamma",dist0="weibull");

## Weibull and Log-normal, different means, same median
# Target group
D1 = rLogNormal(n=1e3,m=0,s=2,p=0.2);
D1$arm = 1;
# Reference group
D0 = rWeibull(n=1e3,a=2,l=sqrt(log(2)),p=0.2);
D0$arm = 0;
# Overall data set
D = rbind(D1,D0);
# Comparison
E = compParaSurv(time=D$time,status=D$status,arm=D$arm,dist1="log-normal",dist0="weibull");

```

contrast-class	<i>Contrast of Survival Distributions.</i>
----------------	--

Description

Defines the object class returned by the comparison function.

Slots

Dist1 Distribution fit to the target group.
 Dist0 Distribution fit to the reference group.
 Model1 Fitted model for the target group.
 Model0 Fitted model for the reference group.
 Contrast Model contrasts.

det	<i>Matrix Determinant</i>
-----	---------------------------

Description

Calculates the determinant of A .

Usage

```
det(A)
```

Arguments

A Numeric matrix.

Value

Scalar.

dIncGamma

Derivatives of the Incomplete Gamma Function

Description

Evaluates derivatives of the upper incomplete gamma function, defined as:

$$\Gamma(\alpha, s) = \int_s^{\infty} u^{\alpha-1} e^{-u} du$$

Usage

```
dIncGamma(a, s, Dirn = "a", Order = 1)
```

Arguments

a Value of the shape α at which to evaluate.

s Value of the lower limit s at which to evaluate.

Dirn Direction in which to differentiate. Select from among "a", "s", and "as".

Order Order of the derivative, if the direction is either "a", or "s". Select from among 1 and 2.

Value

Scalar value of the partial derivative in the direction of interest.

Examples

```
# First partial in shape at (a=1,s=1)
dIncGamma(a=1,s=1,Dirn="a",Order=1);
# Second partial in lower limit at (a=1,s=1)
dIncGamma(a=1,s=1,Dirn="s",Order=2);
# Mixed partial at (a=1,s=1);
dIncGamma(a=1,s=1,Dirn="as");
```

dLogIncGamma

Derivatives of the Log Incomplete Gamma Function

Description

Evaluates derivatives of the log of the upper log incomplete gamma function, defined as:

$$\ln \Gamma(\alpha, s) = \int_s^{\infty} u^{\alpha-1} e^{-u} du$$

Usage

dLogIncGamma(a, s, Dirn = "a", Order = 1)

Arguments

a	Value of the shape α at which to evaluate.
s	Value of lower limit s at which to evaluate.
Dirn	Direction in which to differentiate. Select from among "a", "s", and "as".
Order	Order of the derivative, if the direction is either "a", or "s". Select from among 1 and 2.

Value

Scalar value of the partial derivative in the direction of interest.

Examples

```
# First partial in shape at (a=1,s=1)
dLogIncGamma(a=1,s=1,Dirn="a",Order=1);
# Second partial in lower limit at (a=1,s=1)
dLogIncGamma(a=1,s=1,Dirn="s",Order=2);
# Mixed partial at (a=1,s=1);
dLogIncGamma(a=1,s=1,Dirn="as");
```

dLogNormSurv

Logarithmic Derivatives of the Normal Survival Function

Description

Evaluates logarithmic derivatives of the normal survival, defined as:

$$\ln \Phi(s) = \ln \int_s^{\infty} \frac{e^{-(u^2)/2}}{\sqrt{2\pi}} du$$

Usage

```
dLogNormSurv(s, Order = 1)
```

Arguments

`s` Value of the lower limit s at which to evaluate.
`Order` Order of the derivative, select from among 1 and 2.

Value

Scalar value of the derivative.

estDiff	<i>Difference of Estimates</i>
---------	--------------------------------

Description

Calculate CIs and p-value for the difference of estimated parameters

Usage

```
estDiff(t1, s1, t0, s0, sig = 0.05)
```

Arguments

`t1` Treatment estimate
`s1` Treatment standard error
`t0` Reference estimate
`s0` Reference standard error
`sig` Significance level

Value

Data.frame containing estimated difference, its standard error, lower and upper confidence bounds, and a p-value assessing the null hypothesis of no difference.

estRatio	<i>Ratio of Estimates</i>
----------	---------------------------

Description

Calculate CIs and p-value for the ratio of estimated parameters

Usage

```
estRatio(t1, s1, t0, s0, sig = 0.05)
```

Arguments

t1	Treatment estimate
s1	Treatment standard error
t0	Reference estimate
s0	Reference standard error
sig	Significance level

Value

Data.frame containing estimated ratio, its standard error, lower and upper confidence bounds, and a p-value assessing the null hypothesis that the ratio is unity.

fit-class	<i>Fitted Survival Distribution</i>
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Description

Defines the object class returned by fitting functions.

Slots

Distribution	Fitted distribution.
Parameters	Parameters.
Information	Information components.
Outcome	Properties of the fitted distribution.

Description

Estimates parameters for exponential event times subject to non-informative right censoring. The exponential distribution is parameterized in terms of the rate λ :

$$f(t) = \lambda e^{-\lambda t}, t > 0$$

Usage

```
fit.Exp(time, status, sig = 0.05)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
sig	Significance level, for CIs.

Value

An object of class `fit` containing the following:

Parameters The estimated model parameters.

Information The observed information matrix.

Outcome The fitted mean, median, and variance of the time to event distribution.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
# Simulate
D = rWeibull(n=1e3, a=1, l=2);
# Estimate
M = fitParaSurv(time=D$time, status=D$status, dist="exp");
```

Description

Estimates parameters for gamma event times subject to non-informative right censoring. The gamma distribution is parameterized in terms of the shape α and rate λ :

$$f(t) = \frac{\lambda}{\Gamma(\alpha)} (\lambda t)^{\alpha-1} e^{-\lambda t}, t > 0$$

Usage

```
fit.Gamma(time, status, sig = 0.05, init = NULL, eps = 1e-06,
          maxit = 10, report = F)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
sig	Significance level, for CIs.
init	List of initial parameter values, including the log of the shape parameter "la" and the log of the rate parameter "ll".
eps	Tolerance for Newton-Raphson iterations.
maxit	Maximum number of NR iterations.
report	Report fitting progress?

Value

An object of class `fit` containing the following:

Parameters The estimated shape α and rate λ .

Information The observed information matrix.

Outcome The fitted mean, median, and variance.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
# Simulate
D = rGamma(n=1e3, a=2, l=2);
# Estimate
M = fitParaSurv(time=D$time, status=D$status, dist="gamma");
```

fit.GenGamma

*Generalized Gamma Distribution Parameter Estimation***Description**

Estimates parameters for generalized gamma event times subject to non-informative right censoring. The gamma distribution is parameterized in terms of the shape parameters (α, β) , and the rate λ :

$$f(t) = \frac{\beta\lambda}{\Gamma(\alpha)} (\lambda t)^{\alpha\beta-1} e^{-(\lambda t)^\beta}, t > 0$$

Usage

```
fit.GenGamma(time, status, sig = 0.05, init = NULL, eps = 1e-06,
             maxit = 10, report = F)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
sig	Significance level, for CIs.
init	List of initial parameter values, including the logs of the shape parameters "la" and "lb", and the log of the rate parameter "ll".
eps	Tolerance for Newton-Raphson iterations.
maxit	Maximum number of NR iterations.
report	Report fitting progress?

Value

An object of class `fit` containing the following:

Parameters The estimated shape (α, β) and rate λ parameters.

Information The observed information matrix.

Outcome The fitted mean, median, and variance.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
set.seed(100);
# Simulate
D = rGenGamma(n=1e4, a=2, b=2, l=2);
# Estimate
M = fitParaSurv(time=D$time, status=D$status, dist="gengamma");
```

fit.LogLogistic

*Log-Logistic Distribution Parameter Estimation***Description**

Estimates parameters for log-logistic event times subject to non-informative right censoring. The log-logistic distribution is parameterized in terms of the shape α and rate λ :

$$f(t) = \frac{\alpha\lambda(\lambda t)^{\alpha-1}}{[1 + (\lambda t)^\alpha]^2}, t > 0$$

Usage

```
fit.LogLogistic(time, status, sig = 0.05, init = NULL, eps = 1e-06,
  maxit = 10, report = F)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
sig	Significance level, for CIs.
init	List of initial parameter values, including the log of the shape parameter "la" and the log of the rate parameter "ll".
eps	Tolerance for Newton-Raphson iterations.
maxit	Maximum number of NR iterations.
report	Report fitting progress?

Details

For the log-logistic distribution, the mean is only defined if the shape parameter $\alpha > 1$, and the variance if the shape parameter $\alpha > 2$. Consequently, estimates of the fitted mean and variance are only returned if the estimated shape parameter exceeds these thresholds. For $\alpha \ll 1$, the fitting function may fail.

Value

An object of class `fit` containing the following:

Parameters The estimated location μ and scale σ .

Information The observed information matrix.

Outcome The fitted mean, median, and variance.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
# Simulate
D = rLogLogistic(n=1e3,a=4,l=2);
# Estimate
M = fitParaSurv(time=D$time,status=D$status,dist="log-logistic");
```

fit.LogNormal

*Log-Normal Distribution Parameter Estimation***Description**

Estimates parameters for log-normal event times subject to non-informative right censoring. The log-normal distribution is parameterized in terms of the location μ and scale σ :

$$f(t) = \phi\left(\frac{\ln t - \mu}{\sigma}\right) \frac{1}{t\sigma}, t > 0$$

Usage

```
fit.LogNormal(time, status, sig = 0.05, init = NULL, eps = 1e-06,
maxit = 10, report = F)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
sig	Significance level, for CIs.
init	List of initial parameter values, including the location "m", and the log of the scale parameter "ls".
eps	Tolerance for Newton-Raphson iterations.
maxit	Maximum number of NR iterations.
report	Report fitting progress?

Value

An object of class `fit` containing the following:

Parameters The estimated location μ and scale σ .

Information The observed information matrix.

Outcome The fitted mean, median, and variance.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
# Simulate
D = rLogNormal(n=1e3,m=0,s=1);
# Estimate
M = fitParaSurv(time=D$time,status=D$status,dist="log-normal");
```

fit.Weibull

*Weibull Distribution Parameter Estimation***Description**

Estimates parameters for Weibull event times subject to non-informative right censoring. The Weibull distribution is parameterized in terms of the shape α and rate λ :

$$f(t) = \alpha\lambda^\alpha t^{\alpha-1} e^{-(\lambda t)^\alpha}, t > 0$$

Usage

```
fit.Weibull(time, status, sig = 0.05, init = NULL)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
sig	Significance level, for CIs.
init	List containing an initial value for the shape parameter "a".

Value

An object of class `fit` containing the following:

Parameters The estimated shape α and rate λ .

Information The observed information matrix.

Outcome The fitted mean, median, and variance.

See Also

- Fitting function for parametric survival distributions [fitParaSurv](#)

Examples

```
# Simulate
D = rWeibull(n=1e3,a=2,l=2);
# Estimate
M = fitParaSurv(time=D$time,status=D$status,dist="weibull");
```

fitParaSurv

*Fit Parametric Survival Distribution***Description**

Estimates parametric survival distributions using event times subject to non-informative right censoring. Available distributions include: exponential, gamma, generalized gamma, log-logistic, log-normal, and Weibull.

Usage

```
fitParaSurv(time, status, dist = "weibull", sig = 0.05, init = NULL,
            eps = 1e-06, maxit = 10, report = F)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
dist	Distribution to fit, selected from among: exp, gamma, gengamma, log-logistic, log-normal, and weibull.
sig	Significance level, for CIs.
init	List of initial parameter values. See individual distributions for naming convention.
eps	Tolerance for Newton-Raphson iterations.
maxit	Maximum number of NR iterations.
report	Report fitting progress?

Value

An object of class `fit` containing the following:

Parameters The estimated shape and rate parameters.

Information The observed information matrix.

Outcome The fitted mean, median, and variance.

See Also

- Between group comparison of survival experience [compParaSurv](#)
- Exponential distribution [fit.Exp](#)
- Gamma distribution [fit.Gamma](#)
- Generalized gamma distribution [fit.GenGamma](#)
- Log-logistic distribution [fit.LogLogistic](#)
- Log-normal distribution [fit.LogNormal](#)
- Weibull distribution [fit.Weibull](#)

Examples

```
# Generate censored gamma data
D = rGamma(n=1e3,a=2,l=2,p=0.2);
# Fit gamma distribution
M = fitParaSurv(time=D$time,status=D$status,dist="gamma");

# Generate censored weibull data
D = rWeibull(n=1e3,a=2,l=2,p=0.2);
# Fit weibull distribution
M = fitParaSurv(time=D$time,status=D$status,dist="weibull");
```

init.GenGamma	<i>Initialization for Generalized Gamma</i>
---------------	---

Description

Initializes the parameters for the generalized gamma distribution using moment estimators.

Usage

```
init.GenGamma(tobs, L = 0.01, U = 10)
```

Arguments

tobs	Observed event times.
L	Lower bound on shape parameter α .
U	Upper bound on shape parameter α .

Value

List containing the estimated shape and rate parameters on log scale.

matInv	<i>Matrix Inverse</i>
--------	-----------------------

Description

Calculates A^{-1} .

Usage

```
matInv(A)
```

Arguments

A	Numeric matrix.
---	-----------------

Value

Numeric matrix.

matIP	<i>Matrix Inner Product</i>
-------	-----------------------------

Description

Calculates the product $A'B$.

Usage

matIP(A, B)

Arguments

A	Numeric matrix.
B	Numeric matrix.

Value

Numeric matrix.

matOP	<i>Fast Outer Product</i>
-------	---------------------------

Description

Calculates the outer product XY' .

Usage

matOP(X, Y)

Arguments

X	Numeric matrix.
Y	Numeric matrix.

Value

Numeric matrix.

matQF

Quadratic Form

Description

Calculates the quadratic form $X'AX$.

Usage

matQF(X, A)

Arguments

X Numeric matrix.

A Numeric matrix.

Value

Numeric matrix.

MMP

Matrix Matrix Product

Description

Calculates the product AB .

Usage

MMP(A, B)

Arguments

A Numeric matrix.

B Numeric matrix.

Value

Numeric matrix.

print.contrast	<i>Print Method for a Contrast of Survival Distributions.</i>
----------------	---

Description

Print method for an object of class contrast.

Usage

```
## S3 method for class 'contrast'  
print(x, ...)
```

Arguments

x	A contrast object.
...	Unused.

print.fit	<i>Print Method for Fitted Survival Distributions</i>
-----------	---

Description

Print method for objects of class fit.

Usage

```
## S3 method for class 'fit'  
print(x, ...)
```

Arguments

x	An object of class fit.
...	Unused.

`qLogLogistic`*Quantile Function for the Log-Logistic Distribution*

Description

Quantile function for the log-logistic distribution. See [fit.LogLogistic](#) for the parameterization.

Usage

```
qLogLogistic(p, a = 1, l = 1)
```

Arguments

<code>p</code>	Probability.
<code>a</code>	Shape.
<code>l</code>	Rate.

Value

Scalar quantile.

Examples

```
# Median of standard log-logistic distribution  
qLogLogistic(p=0.5);
```

`qWeibull`*Quantile Function for the Weibull Distribution*

Description

Quantile function for the Weibull distribution. See [fit.Weibull](#) for the parameterization.

Usage

```
qWeibull(p, a = 1, l = 1)
```

Arguments

<code>p</code>	Probability.
<code>a</code>	Shape.
<code>l</code>	Rate.

Value

Scalar quantile.

rGamma

Simulation from the Gamma Distribution

Description

Generates gamma event times with shape parameter α and rate parameter λ . See [fit.Gamma](#) for the parameterization. If a censoring proportion p is provided, the event times are subject to non-informative random right censoring.

Usage

```
rGamma(n, a = 1, l = 1, p = 0)
```

Arguments

n	Sample size.
a	Shape.
l	Rate.
p	Expected censoring proportion.

Value

A data.frame including the observation times and status indicators.

Examples

```
# Gamma event times with shape 2 and rate 2
# Expected censoring proportion of 20%
D = rGamma(n=1e3,a=2,l=2,p=0.2);
```

rGenGamma

Simulation from the Generalized Gamma Distribution

Description

Generates generalized gamma event times with shape parameters (α, β) , and rate parameter λ . See [fit.GenGamma](#) for the parameterization. If a censoring proportion p is provided, the event times are subject to non-informative random right censoring.

Usage

```
rGenGamma(n, a = 1, b = 1, l = 1, p = 0)
```

Arguments

n	Sample size.
a	First shape parameter, α .
b	Second shape parameter, β . For the standard gamma distribution, set $\beta = 1$.
l	Rate.
p	Expected censoring proportion.

Value

A data.frame including the observation times and status indicators.

Examples

```
# Generalized gamma event times with shapes (2,2) and rate 2
# Expected censoring proportion of 20%
D = rGenGamma(n=1e3, a=2, b=2, l=2, p=0.2);
```

rLogLogistic

Simulation from the Log-Logistic Distribution

Description

Generates log-logistic event times with shape parameter α and rate parameter λ . See [fit.LogLogistic](#) for the parameterization. If a censoring proportion p is provided, the event times are subject to non-informative random right censoring.

Usage

```
rLogLogistic(n, a = 4, l = 1, p = 0)
```

Arguments

n	Sample size.
a	Shape.
l	Rate.
p	Expected censoring proportion.

Value

A data.frame including the observation times and status indicators.

Examples

```
# Log-logistic event times with shape 4 and rate 1
# Expected censoring proportion of 20%
D = rLogLogistic(n=1e3, a=4, l=1, p=0.2);
```

`rLogNormal`*Simulation from the Log-Normal Distribution*

Description

Generates log-normal event times with location parameter μ and scale parameter σ . See [fit.LogNormal](#) for the parameterization. If a censoring proportion p is provided, the event times are subject to non-informative random right censoring.

Usage

```
rLogNormal(n, m = 0, s = 1, p = 0)
```

Arguments

<code>n</code>	Sample size.
<code>m</code>	Location.
<code>s</code>	Scale.
<code>p</code>	Expected censoring proportion.

Value

A data.frame including the observation times and status indicators.

Examples

```
# Log-normal event times with location 0 and scale 1
# Expected censoring proportion of 20%
D = rLogNormal(n=1e3,m=0,s=1,p=0.2);
```

`rWeibull`*Simulation from the Weibull Distribution*

Description

Generates Weibull event times with shape parameter α and rate parameter λ . See [fit.Weibull](#) for the parameterization. If a censoring proportion p is provided, the deviates are subject to non-informative random right censoring.

Usage

```
rWeibull(n, a = 1, l = 1, p = 0)
```

Arguments

n	Sample size.
a	Shape.
l	Rate.
p	Expected censoring proportion.

Value

A data.frame including the observation times and status indicators.

Examples

```
# Weibull event times with shape 2 and rate 2
# Expected censoring proportion of 20%
D = rWeibull(n=1e3, a=2, l=2, p=0.2);
```

 SchurC

Schur complement

Description

Calculates the efficient information $I_{bb} - I_{ba}I_{aa}^{-1}I_{ab}$.

Usage

```
SchurC(Ibb, Iaa, Iba)
```

Arguments

Ibb	Information of target parameter
Iaa	Information of nuisance parameter
Iba	Cross information between target and nuisance parameters

Value

Numeric matrix.

show,contrast-method *Show Method for a Contrast of Survival Distributions.*

Description

Show Method for a Contrast of Survival Distributions.

Usage

```
## S4 method for signature 'contrast'  
show(object)
```

Arguments

object An object of class contrast.

show,fit-method *Show Method for Fitted Survival Distributions*

Description

Show Method for Fitted Survival Distributions

Usage

```
## S4 method for signature 'fit'  
show(object)
```

Arguments

object An object of class fit.

survLogLik	<i>Survival Distribution Log Likelihood</i>
------------	---

Description

Evaluation of the log-likelihood for parametric survival distribution.

Usage

```
survLogLik(time, status, dist, theta, log.scale = T)
```

Arguments

time	Observation times.
status	Status indicator, coded as 1 if an event was observed, 0 if censored.
dist	Distribution.
theta	List of parameters, which will vary according to the distribution.
log.scale	Are positive parameters on log-scale?

Value

Scalar value of the log likelihood.

Examples

```
## Not run:
# Simulate
D = rWeibull(n=1e3, a=2, l=2, p=0.2);
# Log likelihood
ll = survLogLik(time=D$time, status=D$status,
  dist="weibull", theta=list("a"=2, "l"=2), log.scale=F);

## End(Not run)
```

Temporal	<i>Temporal: Utilities for Parametric Survival Analysis</i>
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Description

This package performs estimation and inference on parametric survival curves. See [fitParaSurv](#) for maximum likelihood estimation of model parameters. See [compParaSurv](#) for contrasting the survival experience of two treatment arms.

Author(s)

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tr

Matrix Trace

Description

Calculates the trace of a matrix A .

Usage

$\text{tr}(A)$

Arguments

A Numeric matrix.

Value

Scalar.

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