# Package 'PMD'

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Type Package

**Title** Computation of Poisson-Multinomial Distribtuions

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Maintainer Yili Hong <yilihong@vt.edu></yilihong@vt.edu>
Description Applying exact, simulation and approximation methods for computing probability mass functions, cumulative distribution functions of Poisson-Multinomial distributions together with a random number generator to conduct Poisson-Multinomial distribution sampling. The exact method is based on fast Fourier transformation of the characteristic functions of Poisson-Multinomial distributions. Simulation method is built via naive sampling scheme of different multinomial distributions. Approximation method is designed by asymptotic distributions of Poisson-Multinomial distributions, which are normal.
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LinkingTo Rcpp, RcppArmadillo
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Author Yili Hong [aut, ctb], Zhengzhi Lin [aut, cre], Yueyao Wang [aut, ctb]
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dpmd

Probability Mass of Poisson-Multinomial Distributions

## **Description**

Computation of probability mass for Poisson-Multinomial Distributions using exact, simulation, approximation methods. Users are allowed to specified a method and can choose to compute single mass point or all mass points. For simulation method, users can also choose the repeating time to enhance the accuracy of outcomes.

### Usage

```
dpmd(pp, method = "DFT-CF", vec = c(0, 0, 0, 0, 0), B = 100)
```

## **Arguments**

pp A matrix of probabilities. Each row of pp should add up to 1.

method Method selected by user to compute the probability mass. There are totally 4

methods. "DFT-CF": An exact method to calculate all probability mass points of Poisson-Multinomial Distributions via FFT algorithm. "simulation": A simulation method calculating all probability mass points. "NA by demands": An approximation method using Normal approximation to compute the probability for the 'vec' vector input by user. "simulation by demands": The same simulation method as above just to compute single probability mass point as input

by user.

vec Result vector(probability mass point) specified by user. eg. pp is 4 by 3 matrix

then a user might be interested in the probability of getting result: vec=c(0,0,1,2).

B Simulation repeat time.

## Value

For a single mass point, dpmd returns a probability. For all probability mass points of a given pp, it returns a multi-dimensional array. For instance, for the pp matrix in the following example, the value of the array element  $a_{1,2,1} = 0.90$  means the value of probability mass point (0,1,0,2) is 0.90.

# **Examples**

ppmd 3

ppmd

cumulative distribution function of PMN

## **Description**

```
By an input vector \mathbf{x} = (x_1, x_2, ...), this function compute P(X_1 \le x_1, X_2 \le x_2, ...).
```

## Usage

```
ppmd(pp, x, method = "DFT-CF", B = 1000)
```

## **Arguments**

pp input matrix of probabilities

x input result vector

method method selected by users to compute the cumulative mass probabilities.

B repeating time

#### Value

cumulative distribution of given pp.

# **Examples**

rpmd

generate random number from PMD

## **Description**

generate random number from PMD

## Usage

```
rpmd(pp)
```

## Arguments

рр

input matrix of probabilities

# Value

the random number vector generated from PMD.

## **Examples**

```
pp=matrix(c(.1, .1, .1, .7, .1, .3, .3, .3, .5, .2, .1, .2), nrow=3, byrow=TRUE) rpmd(pp)
```

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