

Package ‘supportInt’

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Type Package

Title Calculates Likelihood Support Intervals for Common Data Types

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Description Calculates likelihood based support intervals for several common data types including binomial, Poisson, normal, lm(), and glm(). For the binomial, Poisson, and normal data likelihood intervals are calculated via root finding algorithm. Additional parameters allow the user to specify whether they would like to receive a parametric bootstrap estimate of the confidence level of said support interval. For lm() and glm(), the function returns profile likelihoods for each coefficient in the model.

Imports ProfileLikelihood, splines

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supportInt-package *Calculates Likelihood Support Intervals for Common Data Types*

Description

Calculates likelihood based support intervals for several common data types including binomial, Poisson, normal, lm(), and glm(). For the binomial, Poisson, and normal data likelihood intervals are calculated via root finding algorithm. Additional parameters allow the user to specify whether they would like to receive a parametric bootstrap estimate of the confidence level of said support interval. For lm() and glm(), the function returns profile likelihoods for each coefficient in the model.

Author(s)

Derek Smith

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Examples

```
#Binomial data
binLikSI(4, 10, 8)

#Poisson data returning the estimated confidence level
poisLikSI(4, 8, conf=TRUE)

#GLM coefficients
set.seed(10)
x <- rnorm(50, 0 , 5)
expit <- function(z) exp(z)/(1+exp(z))
p <- expit(.1+ .4*x)
y <- sapply(1:length(p), function(z) rbinom(1,1, p[z]))
glm.obj <- glm(y~x, family="binomial")
glmLikSI(glm.obj, 8)
```

binLikSI

Binomial support interval calculation

Description

This function takes binomial data and using a root finding algorithm to calculate the support interval at the user's specified level. If conf=T, the function will attempt to estimate the confidence level of the support interval using a modified parametric bootstrap approach.

Usage

```
binLikSI(dat, n, level, tol = 0.001, conf = F, B = 500)
```

Arguments

<code>dat</code>	number of success in binomial experiment
<code>n</code>	number of trials in binomial experiment
<code>level</code>	desired support level
<code>tol</code>	tolerance passed to uniroot for finding SI endpoints
<code>conf</code>	Logical, whether or not bootstrap should be performed to estimate confidence level
<code>B</code>	Number of bootstraps to be performed

Value

<code>si</code>	The desired support interval
<code>conf.equiv</code>	The estimated confidence level of the SI

Author(s)

Derek Smith

Examples

```
set.seed(10)
binLikSI(dat=4, n=10, level=9)
binLikSI(dat=4, n=10, level=8, conf=TRUE)
```

`calibSI`

Attempts to find a support interval with the desired confidence level for binomial, poisson or normal data.

Description

This function performs a grid search to find the lowest support level that has the desired coverage probability. It begins by searching a grid of relative likelihood values using a fixed number of parametric bootstraps at each point. The likelihood value is then refined using a golden section based search.

Usage

```
calibSI(dat, n = NULL, family, conf.level = 0.95, B = 2000,
gridlo = 4, gridhi = 20, gridix = 2, tol = 0.03)
```

Arguments

<code>dat</code>	This argument is the number of success if family='binomial', the number of occurrences if family='poisson', or a vector of normal draws if family='normal'.
<code>n</code>	Number of trials if family='binomial'
<code>family</code>	Data type entered, accepts "binomial", "poisson", "pois", "gaussian", "normal"
<code>conf.level</code>	target confidence level
<code>B</code>	Number of parametric bootstraps to use to estimate coverage probability after the initial grid search.
<code>gridlo</code>	Lowest relative likelihood considered in the initial grid search
<code>gridhi</code>	Highest relative likelihood considered in the initial grid search
<code>gridix</code>	size of steps to be checked between gridlo and gridhi in initial search
<code>tol</code>	acceptable error in the likelihood support level returned

Value

<code>si</code>	The support interval estimated to have the desired confidence level
<code>support.level</code>	The relative likelihood defining the interval's endpoints
<code>init.grid</code>	Relative likelihoods checked as part of the initial gridsearch and their estimated coverage

Author(s)

Derek Smith

Examples

```
set.seed(10)
calibSI(rnorm(15), family="normal", conf.level=.95, B=1000)
```

`glmLikSI`

Profile support intervals for glm coefficients

Description

Calculates profile likelihood intervals using data returned from the 'ProfileLikelihood' package for each coefficient of a `glm`.

Usage

```
glmLikSI(glm.mod, level)
```

Arguments

<code>glm.mod</code>	A fit <code>glm</code> object.
<code>level</code>	Desired level of support interval (1/level).

Value

a matrix of support intervals for each coefficient

Author(s)

Derek Smith

Examples

```
set.seed(10)
x <- rnorm(50, 0 , 5)
expit <- function(z) exp(z)/(1+exp(z))
p <- expit(.1+ .4*x)
y <- sapply(1:length(p), function(z) rbinom(1,1, p[z]))
glm.obj <- glm(y~x, family="binomial")
glmLikSI(glm.obj, 8)
```

lmLikSI

Profile support intervals for lm coefficients

Description

Calculates profile likelihood intervals using data returned from the 'ProfileLikelihood' package for each coefficient of a lm.

Usage

```
lmLikSI(lm.mod, level)
```

Arguments

lm.mod	A fit lm object.
level	Desired level of support interval (1/level).

Value

a matrix of support intervals for each coefficient

Author(s)

Derek Smith

Examples

```
set.seed(10)
x <- rnorm(50, 0 , 5)
y <- sapply(1:length(x), function(z) 3+.5*x[z]+rnorm(1, 0, 5))
lm.obj <- lm(y~x)
lmLikSI(lm.obj, 8)
```

normLikSI

*Normal support interval calculation.***Description**

This function calculates support intervals for the mean of normal data. Confidence levels are calculate from the formula if conf=T.

Usage

```
normLikSI(dat, level, tol = 0.001, conf = F, B=500)
```

Arguments

dat	a numeric vector
level	support interval relative likelihood
tol	tolerance passed to uniroot for finding likelihood endpoints
conf	Logical, whether or not confidence levels should be returned.
B	Number of bootstrap iterations to verify coverage level

Value

si	1/level support interval
conf.equiv	Calculated confidence level

Author(s)

Derek Smith

Examples

```
set.seed(10)
normLikSI(rnorm(10), 9)
normLikSI(rnorm(10), 9, conf=TRUE)
```

poisLikSI*Poisson support interval calculation*

Description

This function takes Poisson data and using a root finding algorithm to calculate the support interval at the user's specified level. If conf=T, the function will attempt to estimate the confidence level of the support interval using a parametric bootstrap approach.

Usage

```
poisLikSI(dat, level, tol = 0.001, conf = F, B = 500)
```

Arguments

dat	Numeric; outcome from a Poisson experiment
level	Desired support level
tol	tolerance argument passed to uniroot for finding SI endpoints
conf	Logical; Whether or not to simulate the confidence level of the SI
B	Number of simulations to determine coverage if conf=T

Value

si	1/level support interval
conf.equiv	estimated confidence level of interval returned

Author(s)

Derek Smith

Examples

```
poisLikSI(4, 8)
poisLikSI(4, 8, conf=TRUE)
```

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