Package: PoissonBinomial (via r-universe)

October 28, 2026

Type Package

Title Efficient Computation of Ordinary and Generalised Poisson Binomial Distributions

Version 1.2.8

Date 2026-03-06

Maintainer Florian Junge <florian.junge@mailbox.org>

Language en-GB

Description Efficient implementations of multiple exact and approximate methods as described in Hong (2013) <doi:10.1016/j.csda.2012.10.006>, Biscarri, Zhao & Brunner (2018) <doi:10.1016/j.csda.2018.01.007> and Zhang, Hong & Balakrishnan (2018) <doi:10.1080/00949655.2018.1440294> for computing the probability mass, cumulative distribution and quantile functions, as well as generating random numbers for both the ordinary and generalised Poisson binomial distribution.

License GPL-3

Encoding UTF-8

Imports Rcpp (>= 1.0.11)

LinkingTo Rcpp

SystemRequirements fftw3 (>= 3.3)

Suggests knitr, rmarkdown, microbenchmark

VignetteBuilder knitr

URL https://github.com/fj86/PoissonBinomial

BugReports https://github.com/fj86/PoissonBinomial/issues

RoxygenNote 7.3.3

NeedsCompilation yes

Author Florian Junge [aut, cre] (ORCID: <https://orcid.org/0009-0001-6856-6938>)

Config/pak/sysreqs libfftw3-dev

Repository https://cran.r-universe.dev

Date/Publication 2026-03-06 13:50:13 UTC

RemoteUrl https://github.com/cran/PoissonBinomial

RemoteRef HEAD

RemoteSha 47ad2b35ea4347cccd58a9a7774cc036ecaa02c1

Contents

PoissonBinomial-package 2
GenPoissonBinomial-Distribution 3
PoissonBinomial-Distribution 6

Index **10**

`PoissonBinomial-package`

Efficient Exact and Approximate Implementations for Computing Ordinary and Generalised Poisson Binomial Distributions

---

Description

This package implements various algorithms for computing the probability mass function, the cumulative distribution function, quantiles and random numbers of both ordinary and generalised Poisson binomial distributions.

References

Hong, Y. (2013). On computing the distribution function for the Poisson binomial distribution. *Computational Statistics & Data Analysis*, **59**, pp. 41-51. doi:10.1016/j.csda.2012.10.006

Biscarri, W., Zhao, S. D. and Brunner, R. J. (2018) A simple and fast method for computing the Poisson binomial distribution. *Computational Statistics and Data Analysis*, **31**, pp. 216–222. doi:10.1016/j.csda.2018.01.007

Zhang, M., Hong, Y. and Balakrishnan, N. (2018). The generalized Poisson-binomial distribution and the computation of its distribution function. *Journal of Statistical Computational and Simulation*, **88**(8), pp. 1515-1527. doi:10.1080/00949655.2018.1440294

Author(s)

Maintainer: Florian Junge <florian.junge@mailbox.org> (ORCID)

See Also

Useful links:

- https://github.com/fj86/PoissonBinomial
- Report bugs at https://github.com/fj86/PoissonBinomial/issues

Examples

```
# Functions for ordinary Poisson binomial distributions
set.seed(1)
pp <- c(1, 0, runif(10), 1, 0, 1)
qq <- seq(0, 1, 0.01)

dpbinom(NULL, pp)
ppbinom(7:10, pp, method = "DivideFFT")
qpbinom(qq, pp, method = "Convolve")
rpbinom(10, pp, method = "RefinedNormal")

# Functions for generalised Poisson binomial distributions
va <- rep(5, length(pp))
vb <- 1:length(pp)

dgpbinom(NULL, pp, va, vb, method = "Convolve")
pgpbinom(80:100, pp, va, vb, method = "Convolve")
qgpbinom(qq, pp, va, vb, method = "Convolve")
rgpbinom(100, pp, va, vb, method = "Convolve")
```

GenPoissonBinomial-Distribution

The Generalised Poisson Binomial Distribution

Description

Density, distribution function, quantile function and random generation for the generalised Poisson binomial distribution with probability vector `probs`.

Usage

```
dgpbinom(x, probs, val_p, val_q, wts = NULL, method = "DivideFFT", log = FALSE)

pgpbinom(
  x,
  probs,
  val_p,
  val_q,
  wts = NULL,
  method = "DivideFFT",
  lower.tail = TRUE,
  log.p = FALSE
)

qgpbinom(
  p,
  probs,
  val_p,
```

```
  val_q,
  wts = NULL,
  method = "DivideFFT",
  lower.tail = TRUE,
  log.p = FALSE
)

rgpbinom(
  n,
  probs,
  val_p,
  val_q,
  wts = NULL,
  method = "DivideFFT",
  generator = "Sample"
)
```

Arguments

x	Either a vector of observed sums or NULL. If NULL, probabilities of all possible observations are returned.
probs	Vector of probabilities of success of each Bernoulli trial.
val_p	Vector of values that each trial produces with probability in probs.
val_q	Vector of values that each trial produces with probability in 1 - probs.
wts	Vector of non-negative integer weights for the input probabilities.
method	Character string that specifies the method of computation and must be one of "DivideFFT", "Convolve", "Characteristic", "Normal" or "RefinedNormal" (abbreviations are allowed).
log, log.p	Logical value indicating if results are given as logarithms.
lower.tail	Logical value indicating if results are $P[X \leq x]$ (if TRUE; default) or $P[X > x]$ (if FALSE).
p	Vector of probabilities for computation of quantiles.
n	Number of observations. If length(n) > 1, the length is taken to be the number required.
generator	Character string that specifies the random number generator and must either be "Sample" or "Bernoulli" (abbreviations are allowed).

Details

See the references for computational details. The *Divide and Conquer* ("DivideFFT") and *Direct Convolution* ("Convolve") algorithms are derived and described in Biscarri, Zhao & Brunner (2018). They have been modified for use with the generalised Poisson binomial distribution. The *Discrete Fourier Transformation of the Characteristic Function* ("Characteristic") is derived in Zhang, Hong & Balakrishnan (2018), the *Normal Approach* ("Normal") and the *Refined Normal Approach* ("RefinedNormal") are described in Hong (2013). They were slightly adapted for the generalised Poisson binomial distribution.

In some special cases regarding the values of probs, the method parameter is ignored (see Introduction vignette).

Random numbers can be generated in two ways. The "Sample" method uses R's sample function to draw random values according to their probabilities that are calculated by dgpbinom. The "Bernoulli" procedure ignores the method parameter and simulates Bernoulli-distributed random numbers according to the probabilities in probs and sums them up. It is a bit slower than the "Sample" generator, but may yield better results, as it allows to obtain observations that cannot be generated by the "Sample" procedure, because dgpbinom may compute 0-probabilities, due to rounding, if the length of probs is large and/or its values contain a lot of very small values.

Value

dgpbinom gives the density, pgpbinom computes the distribution function, qgpbinom gives the quantile function and rgpbinom generates random deviates.

For rgpbinom, the length of the result is determined by n, and is the lengths of the numerical arguments for the other functions.

References

Hong, Y. (2018). On computing the distribution function for the Poisson binomial distribution. *Computational Statistics & Data Analysis*, **59**, pp. 41-51. doi:10.1016/j.csda.2012.10.006

Biscarri, W., Zhao, S. D. and Brunner, R. J. (2018) A simple and fast method for computing the Poisson binomial distribution. *Computational Statistics and Data Analysis*, **31**, pp. 216–222. doi:10.1016/j.csda.2018.01.007

Zhang, M., Hong, Y. and Balakrishnan, N. (2018). The generalized Poisson-binomial distribution and the computation of its distribution function. *Journal of Statistical Computational and Simulation*, **88**(8), pp. 1515-1527. doi:10.1080/00949655.2018.1440294

Examples

```
set.seed(1)
pp <- c(1, 0, runif(10), 1, 0, 1)
qq <- seq(0, 1, 0.01)
va <- rep(5, length(pp))
vb <- 1:length(pp)

dgpbinom(NULL, pp, va, vb, method = "DivideFFT")
pgpbinom(75:100, pp, va, vb, method = "DivideFFT")
qgpbinom(qq, pp, va, vb, method = "DivideFFT")
rgpbinom(100, pp, va, vb, method = "DivideFFT")

dgpbinom(NULL, pp, va, vb, method = "Convolve")
pgpbinom(75:100, pp, va, vb, method = "Convolve")
qgpbinom(qq, pp, va, vb, method = "Convolve")
rgpbinom(100, pp, va, vb, method = "Convolve")

dgpbinom(NULL, pp, va, vb, method = "Characteristic")
pgpbinom(75:100, pp, va, vb, method = "Characteristic")
qgpbinom(qq, pp, va, vb, method = "Characteristic")
rgpbinom(100, pp, va, vb, method = "Characteristic")
```

```
dgpbinom(NULL, pp, va, vb, method = "Normal")
pgpbinom(75:100, pp, va, vb, method = "Normal")
qgpbinom(qq, pp, va, vb, method = "Normal")
rgpbinom(100, pp, va, vb, method = "Normal")

dgpbinom(NULL, pp, va, vb, method = "RefinedNormal")
pgpbinom(75:100, pp, va, vb, method = "RefinedNormal")
qgpbinom(qq, pp, va, vb, method = "RefinedNormal")
rgpbinom(100, pp, va, vb, method = "RefinedNormal")
```

PoissonBinomial-Distribution

The Poisson Binomial Distribution

Description

Density, distribution function, quantile function and random generation for the Poisson binomial distribution with probability vector probs.

Usage

```
dpbinom(x, probs, wts = NULL, method = "DivideFFT", log = FALSE)

ppbinom(
  x,
  probs,
  wts = NULL,
  method = "DivideFFT",
  lower.tail = TRUE,
  log.p = FALSE
)

qpbinom(
  p,
  probs,
  wts = NULL,
  method = "DivideFFT",
  lower.tail = TRUE,
  log.p = FALSE
)

rpbinom(n, probs, wts = NULL, method = "DivideFFT", generator = "Sample")
```

Arguments

x	Either a vector of observed numbers of successes or NULL. If NULL, probabilities of all possible observations are returned.

probs	Vector of probabilities of success of each Bernoulli trial.
wts	Vector of non-negative integer weights for the input probabilities.
method	Character string that specifies the method of computation and must be one of "DivideFFT", "Convolve", "Characteristic", "Recursive", "Mean", "GeoMean", "GeoMeanCounter", "Poisson", "Normal" or "RefinedNormal" (abbreviations are allowed).
log, log.p	Logical value indicating if results are given as logarithms.
lower.tail	Logical value indicating if results are $P[X \leq x]$ (if TRUE; default) or $P[X > x]$ (if FALSE).
p	Vector of probabilities for computation of quantiles.
n	Number of observations. If length(n) > 1, the length is taken to be the number required.
generator	Character string that specifies the random number generator and must either be "Sample" (default) or "Bernoulli" (abbreviations are allowed). See Details for more information.

Details

See the references for computational details. The *Divide and Conquer* ("DivideFFT") and *Direct Convolution* ("Convolve") algorithms are derived and described in Biscarri, Zhao & Brunner (2018). The *Discrete Fourier Transformation of the Characteristic Function* ("Characteristic"), the *Recursive Formula* ("Recursive"), the *Poisson Approximation* ("Poisson"), the *Normal Approach* ("Normal") and the *Refined Normal Approach* ("RefinedNormal") are described in Hong (2013). The calculation of the *Recursive Formula* was modified to overcome the excessive memory requirements of Hong's implementation.

The "Mean" method is a naive binomial approach using the arithmetic mean of the probabilities of success. Similarly, the "GeoMean" and "GeoMeanCounter" procedures are binomial approximations, too, but they form the geometric mean of the probabilities of success ("GeoMean") and their counter probabilities ("GeoMeanCounter"), respectively.

In some special cases regarding the values of probs, the method parameter is ignored (see Introduction vignette).

Random numbers can be generated in two ways. The "Sample" method uses R's sample function to draw random values according to their probabilities that are calculated by dgpbinom. The "Bernoulli" procedure ignores the method parameter and simulates Bernoulli-distributed random numbers according to the probabilities in probs and sums them up. It is a bit slower than the "Sample" generator, but may yield better results, as it allows to obtain observations that cannot be generated by the "Sample" procedure, because dgpbinom may compute 0-probabilities, due to rounding, if the length of probs is large and/or its values contain a lot of very small values.

Value

dpbinom gives the density, ppbinom computes the distribution function, qpbinom gives the quantile function and rpbinom generates random deviates.

For rpbinom, the length of the result is determined by n, and is the lengths of the numerical arguments for the other functions.

References

Hong, Y. (2013). On computing the distribution function for the Poisson binomial distribution. *Computational Statistics & Data Analysis*, **59**, pp. 41-51. doi:10.1016/j.csda.2012.10.006

Biscarri, W., Zhao, S. D. and Brunner, R. J. (2018) A simple and fast method for computing the Poisson binomial distribution. *Computational Statistics and Data Analysis*, **31**, pp. 216–222. doi:10.1016/j.csda.2018.01.007

Examples

```
set.seed(1)
pp <- c(0, 0, runif(995), 1, 1, 1)
qq <- seq(0, 1, 0.01)

dpbinom(NULL, pp, method = "DivideFFT")
ppbinom(450:550, pp, method = "DivideFFT")
qpbinom(qq, pp, method = "DivideFFT")
rpbinom(100, pp, method = "DivideFFT")

dpbinom(NULL, pp, method = "Convolve")
ppbinom(450:550, pp, method = "Convolve")
qpbinom(qq, pp, method = "Convolve")
rpbinom(100, pp, method = "Convolve")

dpbinom(NULL, pp, method = "Characteristic")
ppbinom(450:550, pp, method = "Characteristic")
qpbinom(qq, pp, method = "Characteristic")
rpbinom(100, pp, method = "Characteristic")

dpbinom(NULL, pp, method = "Recursive")
ppbinom(450:550, pp, method = "Recursive")
qpbinom(qq, pp, method = "Recursive")
rpbinom(100, pp, method = "Recursive")

dpbinom(NULL, pp, method = "Mean")
ppbinom(450:550, pp, method = "Mean")
qpbinom(qq, pp, method = "Mean")
rpbinom(100, pp, method = "Mean")

dpbinom(NULL, pp, method = "GeoMean")
ppbinom(450:550, pp, method = "GeoMean")
qpbinom(qq, pp, method = "GeoMean")
rpbinom(100, pp, method = "GeoMean")

dpbinom(NULL, pp, method = "GeoMeanCounter")
ppbinom(450:550, pp, method = "GeoMeanCounter")
qpbinom(qq, pp, method = "GeoMeanCounter")
rpbinom(100, pp, method = "GeoMeanCounter")

dpbinom(NULL, pp, method = "Poisson")
ppbinom(450:550, pp, method = "Poisson")
qpbinom(qq, pp, method = "Poisson")
```

```
rpbinom(100, pp, method = "Poisson")

dpbinom(NULL, pp, method = "Normal")
ppbinom(450:550, pp, method = "Normal")
qpbinom(qq, pp, method = "Normal")
rpbinom(100, pp, method = "Normal")

dpbinom(NULL, pp, method = "RefinedNormal")
ppbinom(450:550, pp, method = "RefinedNormal")
qpbinom(qq, pp, method = "RefinedNormal")
rpbinom(100, pp, method = "RefinedNormal")
```

Index

dgpbinom
(GenPoissonBinomial-Distribution), 3
dpbinom (PoissonBinomial-Distribution), 6

GenPoissonBinomial-Distribution, 3

pgpbinom
(GenPoissonBinomial-Distribution), 3
PoissonBinomial
(PoissonBinomial-package), 2
PoissonBinomial-Distribution, 6
PoissonBinomial-package, 2
ppbinom (PoissonBinomial-Distribution), 6

qgpbinom
(GenPoissonBinomial-Distribution), 3
qpbinom (PoissonBinomial-Distribution), 6

rgpbinom
(GenPoissonBinomial-Distribution), 3
rpbinom (PoissonBinomial-Distribution), 6