# Package 'LBI'

## August 21, 2025

Version 0.2.3

Title Likelihood Based Inference

<b>Description</b> Maximum likelihood estimation and likelihood ratio test are essential for modern statics. This package supports in calculating likelihood based inference. Reference: Pawitan Y. (2001, ISBN:0-19-850765-8).	tis-
<b>Depends</b> R (>= $3.0.0$ )	
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LBI-package

Likelihood Based Inference

#### **Description**

It conducts likelihood based inference.

#### **Details**

Modern likelihood concept and maximum likelihood estimation are established by Fisher RA, while Likelihood Ratio Test (LRT) is established by Neyman J. Post-Fisher methods - generalized linear model, survival analysis, and mixed effects model - are all likelihood based. Inferences from the perspective of Fisherian and pure likelihoodist are suggested here.

#### Author(s)

Kyun-Seop Bae <k@acr.kr>

#### References

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- 2. Edwards AWF. Likelihood. 1972.
- 3. Fisher RA. Statistical Methods and Scientific Inference. 3e. 1973.
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- 6. Royall R. Statistical Evidence. 1997.
- 7. Pinheiro JC, Bates DM. Mixed Effects Models in S and S-PLUS. 2000.
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- 9. Lehmann EL. Fisher, Nayman, and the Creation of Classical Statistics. 2011.
- 10. Rohde CA. Introductory Statistical Inference with the Likelihood Function. 2014.
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LBCIvar

Likelihood Based Confidence Interval of sd and variance assuming Normal Distribution

#### Description

Likelihood based confidence interval of sd and variance assuming normal distribution. It usually shows narrower interval than conventional chi-square interval. This uses estimated likelihood, not profile likelihood.

#### Usage

LBCIvar(x, conf.level=0.95)

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#### **Arguments**

x a vector of observation conf.level confidence level

#### **Details**

It calculates (same height) likelihood based confidence interval of sd and variance assuming normal distribution in one group. The likelihood interval is asymmetric and there is no standard error in the output.

#### Value

PE	maximum likelihood estimate
LL	lower limit of likelihood interval
UL	upper limit of likelihood interval

#### Author(s)

Kyun-Seop Bae k@acr.kr

## **Examples**

```
LBCIvar(lh) (length(lh) - 1)*var(lh)/qchisq(c(0.975, 0.025), length(lh) - 1)
```

LIbin

Likelihood Interval for a Proportion or a Binomial Distribution

## **Description**

Likelihood interval of a proportion in one group

#### Usage

```
LIbin(y, n, k, conf.level=0.95, eps=1e-8)
```

#### **Arguments**

y positive event count of a group

n total count of a group

k 1/k likelihood interval will be calculated

conf.level approximately corresponding confidence level. If k is specified, this is ignored.

eps Values less than eps are considered as 0.

#### **Details**

It calculates likelihood interval of a proportion in one group. The likelihood interval is asymmetric and there is no standard error in the output. If you need percent scale, multiply the output by 100.

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#### Value

у	positive (concerning) event count
n	total trial count
PE	maximum likelihood estimate on the proportion
LL	lower limit of likelihood interval
UL	upper limit of likelihood interval

#### Author(s)

Kyun-Seop Bae k@acr.kr

#### References

Fisher RA. Statistical methods and scientific inference. 3e. 1973. pp68-76.

#### See Also

```
binom.test, prop.test
```

## **Examples**

```
LIbin(3, 14, k=2)

LIbin(3, 14, k=5)

LIbin(3, 14, k=15)

LIbin(3, 14)

# binom.test(3, 14)

# prop.test(3, 14)
```

LInorm

Likelihood Interval of mean, sd and variance assuming Normal Distribution

## Description

Likelihood interval of mean and sd assuming normal distribution. This is estimated likelihood interval, not profile likelihood interval.

## Usage

```
LInorm(x, k, conf.level=0.95, PLOT="", LOCATE=FALSE, Resol=201)
```

## Arguments

X	a vector of observation
k	1/k likelihood interval will be calculated
conf.level	approximately corresponding confidence level. If k is specified, this is ignored.
PLOT	"1d" for profile plot or "2d" for contour plot.
LOCATE	use locater. This works only with PLOT="2D" option.
Resol	resolution for plot. This works only with PLOT=TRUE option.

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#### **Details**

It calculates likelihood interval of mean and sd assuming normal distribution in one group. There is no standard error in the output.

#### Value

PE maximum likelihood estimate

LL lower limit of likelihood interval

UL upper limit of likelihood interval

#### Author(s)

Kyun-Seop Bae k@acr.kr

#### **Examples**

```
x = c(-5.3, -4.5, -1.0, -0.7, 3.7, 3.9, 4.2, 5.5, 6.8, 7.4, 9.3)

LInorm(x, k=1/0.15) # Pawitan Ex10-9 p289

LInorm(x)

LInorm(x, PLOT="1d")

LInorm(x, PLOT="2d", LOCATE=TRUE)
```

LInormVar

Likelihood Interval of sd and variance assuming Normal Distribution

#### **Description**

Likelihood interval of sd and variance assuming normal distribution. This is estimated likelihood interval, not profile likelihood interval.

## Usage

```
LInormVar(x, k, conf.level=0.95)
```

## **Arguments**

x a vector of observation

k 1/k likelihood interval will be calculated

conf.level approximately corresponding confidence level. If k is specified, this is ignored.

## **Details**

It calculates likelihood interval of sd and variance assuming normal distribution in one group. The likelihood interval is asymmetric and there is no standard error in the output.

#### Value

PE	maximum likelihood estimate
LL	lower limit of likelihood interval
UL	upper limit of likelihood interval

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#### Author(s)

Kyun-Seop Bae k@acr.kr

#### **Examples**

```
x = c(-5.3, -4.5, -1.0, -0.7, 3.7, 3.9, 4.2, 5.5, 6.8, 7.4, 9.3)
LInormVar(x, k=1/0.15) # Pawitan Ex10-9 p289
LInormVar(x)
```

LIpois

Likelihood Interval of the Mean assuming Poisson Distribution

## Description

Likelihood interval of lambda assuming Poisson distribution.

#### Usage

```
LIpois(x, k, n = 1, conf.level = 0.95, eps = 1e-8)
```

## Arguments

X	raw data vector or a mean value. If the length of x is 1, x is considered as a
	mean.
k	1/k likelihood interval will be calculated.
n	number of observations. If the length of x is 1, x is considered as the mean.
conf.level	approximately corresponding confidence level. If k is specified, this is ignored.
eps	estimated values less than this eps are considered as 0.

## **Details**

It calculates likelihood interval of mean(lambda) assuming Poisson distribution. The likelihood interval is asymmetric and there is no standard error in the output.

#### Value

PE	maximum likelihood estimate on the lambda
LL	lower limit of likelihood interval
UL	upper limit of likelihood interval
n	number of observations
k	1/k likelihood interval provided
logk	log(k) of k value
maxLL without fa	actorial
	maximum log likelihood without factorial part

#### Author(s)

Kyun-Seop Bae k@acr.kr

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#### **Examples**

```
LIpois(4, k=15)
                      # Fisher
  LIpois(4, k=exp(2)) # Edwards
  LIpois(4, k=1/0.15) # Pawitan
  LIpois(4, k=8)
                      # Rhode
 LIpois(4, n=4)
                      # Bae
 LIpois(4)
                      # Bae
# poisson.test(4)
 LIpois(4, k=32)
                      # 0.7454614 11.7893612
  LIpois(2.1, n=60)
                      # 1.750222
                                   2.493533
```

LIvar

Likelihood Interval of variance and sd assuming Normal Distribution with sample mean and sample size

## Description

Likelihood interval of sd and variance assuming normal distribution. This is estimated likelihood interval, not profile likelihood interval.

#### Usage

```
LIvar(s1, n1, k, conf.level=0.95)
```

#### **Arguments**

s1 standard deviation of the sample

n1 sample size

k 1/k likelihood interval will be calculated

conf.level approximately corresponding confidence level. If k is specified, this is ignored.

#### Details

It calculates likelihood interval of sd and variance assuming normal distribution in one group. The likelihood interval is asymmetric and there is no standard error in the output.

#### Value

PE maximum likelihood estimate on the population variance

LL lower limit of likelihood interval

UL upper limit of likelihood interval

#### Author(s)

Kyun-Seop Bae k@acr.kr

```
x = c(-5.3, -4.5, -1.0, -0.7, 3.7, 3.9, 4.2, 5.5, 6.8, 7.4, 9.3)

LInormVar(x)

LIvar(sd(x), length(x))
```

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LIvRatio	Likelihood Interval of the ratio of two variances from two groups

## Description

Likelihood interval of the ratio of two variances from two groups assuming normal distribution. Likelihood interval usually gives a narrower interval when the likelihood function is asymmetric.

#### Usage

```
LIvRatio(x, y, k, conf.level=0.95)
```

#### **Arguments**

X	observations from the first group, the test group, used for the numerator
У	observations from the second group, the control group, used for the denominator
k	1/k likelihood interval will be provided
conf.level	approximate confidence level

#### **Details**

It calculates likelihood interval of the ratio of two variances from two groups. Likelihood interval usually gives a narrower interval when the likelihood function is asymmetric.

## Value

PE	maximum likelihood estimate on the ratio
LL	lower limit of likelihood interval
UL	upper limit of likelihood interval
logk	log(k) value used for LI
maxLL	maximum log likelihood
conf.level	approximate confidence level

#### Author(s)

```
Kyun-Seop Bae k@acr.kr
```

```
LIvRatio(mtcars$drat, mtcars$wt)
var.test(mtcars$drat, mtcars$wt)
LIvRatio(mtcars$qsec, mtcars$wt)
var.test(mtcars$qsec, mtcars$wt)
LIvtest(sd(mtcars$qsec), nrow(mtcars), sd(mtcars$wt), nrow(mtcars))
```

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LIvtest	Likelihood Interval of variance and sd assuming Normal Distribution using means and SDs

## Description

Likelihood interval of variance and sd assuming normal distribution. This is estimated likelihood interval, not profile likelihood interval.

## Usage

```
LIvtest(s1, n1, s2, n2, k, conf.level=0.95)
```

## Arguments

s1	sample standard deviation of the first group
n1	sample size of the first group
s2	sample standard deviation of the second group
n2	sample size of the second group
k	1/k likelihood interval will be calculated
conf.level	approximate confidence level. If k is specified, this is ignored.

## **Details**

It calculates likelihood interval of variance and sd using sufficient statistics. There is no standard error in the output.

## Value

PE	maximum likelihood estimate on the ratio
LL	lower limit of likelihood interval
UL	upper limit of likelihood interval
logk	log(k) value used for LI
maxLL	maximum log likelihood
conf.level	approximate confidence level

## Author(s)

```
Kyun-Seop Bae k@acr.kr
```

```
LIvtest(10.5, 3529, 8.9, 5190)
LIvtest(3, 10, 2, 10)
LIvtest(3, 10, 2, 10, k=15)
```

10 LRT

LRT	Likelihood Ratio Test	

#### **Description**

Likelihood ratio test with given fitting results, sample size, number of parameters, log-likelihoods, and alpha

## Usage

```
LRT(n, pFull, pReduced, logLikFull, logLikReduced, alpha=0.05, Wilks=FALSE)
```

#### **Arguments**

n number of observations

pFull number of parameters of full model
pReduced number of parameters of reduced model

logLikFull log likelihood of full model logLikReduced log likelihood of reduced model

alpha alpha value for type I error, significance level

Wilks if TRUE, Wilks theorem (chi-square distribution) will be used, otherwise F dis-

tribution will be used.

#### Details

It performs likelihood ratio test with given fitting results. The default test is using F distribution. For small n (i.e. less than 100), you need to use F distribution. If the residuals are normally distributed, the delta -2 log likelihood (the difference between -2LL, the objective function value of each model) follows exactly an F-distribution, independent of sample size. When the distribution of the residuals is not normal (no matter what the distribution of the residuals is), it approaches a chi-square distribution as sample size increases (Wilks' theorem). The extreme distribution of the F-distribution (when the degrees of freedom in the denominator go to infinity) is chi-square distribution. The p-value from the F-distribution is slightly larger than the p-value from the chi-square distribution, meaning the F-distribution is more conservative. The difference decreases as sample size increases.

#### Value

ations
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paraFull number of parameters of full model
paraReduced number of parameters of reduced model

deltaPara difference of parameter counts

cutoff cutoff, threshold, critical value of log-likelihood for the test

deltaLogLik difference of log likelihood, if negative 0 is used.

Chisq or Fval statistics according to the used distribution Chi-square of F pval p-value of null hypothesis. i.e. the reduced model is better.

Verdict the model preferred.

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#### Author(s)

Kyun-Seop Bae k@acr.kr

#### References

 Ruppert D, Cressie N, Carroll RJ. A Transformation/Weighting Model For Estimating Michaelis-Menten Parameters. School of Operations Research and Industrial Engineering, College of Engineering, Cornell University. Technical Report No. 796. May 1988.

- 2. Scheffé H. The Analysis of Variance. Wiley. 1959.
- 3. Wilks SS. The Large-Sample Distribution of the Likelihood Ratio for Testing Composite Hypotheses. *Annals Math. Statist.* 1938;9:60-62

#### **Examples**

```
LRT(20, 4, 2, -58.085, -60.087)
LRT(20, 4, 2, -58.085, -60.087, Wilks=TRUE)
LRT(20, 4, 2, -57.315, -66.159)
LRT(20, 4, 2, -57.315, -66.159, Wilks=TRUE)

r1 = lm(mpg ~ disp + drat + wt, mtcars)
r2 = lm(mpg ~ disp + drat, mtcars)
anova(r2, r1)
LRT(nrow(mtcars), r1$rank, r2$rank, logLik(r1), logLik(r2))
```

**OneTwo** 

Likelihood Ratio Test for One group vs Two group gaussian mixture model

## Description

With a given vector, it performs likelihood ratio test which model - one or two group - is better.

## Usage

```
OneTwo(x, alpha=0.05)
```

#### **Arguments**

x a vector of numbers

alpha alpha value for type I error, significance level

## **Details**

It performs likelihood ratio test using both F distribution and Chi-square distribution (by Wilks' theorem).

#### Value

Estimate n, Mean, SD for each group assumption and prior probability of each group in

two group model

Delta delta number of parameters and log-likelihoods

Statistic Statistics from both the F distribution and Chi-square distribution. Cutoff is in

terms of log-likelihood not the statistic.

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## Author(s)

Kyun-Seop Bae k@acr.kr

#### **Examples**

```
OneTwo(c(7, 5, 17, 13, 16, 5, 7, 3, 8, 10, 8, 14, 14, 11, 14, 17, 2, 12, 15, 19))
OneTwo(c(5, 3, 0, 6, 5, 2, 6, 6, 4, 4, 15, 13, 18, 18, 19, 14, 19, 13, 19, 18))
```

ORLI

Odds Ratio and its Likelihood Interval between two groups without strata

## Description

Odds ratio and its likelihood interval between two groups without stratification

## Usage

```
ORLI(y1, n1, y2, n2, conf.level=0.95, k, eps=1e-8)
```

## Arguments

y1	positive event count of test (the first) group
n1	total count of the test (the first) group. Maximum allowable value is 1e8.
y2	positive event count of control (the second) group
n2	total count of control (the second) group. Maximum allowable value is 1e8.
conf.level	approximate confidence level to calculate k when k is missing.
k	1/k likelihood interval will be provided
eps	absolute value less than eps is regarded as negligible

## **Details**

It calculates risk (proportion) difference and its likelihood interval between the two groups. The likelihood interval is asymmetric, and there is no standard error in the output. This does not support stratification.

## Value

There is no standard error.

odd1 odd from the first group, y1/(n1 - y1) odd2 odd from the second group, y2/(n2 - y2)

OR odds ratio, odd1/odd2

lower likelihood limit of OR upper upper likelihood limit of OR

#### Author(s)

Kyun-Seop Bae k@acr.kr

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#### **Examples**

```
ORLI(7, 10, 3, 10)
ORLI(3, 10, 7, 10)
```

RDLI

Risk (Proportion) Difference and its Likelihood Interval between two groups without strata

## Description

Risk difference and its likelihood interval between two groups without stratification

## Usage

```
RDLI(y1, n1, y2, n2, conf.level=0.95, k, eps=1e-8)
```

#### **Arguments**

y1	positive event count of test (the first) group
n1	total count of the test (the first) group. Maximum allowable value is 1e8.
y2	positive event count of control (the second) group
n2	total count of control (the second) group. Maximum allowable value is 1e8.
conf.level	approximate confidence level to calculate k when k is missing.
k	1/k likelihood interval will be provided
eps	absolute value less than eps is regarded as negligible

#### **Details**

It calculates risk (proportion) difference and its likelihood interval between the two groups. The likelihood interval is asymmetric, and there is no standard error in the output. This does not support stratification.

#### Value

There is no standard error.

```
p1 proportion from the first group, y1/n1
p2 proportion from the second group, y2/n2
RD risk difference, p1 - p2
lower lower likelihood limit of RD
upper upper likelihood limit of RD
```

## Author(s)

Kyun-Seop Bae k@acr.kr

```
RDLI(7, 10, 3, 10)
RDLI(3, 10, 7, 10)
```

14 RRLI

RRLI	Relative Risk and its Likelihood Interval between two groups without strata

## Description

Relative risk and its likelihood interval between two groups without stratification

## Usage

```
RRLI(y1, n1, y2, n2, conf.level=0.95, k, eps=1e-8)
```

## Arguments

y1	positive event count of test (the first) group
n1	total count of the test (the first) group. Maximum allowable value is 1e8.
y2	positive event count of control (the second) group
n2	total count of control (the second) group. Maximum allowable value is 1e8.
conf.level	approximate confidence level to calculate k when k is missing.
k	1/k likelihood interval will be provided
eps	absolute value less than eps is regarded as negligible

#### **Details**

It calculates relative risk and its likelihood interval between the two groups. The likelihood interval is asymmetric, and there is no standard error in the output. This does not support stratification.

#### Value

There is no standard error.

p1	proportion from the first group, y1/n1
p2	proportion from the second group, y2/n2
	1

RR relative risk, p1/p2

lower likelihood limit of RR upper upper likelihood limit of RR

## Author(s)

Kyun-Seop Bae k@acr.kr

```
RRLI(7, 10, 3, 10)
RRLI(3, 10, 7, 10)
```

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