

Package ‘boodist’

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Title Some Distributions from the 'Boost' Library and More

Version 1.0.0

Description Make some distributions from the 'C++' library 'Boost' available in 'R'. In addition, the normal-inverse Gaussian distribution and the generalized inverse Gaussian distribution are provided. The distributions are represented by 'R6' classes. The method to sample from the generalized inverse Gaussian distribution is the one given in ``Random variate generation for the generalized inverse Gaussian distribution" Luc Devroye (2012) <[doi:10.1007/s11222-012-9367-z](https://doi.org/10.1007/s11222-012-9367-z)>.

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URL <https://github.com/stla/boodist>

BugReports <https://github.com/stla/boodist/issues>

Imports R6, Rcpp, RcppNumerical, stats

LinkingTo BH, Rcpp, RcppEigen, RcppNumerical

Suggests plotly

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Author Stéphane Laurent [aut, cre]

Maintainer Stéphane Laurent <laurent_step@outlook.fr>

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findChi2df	<i>Find degrees of freedom</i>
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Description

Find the degrees of freedom parameter of a non-central Chi-squared distribution given a quantile, its corresponding probability, and the non-centrality parameter.

Usage

```
findChi2df(ncp, q, p)
```

Arguments

ncp	non-centrality parameter, a non-negative number
q	a quantile
p	probability corresponding to the quantile q

Value

The degrees of freedom parameter of the non-central Chi-squared distribution with non-centrality parameter ncp and with cumulative probability p at the quantile q.

Examples

```
library(boodist)
nu <- findChi2df(ncp = 10, q = 3, p = 0.1)
pchisq(3, df = nu, ncp = 10) # should be 0.1
```

findChi2ncp	<i>Find non-centrality parameter</i>
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Description

Find the non-centrality parameter of a Chi-squared distribution given a quantile, its corresponding probability, and the degrees of freedom.

Usage

```
findChi2ncp(df, q, p)
```

Arguments

df	degrees of freedom, a positive number
q	a quantile
p	probability corresponding to the quantile q

Value

The non-centrality parameter of the Chi-squared distribution with degrees of freedom parameter df and with cumulative probability p at the quantile q.

Examples

```
library(boodist)
ncp <- findChi2ncp(df = 1, q = 3, p = 0.1)
pchisq(3, df = 1, ncp = ncp) # should be 0.1
```

GeneralizedInverseGaussian	<i>Generalized inverse Gaussian distribution</i>
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Description

A R6 class to represent a generalized inverse Gaussian distribution.

Details

See [Wikipedia](#).

Active bindings

theta Get or set the value of theta.
eta Get or set the value of eta.
lambda Get or set the value of lambda.

Methods

Public methods:

- `GeneralizedInverseGaussian$new()`
- `GeneralizedInverseGaussian$d()`
- `GeneralizedInverseGaussian$p()`
- `GeneralizedInverseGaussian$q()`
- `GeneralizedInverseGaussian$r()`
- `GeneralizedInverseGaussian$mean()`
- `GeneralizedInverseGaussian$mode()`
- `GeneralizedInverseGaussian$sd()`
- `GeneralizedInverseGaussian$variance()`
- `GeneralizedInverseGaussian$clone()`

Method `new()`: New generalized inverse Gaussian distribution.

Usage:

```
GeneralizedInverseGaussian$new(theta, eta, lambda)
```

Arguments:

theta concentration parameter, >0

eta scale parameter, >0

lambda parameter (denoted by p on Wikipedia)

Returns: A `GeneralizedInverseGaussian` object.

Method `d()`: Density function of the generalized inverse Gaussian distribution.

Usage:

```
GeneralizedInverseGaussian$d(x, log = FALSE)
```

Arguments:

x vector of positive numbers

log Boolean, whether to return the log-density

Returns: The density or the log-density evaluated at x.

Method `p()`: Cumulative distribution function of the generalized inverse Gaussian distribution.

Usage:

```
GeneralizedInverseGaussian$p(q)
```

Arguments:

q numeric vector of quantiles (≥ 0)

Returns: The cumulative probabilities corresponding to q, with two attributes (see the **Note**).

Method `q()`: Quantile function of the generalized inverse Gaussian distribution.

Usage:

```
GeneralizedInverseGaussian$q(p, bounds = NULL)
```

Arguments:

`p` numeric vector of probabilities
`bounds` bounds enclosing the quantiles to be found (see the **Note**), or NULL for automatic bounds
Returns: The quantiles corresponding to `p`.

Method `r()`: Sampling from the generalized inverse Gaussian distribution.

Usage:
`GeneralizedInverseGaussian$r(n)`

Arguments:
`n` number of simulations

Returns: A numeric vector of length `n`.

Method `mean()`: Mean of the generalized inverse Gaussian distribution.

Usage:
`GeneralizedInverseGaussian$mean()`

Returns: The mean of the generalized inverse Gaussian distribution.

Method `mode()`: Mode of the generalized inverse Gaussian distribution.

Usage:
`GeneralizedInverseGaussian$mode()`

Returns: The mode of the generalized inverse Gaussian distribution.

Method `sd()`: Standard deviation of the generalized inverse Gaussian distribution.

Usage:
`GeneralizedInverseGaussian$sd()`

Returns: The standard deviation of the generalized inverse Gaussian distribution.

Method `variance()`: Variance of the generalized inverse Gaussian distribution.

Usage:
`GeneralizedInverseGaussian$variance()`

Returns: The variance of the generalized inverse Gaussian distribution.

Method `clone()`: The objects of this class are cloneable with this method.

Usage:
`GeneralizedInverseGaussian$clone(deep = FALSE)`

Arguments:
`deep` Whether to make a deep clone.

Note

The cumulative distribution function is evaluated by integrating the density function (in C++). Its returned value has two attributes: a numeric vector "error_estimate" and an integer vector "error_code". The error code is 0 if no problem is detected. If an error code is not 0, a warning is thrown. The quantile function is evaluated by root-finding and then the user must provide some bounds enclosing the values of the quantiles or choose the automatic bounds. A maximum number of iterations is fixed in the root-finding algorithm. If it is reached, a warning is thrown.

Examples

```

if(require("plotly")) {
  library(boodist)

  x_ <- seq(0, 3, length.out = 100L)
  lambda_ <- seq(-1, 1, length.out = 100L)
  dsty <- vapply(lambda_, function(lambda) {
    GeneralizedInverseGaussian$new(theta = 1, eta = 1, lambda)$d(x_)
  }, numeric(length(x_)))
  #
  txt <- matrix(NA_character_, nrow = length(x_), ncol = length(lambda_))
  for(i in 1L:nrow(txt)) {
    for(j in 1L:ncol(txt)) {
      txt[i, j] <- paste0(
        "x: ", formatC(x_[i]),
        "<br> lambda: ", formatC(lambda_[j]),
        "<br> density: ", formatC(dsty[i, j])
      )
    }
  }
  #
  plot_ly(
    x = ~lambda_, y = ~x_, z = ~dsty, type = "surface",
    text = txt, hoverinfo = "text", showscale = FALSE
  ) %>% layout(
    title = "Generalized inverse Gaussian distribution",
    margin = list(t = 40, r = 5, b = 5, l = 5),
    scene = list(
      xaxis = list(
        title = "lambda"
      ),
      yaxis = list(
        title = "x"
      ),
      zaxis = list(
        title = "density"
      )
    )
  )
}

```

Gumbel

Gumbel distribution

Description

A R6 class to represent a Gumbel distribution.

Details

See [Wikipedia](#).

Active bindings

- a Get or set the value of a.
- b Get or set the value of b.

Methods**Public methods:**

- `Gumbel$new()`
- `Gumbel$d()`
- `Gumbel$p()`
- `Gumbel$q()`
- `Gumbel$r()`
- `Gumbel$mean()`
- `Gumbel$median()`
- `Gumbel$mode()`
- `Gumbel$sd()`
- `Gumbel$variance()`
- `Gumbel$skewness()`
- `Gumbel$kurtosis()`
- `Gumbel$kurtosisExcess()`
- `Gumbel$clone()`

Method `new()`: New Gumbel distribution.

Usage:

```
Gumbel$new(a, b)
```

Arguments:

a location parameter

b scale parameter, >0

Returns: A Gumbel object.

Method `d()`: Density function of the Gumbel distribution.

Usage:

```
Gumbel$d(x, log = FALSE)
```

Arguments:

x numeric vector

log Boolean, whether to return the logarithm of the density

Returns: The density or the log-density evaluated at x.

Method `p()`: Cumulative distribution function of the Gumbel distribution.

Usage:

```
Gumbel$p(q, lower = TRUE)
```

Arguments:

q numeric vector of quantiles
lower Boolean, whether to deal with the lower tail
Returns: The cumulative probabilities corresponding to q.

Method q(): Quantile function of the Gumbel distribution.

Usage:
Gumbel\$q(p, lower = TRUE)
Arguments:
p numeric vector of probabilities
lower Boolean, whether to deal with the lower tail
Returns: The quantiles corresponding to p.

Method r(): Sampling from the Gumbel distribution.

Usage:
Gumbel\$r(n)
Arguments:
n number of simulations
Returns: A numeric vector of length n.

Method mean(): Mean of the Gumbel distribution.

Usage:
Gumbel\$mean()
Returns: The mean of the Gumbel distribution.

Method median(): Median of the Gumbel distribution.

Usage:
Gumbel\$median()
Returns: The median of the Gumbel distribution.

Method mode(): Mode of the Gumbel distribution.

Usage:
Gumbel\$mode()
Returns: The mode of the Gumbel distribution.

Method sd(): Standard deviation of the Gumbel distribution.

Usage:
Gumbel\$sd()
Returns: The standard deviation of the Gumbel distribution.

Method variance(): Variance of the Gumbel distribution.

Usage:
Gumbel\$variance()

Returns: The variance of the Gumbel distribution.

Method skewness(): Skewness of the Gumbel distribution.

Usage:

```
Gumbel$skewness()
```

Returns: The skewness of the Gumbel distribution.

Method kurtosis(): Kurtosis of the Gumbel distribution.

Usage:

```
Gumbel$kurtosis()
```

Returns: The kurtosis of the Gumbel distribution.

Method kurtosisExcess(): Kurtosis excess of the Gumbel distribution.

Usage:

```
Gumbel$kurtosisExcess()
```

Returns: The kurtosis excess of the Gumbel distribution.

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
Gumbel$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

Hyperexponential *Hyperexponential distribution*

Description

A R6 class to represent a hyperexponential distribution.

Details

See [Wikipedia](#).

Active bindings

probs Get or set the value of probs.

rates Get or set the value of rates.

Methods**Public methods:**

- `Hyperexponential$new()`
- `Hyperexponential$d()`
- `Hyperexponential$p()`
- `Hyperexponential$q()`
- `Hyperexponential$r()`
- `Hyperexponential$mean()`
- `Hyperexponential$mode()`
- `Hyperexponential$sd()`
- `Hyperexponential$variance()`
- `Hyperexponential$skewness()`
- `Hyperexponential$kurtosis()`
- `Hyperexponential$kurtosisExcess()`
- `Hyperexponential$clone()`

Method `new()`: New hyperexponential distribution.

Usage:

```
Hyperexponential$new(probs, rates)
```

Arguments:

`probs` probabilities (weights), a vector of positive numbers

`rates` rate parameters, vector of positive numbers of the same length as the `probs` vector

Returns: A Hyperexponential object.

Method `d()`: Density function of the hyperexponential distribution.

Usage:

```
Hyperexponential$d(x)
```

Arguments:

`x` vector of positive numbers

Returns: The density evaluated at `x`.

Method `p()`: Cumulative distribution function of the hyperexponential distribution.

Usage:

```
Hyperexponential$p(q, lower = TRUE)
```

Arguments:

`q` numeric vector of quantiles

`lower` Boolean, whether to deal with the lower tail

Returns: The cumulative probabilities corresponding to `q`.

Method `q()`: Quantile function of the hyperexponential distribution.

Usage:

Hyperexponential\$q(p, lower = TRUE)

Arguments:

p numeric vector of probabilities

lower Boolean, whether to deal with the lower tail

Returns: The quantiles corresponding to p.

Method r(): Sampling from the hyperexponential distribution.

Usage:

Hyperexponential\$r(n)

Arguments:

n number of simulations

Returns: A numeric vector of length n.

Method mean(): Mean of the hyperexponential distribution.

Usage:

Hyperexponential\$mean()

Returns: The mean of the hyperexponential distribution.

Method mode(): Mode of the hyperexponential distribution.

Usage:

Hyperexponential\$mode()

Returns: The mode of the hyperexponential distribution.

Method sd(): Standard deviation of the hyperexponential distribution.

Usage:

Hyperexponential\$sd()

Returns: The standard deviation of the hyperexponential distribution.

Method variance(): Variance of the hyperexponential distribution.

Usage:

Hyperexponential\$variance()

Returns: The variance of the hyperexponential distribution.

Method skewness(): Skewness of the hyperexponential distribution.

Usage:

Hyperexponential\$skewness()

Returns: The skewness of the hyperexponential distribution.

Method kurtosis(): Kurtosis of the hyperexponential distribution.

Usage:

Hyperexponential\$kurtosis()

Returns: The kurtosis of the hyperexponential distribution.

Method `kurtosisExcess()`: Kurtosis excess of the hyperexponential distribution.

Usage:

```
Hyperexponential$kurtosisExcess()
```

Returns: The kurtosis excess of the hyperexponential distribution.

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
Hyperexponential$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

InverseGamma

Inverse Gamma distribution

Description

A R6 class to represent an inverse Gamma distribution.

Details

See [Wikipedia](#).

Active bindings

`alpha` Get or set the value of alpha.

`beta` Get or set the value of beta.

Methods

Public methods:

- `InverseGamma$new()`
- `InverseGamma$d()`
- `InverseGamma$p()`
- `InverseGamma$q()`
- `InverseGamma$r()`
- `InverseGamma$mean()`
- `InverseGamma$median()`
- `InverseGamma$mode()`
- `InverseGamma$sd()`
- `InverseGamma$variance()`
- `InverseGamma$skewness()`
- `InverseGamma$kurtosis()`
- `InverseGamma$kurtosisExcess()`

- [InverseGamma\\$clone\(\)](#)

Method `new()`: New inverse Gamma distribution.

Usage:

```
InverseGamma$new(alpha, beta)
```

Arguments:

alpha shape parameter, >0

beta scale parameter, >0

Returns: An inverseGamma object.

Method `d()`: Density function of the inverse Gamma distribution.

Usage:

```
InverseGamma$d(x, log = FALSE)
```

Arguments:

x vector of positive numbers

log Boolean, whether to return the logarithm of the density

Returns: The density or the log-density evaluated at x.

Method `p()`: Cumulative distribution function of the inverse Gamma distribution.

Usage:

```
InverseGamma$p(q, lower = TRUE)
```

Arguments:

q numeric vector of quantiles

lower Boolean, whether to deal with the lower tail

Returns: The cumulative probabilities corresponding to q.

Method `q()`: Quantile function of the inverse Gamma distribution.

Usage:

```
InverseGamma$q(p, lower = TRUE)
```

Arguments:

p numeric vector of probabilities

lower Boolean, whether to deal with the lower tail

Returns: The quantiles corresponding to p.

Method `r()`: Sampling from the inverse Gamma distribution.

Usage:

```
InverseGamma$r(n)
```

Arguments:

n number of simulations

Returns: A numeric vector of length n.

Method `mean()`: Mean of the inverse Gamma distribution.

Usage:

```
InverseGamma$mean()
```

Returns: The mean of the inverse Gamma distribution.

Method median(): Median of the inverse Gamma distribution.

Usage:

```
InverseGamma$median()
```

Returns: The median of the inverse Gamma distribution.

Method mode(): Mode of the inverse Gamma distribution.

Usage:

```
InverseGamma$mode()
```

Returns: The mode of the inverse Gamma distribution.

Method sd(): Standard deviation of the inverse Gamma distribution.

Usage:

```
InverseGamma$sd()
```

Returns: The standard deviation of the inverse Gamma distribution.

Method variance(): Variance of the inverse Gamma distribution.

Usage:

```
InverseGamma$variance()
```

Returns: The variance of the inverse Gamma distribution.

Method skewness(): Skewness of the inverse Gamma distribution.

Usage:

```
InverseGamma$skewness()
```

Returns: The skewness of the inverse Gamma distribution.

Method kurtosis(): Kurtosis of the inverse Gamma distribution.

Usage:

```
InverseGamma$kurtosis()
```

Returns: The kurtosis of the inverse Gamma distribution.

Method kurtosisExcess(): Kurtosis excess of the inverse Gamma distribution.

Usage:

```
InverseGamma$kurtosisExcess()
```

Returns: The kurtosis excess of the inverse Gamma distribution.

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
InverseGamma$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

Examples

```

if(require("plotly")) {
  x_ <- seq(0, 2, length.out = 100L)
  alpha_ <- seq(0.5, 2.5, length.out = 100L)
  dsty <- vapply(alpha_, function(alpha) {
    InverseGamma$new(alpha, beta = 1)$d(x_)
  }, numeric(length(x_)))
  #
  txt <- matrix(NA_character_, nrow = length(x_), ncol = length(alpha_))
  for(i in 1L:nrow(txt)) {
    for(j in 1L:ncol(txt)) {
      txt[i, j] <- paste0(
        "x: ", formatC(x_[i]),
        "<br> alpha: ", formatC(alpha_[j]),
        "<br> density: ", formatC(dsty[i, j])
      )
    }
  }
  #
  plot_ly(
    x = ~alpha_, y = ~x_, z = ~dsty, type = "surface",
    text = txt, hoverinfo = "text", showscale = FALSE
  ) %>% layout(
    title = "Inverse Gamma distribution",
    margin = list(t = 40, r = 5, b = 5, l = 5),
    scene = list(
      xaxis = list(
        title = "alpha"
      ),
      yaxis = list(
        title = "x"
      ),
      zaxis = list(
        title = "density"
      )
    )
  )
}

```

InverseGaussian

Inverse Gaussian distribution

Description

A R6 class to represent an inverse Gaussian distribution.

Details

See [Wikipedia](#).

Active bindings

mu Get or set the value of mu.

lambda Get or set the value of lambda.

Methods**Public methods:**

- `InverseGaussian$new()`
- `InverseGaussian$d()`
- `InverseGaussian$p()`
- `InverseGaussian$q()`
- `InverseGaussian$r()`
- `InverseGaussian$mean()`
- `InverseGaussian$median()`
- `InverseGaussian$mode()`
- `InverseGaussian$sd()`
- `InverseGaussian$variance()`
- `InverseGaussian$skewness()`
- `InverseGaussian$kurtosis()`
- `InverseGaussian$kurtosisExcess()`
- `InverseGaussian$clone()`

Method `new()`: New inverse Gaussian distribution.

Usage:

```
InverseGaussian$new(mu, lambda)
```

Arguments:

mu parameter, the mean, >0

lambda shape parameter, >0

Returns: An inverseGaussian object.

Method `d()`: Density function of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$d(x, log = FALSE)
```

Arguments:

x vector of positive numbers

log Boolean, whether to return the logarithm of the density

Returns: The density or the log-density evaluated at x.

Method `p()`: Cumulative distribution function of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$p(q, lower = TRUE)
```

Arguments:

q numeric vector of quantiles
lower Boolean, whether to deal with the lower tail

Returns: The cumulative probabilities corresponding to q.

Method q(): Quantile function of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$q(p, lower = TRUE)
```

Arguments:

p numeric vector of probabilities
lower Boolean, whether to deal with the lower tail

Returns: The quantiles corresponding to p.

Method r(): Sampling from the inverse Gaussian distribution.

Usage:

```
InverseGaussian$r(n)
```

Arguments:

n number of simulations

Returns: A numeric vector of length n.

Method mean(): Mean of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$mean()
```

Returns: The mean of the inverse Gaussian distribution.

Method median(): Median of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$median()
```

Returns: The median of the inverse Gaussian distribution.

Method mode(): Mode of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$mode()
```

Returns: The mode of the inverse Gaussian distribution.

Method sd(): Standard deviation of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$sd()
```

Returns: The standard deviation of the inverse Gaussian distribution.

Method variance(): Variance of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$variance()
```

Returns: The variance of the inverse Gaussian distribution.

Method skewness(): Skewness of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$skewness()
```

Returns: The skewness of the inverse Gaussian distribution.

Method kurtosis(): Kurtosis of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$kurtosis()
```

Returns: The kurtosis of the inverse Gaussian distribution.

Method kurtosisExcess(): Kurtosis excess of the inverse Gaussian distribution.

Usage:

```
InverseGaussian$kurtosisExcess()
```

Returns: The kurtosis excess of the inverse Gaussian distribution.

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
InverseGaussian$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

NormalInverseGaussian *Normal-inverse Gaussian distribution*

Description

A R6 class to represent a normal-inverse Gaussian distribution.

Details

See [Wikipedia](#).

Active bindings

mu Get or set the value of mu.

alpha Get or set the value of alpha.

beta Get or set the value of beta.

delta Get or set the value of delta.

Methods**Public methods:**

- `NormalInverseGaussian$new()`
- `NormalInverseGaussian$d()`
- `NormalInverseGaussian$p()`
- `NormalInverseGaussian$q()`
- `NormalInverseGaussian$r()`
- `NormalInverseGaussian$mean()`
- `NormalInverseGaussian$sd()`
- `NormalInverseGaussian$variance()`
- `NormalInverseGaussian$skewness()`
- `NormalInverseGaussian$kurtosis()`
- `NormalInverseGaussian$kurtosisExcess()`
- `NormalInverseGaussian$clone()`

Method `new()`: New normal-inverse Gaussian distribution.

Usage:

```
NormalInverseGaussian$new(mu, alpha, beta, delta)
```

Arguments:

mu location parameter

alpha tail heaviness parameter, >0

beta asymmetry parameter

delta scale parameter, >0

Returns: A NormalInverseGaussian object.

Method `d()`: Density function of the normal-inverse Gaussian distribution.

Usage:

```
NormalInverseGaussian$d(x, log = FALSE)
```

Arguments:

x numeric vector

log Boolean, whether to return the logarithm of the density

Returns: The density or the log-density evaluated at x.

Method `p()`: Cumulative distribution function of the normal-inverse Gaussian distribution.

Usage:

```
NormalInverseGaussian$p(q)
```

Arguments:

q numeric vector of quantiles

Returns: The cumulative probabilities corresponding to q, with two attributes (see the **Note**).

Method `q()`: Quantile function of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$q(p, bounds = NULL)

Arguments:

p numeric vector of probabilities

bounds bounds enclosing the quantiles to be found (see the **Note**), or NULL for automatic bounds

Returns: The quantiles corresponding to p.

Method r(): Sampling from the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$r(n)

Arguments:

n number of simulations

Returns: A numeric vector of length n.

Method mean(): Mean of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$mean()

Returns: The mean of the normal-inverse Gaussian distribution.

Method sd(): Standard deviation of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$sd()

Returns: The standard deviation of the normal-inverse Gaussian distribution.

Method variance(): Variance of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$variance()

Returns: The variance of the normal-inverse Gaussian distribution.

Method skewness(): Skewness of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$skewness()

Returns: The skewness of the normal-inverse Gaussian distribution.

Method kurtosis(): Kurtosis of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$kurtosis()

Returns: The kurtosis of the normal-inverse Gaussian distribution.

Method kurtosisExcess(): Kurtosis excess of the normal-inverse Gaussian distribution.

Usage:

NormalInverseGaussian\$kurtosisExcess()

Returns: The kurtosis excess of the normal-inverse Gaussian distribution.

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
NormalInverseGaussian$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

Note

The cumulative distribution function is evaluated by integrating the density function (in C++). Its returned value has two attributes: a numeric vector "error_estimate" and an integer vector "error_code". The error code is 0 if no problem is detected. If an error code is not 0, a warning is thrown. The quantile function is evaluated by root-finding and then the user must provide some bounds enclosing the values of the quantiles or choose the automatic bounds. A maximum number of iterations is fixed in the root-finding algorithm. If it is reached, a warning is thrown.

SkewNormal

Skew normal distribution

Description

A R6 class to represent a skew normal distribution.

Details

See [Wikipedia](#).

Active bindings

`xi` Get or set the value of `xi`.

`omega` Get or set the value of `omega`.

`alpha` Get or set the value of `alpha`.

Methods

Public methods:

- `SkewNormal$new()`
- `SkewNormal$d()`
- `SkewNormal$p()`
- `SkewNormal$q()`
- `SkewNormal$r()`
- `SkewNormal$mean()`
- `SkewNormal$mode()`
- `SkewNormal$sd()`

- `SkewNormal$variance()`
- `SkewNormal$skewness()`
- `SkewNormal$skurtosis()`
- `SkewNormal$skurtosisExcess()`
- `SkewNormal$clone()`

Method `new()`: New skew normal distribution.

Usage:

`SkewNormal$new(xi, omega, alpha)`

Arguments:

`xi` location parameter

`omega` scale parameter, >0

`alpha` shape parameter

Returns: A `SkewNormal` object.

Method `d()`: Density function of the skew normal distribution.

Usage:

`SkewNormal$d(x)`

Arguments:

`x` numeric vector

Returns: The density evaluated at `x`.

Method `p()`: Cumulative distribution function of the skew normal distribution.

Usage:

`SkewNormal$p(q, lower = TRUE)`

Arguments:

`q` numeric vector of quantiles

`lower` Boolean, whether to deal with the lower tail

Returns: The cumulative probabilities corresponding to `q`.

Method `q()`: Quantile function of the skew normal distribution.

Usage:

`SkewNormal$q(p, lower = TRUE)`

Arguments:

`p` numeric vector of probabilities

`lower` Boolean, whether to deal with the lower tail

Returns: The quantiles corresponding to `p`.

Method `r()`: Sampling from the skew normal distribution.

Usage:

`SkewNormal$r(n)`

Arguments:

n number of simulations

Returns: A numeric vector of length n.

Method mean(): Mean of the skew normal distribution.

Usage:

SkewNormal\$mean()

Returns: The mean of the skew normal distribution.

Method mode(): Mode of the skew normal distribution.

Usage:

SkewNormal\$mode()

Returns: The mode of the skew normal distribution.

Method sd(): Standard deviation of the skew normal distribution.

Usage:

SkewNormal\$sd()

Returns: The standard deviation of the skew normal distribution.

Method variance(): Variance of the skew normal distribution.

Usage:

SkewNormal\$variance()

Returns: The variance of the skew normal distribution.

Method skewness(): Skewness of the skew normal distribution.

Usage:

SkewNormal\$skewness()

Returns: The skewness of the skew normal distribution.

Method kurtosis(): Kurtosis of the skew normal distribution.

Usage:

SkewNormal\$kurtosis()

Returns: The kurtosis of the skew normal distribution.

Method kurtosisExcess(): Kurtosis excess of the skew normal distribution.

Usage:

SkewNormal\$kurtosisExcess()

Returns: The kurtosis excess of the skew normal distribution.

Method clone(): The objects of this class are cloneable with this method.

Usage:

SkewNormal\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

Student

Non-central Student distribution

Description

A R6 class to represent a non-central Student distribution.

Active bindings

nu Get or set the value of nu.

delta Get or set the value of delta.

Methods

Public methods:

- `Student$new()`
- `Student$d()`
- `Student$p()`
- `Student$q()`
- `Student$r()`
- `Student$mean()`
- `Student$median()`
- `Student$mode()`
- `Student$sd()`
- `Student$variance()`
- `Student$skewness()`
- `Student$kurtosis()`
- `Student$kurtosisExcess()`
- `Student$clone()`

Method `new()`: New Student distribution.

Usage:

`Student$new(nu, delta)`

Arguments:

nu degrees of freedom parameter, >0

delta non-centrality parameter

Returns: A Student object.

Method `d()`: Density function of the Student distribution.

Usage:

`Student$d(x)`

Arguments:

x numeric vector

Returns: The density evaluated at x.

Method p(): Cumulative distribution function of the Student distribution.

Usage:

Student\$p(q, lower = TRUE)

Arguments:

q numeric vector of quantiles

lower Boolean, whether to deal with the lower tail

Returns: The cumulative probabilities corresponding to q.

Method q(): Quantile function of the Student distribution.

Usage:

Student\$q(p, lower = TRUE)

Arguments:

p numeric vector of probabilities

lower Boolean, whether to deal with the lower tail

Returns: The quantiles corresponding to p.

Method r(): Sampling from the Student distribution.

Usage:

Student\$r(n)

Arguments:

n number of simulations

Returns: A numeric vector of length n.

Method mean(): Mean of the Student distribution.

Usage:

Student\$mean()

Returns: The mean of the Student distribution.

Method median(): Median of the Student distribution.

Usage:

Student\$median()

Returns: The median of the Student distribution.

Method mode(): Mode of the Student distribution.

Usage:

Student\$mode()

Returns: The mode of the Student distribution.

Method sd(): Standard deviation of the Student distribution.

Usage:

Student\$sd()

Returns: The standard deviation of the Student distribution.

Method variance(): Variance of the Student distribution.

Usage:

Student\$variance()

Returns: The variance of the Student distribution.

Method skewness(): Skewness of the Student distribution.

Usage:

Student\$skewness()

Returns: The skewness of the Student distribution.

Method kurtosis(): Kurtosis of the Student distribution.

Usage:

Student\$kurtosis()

Returns: The kurtosis of the Student distribution.

Method kurtosisExcess(): Kurtosis excess of the Student distribution.

Usage:

Student\$kurtosisExcess()

Returns: The kurtosis excess of the Student distribution.

Method clone(): The objects of this class are cloneable with this method.

Usage:

Student\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

Note

The non-centrality parameter of the Student distribution in the **stats** package is limited to $\text{abs}(ncp) \leq 37.62$. The present implementation allows a larger range.

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