

Package ‘logitnorm’

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Title Functions for the Logitnormal Distribution

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Description Density, distribution, quantile and random generation function for the logitnormal distribution. Estimation of the mode and the first two moments. Estimation of distribution parameters.

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logitnorm-package *Utilities for the logitnormal distribution in R*

Description

Utilities for the logitnormal distribution in R

- Density, distribution, quantile and random generation function.
- Estimation of the mode and the first two moments.
- Estimation of distribution parameters from observations.

Details

The package provides the main distribution functions:

- density `dlogitnorm`,
- distribution `plogitnorm`,
- quantile `qlogitnorm`, and
- random generation function `rlogitnorm`.

Transformation functions

- $(0,1) \rightarrow (-\infty, \infty)$: `logit`
- $(-\infty, \infty) \rightarrow (0,1)$: `invlogit`

Moments and mode

- Expected value and variance: `momentsLogitnorm`
- Mode: `modeLogitnorm`

Estimating parameters

- from mode and upper quantile: `twCoefLogitnormMLE`
- from mode and constraint to be unimodal and maximally flat: `twCoefLogitnormMLEflat`
- from median and upper quantile: `twCoefLogitnorm`
- from expected value, i.e. mean and upper quantile: `twCoefLogitnormE`
- from a confidence interval which is symmetric at normal scale: `twCoefLogitnormCi`
- from prescribed quantiles: `twCoefLogitnormN`

Have a look at the package vignettes.

Author(s)

Thomas Wutzler

References

Frederic, P. & Lad, F. (2008) Two Moments of the Logitnormal Distribution. Communications in Statistics-Simulation and Computation, 37, 1263-1269

dlogitnorm	<i>dlogitnorm</i>
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Description

Density function of logitnormal distribution

Usage

```
dlogitnorm(x, mu = 0, sigma = 1, log = FALSE,  
...)
```

Arguments

x	vector of quantiles
mu	scale distribution parameter
sigma	location distribution parameter
log	if TRUE, the log-density is returned
...	further arguments passed to dnorm : mean, and sd for mu and sigma respectively.

Details

Logitnorm distribution • density function: [dlogitnorm](#)

- distribution function: [plogitnorm](#)
- quantile function: [qlogitnorm](#)
- random generation function: [rlogitnorm](#)

The function is only defined in interval (0,1), but the density returns 0 outside the support region.

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

invlogit	<i>invlogit</i>
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Description

Transforming (-Inf,Inf) to original scale (0,1)

Usage

```
invlogit(q, ...)
```

Arguments

q	quantile
...	further arguments to plogis

Details

function $f(z) = \frac{e^z}{e^z+1} = \frac{1}{1+e^{-z}}$

Author(s)

Thomas Wutzler

See Also

[logit](#)

[logitnorm](#)

logit	<i>logit</i>
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Description

Transforming (0,1) to normal scale (-Inf Inf)

Usage

```
logit(p, ...)
```

Arguments

p	percentile
...	further arguments to qlogis

Details

$$\text{function } \text{logit}(p) = \log\left(\frac{p}{1-p}\right) = \log(p) - \log(1-p)$$

Author(s)

Thomas Wutzler

See Also

[invlogit](#)

[logitnorm](#)

modeLogitnorm

modeLogitnorm

Description

Mode of the logitnormal distribution by numerical optimization

Usage

```
modeLogitnorm(mu, sigma, tol = invlogit(mu)/1000)
```

Arguments

mu	parameter mu
sigma	parameter sigma
tol	precisions of the estimate

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

momentsLogitnorm	<i>momentsLogitnorm</i>
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Description

First two moments of the logitnormal distribution by numerical integration

Usage

```
momentsLogitnorm(mu, sigma, abs.tol = 0,  
  ...)
```

Arguments

mu	parameter mu
sigma	parameter sigma
abs.tol	changing default to integrate
...	further parameters to the integrate function

Value

named numeric vector with components

- mean: expected value, i.e. first moment
- var: variance, i.e. second moment

Author(s)

Thomas Wutzler

Examples

```
(res <- momentsLogitnorm(4,1))  
(res <- momentsLogitnorm(5,0.1))
```

plogitnorm

plogitnorm

Description

Distribution function for logitnormal distribution

Usage

```
plogitnorm(q, mu = 0, sigma = 1, ...)
```

Arguments

q	vector of quantiles
mu	location distribution parameter
sigma	scale distribution parameter
...	further arguments to pnorm

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

qlogitnorm

qlogitnorm

Description

Quantiles of logitnormal distribution.

Usage

```
qlogitnorm(p, mu = 0, sigma = 1, ...)
```

Arguments

p	vector of probabilities
mu	location distribution parameter
sigma	scale distribution parameter
...	further arguments to plogis

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

rlogitnorm

rlogitnorm

Description

Random number generation for logitnormal distribution

Usage

```
rlogitnorm(n, mu = 0, sigma = 1, ...)
```

Arguments

n	number of observations
mu	distribution parameter
sigma	distribution parameter
...	arguments to rnorm

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

twCoefLogitnorm	<i>twCoefLogitnorm</i>
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Description

Estimating coefficients of logitnormal distribution from median and upper quantile

Usage

```
twCoefLogitnorm(median, quant, perc = 0.975)
```

Arguments

median	numeric vector: the median of the density function
quant	numeric vector: the upper quantile value
perc	numeric vector: the probability for which the quantile was specified

Value

numeric matrix with columns c("mu", "sigma") rows correspond to rows in median, quant, and perc

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

Examples

```
# estimate the parameters, with median at 0.7 and upper quantile at 0.9
med = 0.7; upper = 0.9
med = 0.2; upper = 0.4
(theta <- twCoefLogitnorm(med,upper))

x <- seq(0,1,length.out = 41)[-c(1,41)] # plotting grid
px <- plogitnorm(x,mu = theta[1],sigma = theta[2]) #percentiles function
plot(px~x); abline(v = c(med,upper),col = "gray"); abline(h = c(0.5,0.975),col = "gray")

dx <- dlogitnorm(x,mu = theta[1],sigma = theta[2]) #density function
plot(dx~x); abline(v = c(med,upper),col = "gray")

# vectorized
(theta <- twCoefLogitnorm(seq(0.4,0.8,by = 0.1),0.9))

.tmp.f <- function(){
  # xr = rlogitnorm(1e5, mu = theta["mu"], sigma = theta["sigma"])
```

```

# median(xr)
invlogit(theta["mu"])
qlogitnorm(0.975, mu = theta["mu"], sigma = theta["sigma"])
}

```

twCoefLogitnormCi	<i>twCoefLogitnormCi</i>
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Description

Calculates mu and sigma of the logitnormal distribution from lower and upper quantile, i.e. confidence interval.

Usage

```
twCoefLogitnormCi(lower, upper, perc = 0.975,
  sigmaFac = qnorm(perc), isTransScale = FALSE)
```

Arguments

lower	value at the lower quantile, i.e. practical minimum
upper	value at the upper quantile, i.e. practical maximum
perc	numeric vector: the probability for which the quantile was specified
sigmaFac	sigmaFac = 2 is 95% sigmaFac = 2.6 is 99% interval
isTransScale	if true lower and upper are already on logit scale

Value

named numeric vector: mu and sigma parameter of the logitnormal distribution.

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

Examples

```

mu = 2
sd = c(1,0.8)
p = 0.99
lower <- l <- qlogitnorm(1 - p, mu, sd ) # p-confidence interval
upper <- u <- qlogitnorm(p, mu, sd ) # p-confidence interval
cf <- twCoefLogitnormCi(lower,upper, perc = p)
all.equal( cf[, "mu"] , c(mu,mu) )
all.equal( cf[, "sigma"] , sd )

```

twCoefLogitnormE	<i>twCoefLogitnormE</i>
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Description

Estimating coefficients of logitnormal distribution from expected value, i.e. mean, and upper quantile.

Usage

```
twCoefLogitnormE(mean, quant, perc = c(0.975),
  method = "BFGS", theta0 = c(mu = 0, sigma = 1),
  returnDetails = FALSE, ...)
```

Arguments

mean	the expected value of the density function
quant	the quantile values
perc	the probabilities for which the quantiles were specified
method	method of optimization (see optim)
theta0	starting parameters
returnDetails	if TRUE, the full output of optim is returned with attribute resOptim
...	further arguments to optim

Value

named numeric matrix with estimated parameters of the logitnormal distribution. colnames: c("mu", "sigma")

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

Examples

```
# estimate the parameters
(thetaE <- twCoefLogitnormE(0.7,0.9))

x <- seq(0,1,length.out = 41)[-c(1,41)] # plotting grid
px <- plogitnorm(x,mu = thetaE[1],sigma = thetaE[2]) #percentiles function
plot(px~x); abline(v = c(0.7,0.9),col = "gray"); abline(h = c(0.5,0.975),col = "gray")
dx <- dlogitnorm(x,mu = thetaE[1],sigma = thetaE[2]) #density function
plot(dx~x); abline(v = c(0.7,0.9),col = "gray")
```

```
z <- rlogitnorm(1e5, mu = thetaE[1],sigma = thetaE[2])
mean(z) # about 0.7

# vectorized
(theta <- twCoefLogitnormE(mean = seq(0.4,0.8,by = 0.1),quant = 0.9))
```

```
twCoefLogitnormMLE      twCoefLogitnormMLE
```

Description

Estimating coefficients of logitnormal distribution from mode and upper quantile

Usage

```
twCoefLogitnormMLE(mle, quant, perc = 0.999)
```

Arguments

mle	numeric vector: the mode of the density function
quant	numeric vector: the upper quantile value
perc	numeric vector: the probability for which the quantile was specified

Value

numeric matrix with columns c("mu", "sigma") rows correspond to rows in mle, quant, and perc

Author(s)

Thomas Wutzler

See Also

[logitnorm](#)

Examples

```
# estimate the parameters, with mode 0.7 and upper quantile 0.9
mode = 0.7; upper = 0.9
(theta <- twCoefLogitnormMLE(mode,upper))
x <- seq(0,1,length.out = 41)[-c(1,41)] # plotting grid
px <- plogitnorm(x,mu = theta[1],sigma = theta[2]) #percentiles function
plot(px~x); abline(v = c(mode,upper),col = "gray"); abline(h = c(0.999),col = "gray")
dx <- dlogitnorm(x,mu = theta[1],sigma = theta[2]) #density function
plot(dx~x); abline(v = c(mode,upper),col = "gray")
# vectorized
(theta <- twCoefLogitnormMLE(mle = seq(0.4,0.8,by = 0.1),quant = upper))
# flat
(theta <- twCoefLogitnormMLEFlat(mode))
```

```
twCoefLogitnormMLEFlat
      twCoefLogitnormMLEFlat
```

Description

Estimating coefficients of a maximally flat unimodal logitnormal distribution given the mode

Usage

```
twCoefLogitnormMLEFlat(mle)
```

Arguments

mle numeric vector: the mode of the density function

Author(s)

Thomas Wutzler

```
twCoefLogitnormN        twCoefLogitnormN
```

Description

Estimating coefficients from a vector of quantiles and percentiles (non-vectorized).

Usage

```
twCoefLogitnormN(quant, perc = c(0.5, 0.975),
  method = "BFGS", theta0 = c(mu = 0, sigma = 1),
  returnDetails = FALSE, ...)
```

Arguments

quant the quantile values
 perc the probabilities for which the quantiles were specified
 method method of optimization (see [optim](#))
 theta0 starting parameters
 returnDetails if TRUE, the full output of [optim](#) is returned instead of only entry par
 ... further parameters passed to [optim](#), e.g. `control = list(maxit = 1000)`

Value

named numeric vector with estimated parameters of the logitnormal distribution. names: `c("mu", "sigma")`

Author(s)

Thomas Wutzler

See Also[logitnorm](#)**Examples**

```
# experiment of re-estimation the parameters from generated observations
thetaTrue <- c(mu = 0.8, sigma = 0.7)
obsTrue <- rlogitnorm(thetaTrue["mu"],thetaTrue["sigma"], n = 500)
obs <- obsTrue + rnorm(100, sd = 0.05) # some observation uncertainty
plot(density(obsTrue),col = "blue"); lines(density(obs))

# re-estimate parameters based on the quantiles of the observations
(theta <- twCoefLogitnorm( median(obs), quantile(obs,probs = 0.9), perc = 0.9))

# add line of estimated distribution
x <- seq(0,1,length.out = 41)[-c(1,41)] # plotting grid
dx <- dlogitnorm(x,mu = theta[1],sigma = theta[2])
lines( dx ~ x, col = "orange")
```

twSigmaLogitnorm	<i>twSigmaLogitnorm</i>
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Description

Estimating coefficients of logitnormal distribution from mode and given mu

Usage

```
twSigmaLogitnorm(mle, mu = 0)
```

Arguments

mle	numeric vector: the mode of the density function
mu	for mu = 0 the distribution will be the flattest case (maybe bimodal)

Details

For a mostly flat unimodal distribution use [twCoefLogitnormMLE\(mle,0\)](#)

Value

numeric matrix with columns c("mu", "sigma") rows correspond to rows in mle and mu

Author(s)

Thomas Wutzler

See Also[logitnorm](#)**Examples**

```
mle <- 0.8
(theta <- twSigmaLogitnorm(mle))
#
x <- seq(0,1,length.out = 41)[-c(1,41)] # plotting grid
px <- plogitnorm(x,mu = theta[1],sigma = theta[2]) #percentiles function
plot(px~x); abline(v = c(mle),col = "gray")
dx <- dlogitnorm(x,mu = theta[1],sigma = theta[2]) #density function
plot(dx~x); abline(v = c(mle),col = "gray")
# vectorized
(theta <- twSigmaLogitnorm(mle = seq(0.401,0.8,by = 0.1)))
```

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