

Package ‘NonNorMvtDist’

October 12, 2022

Type Package

Title Multivariate Lomax (Pareto Type II) and Its Related Distributions

Version 1.0.2

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Description Implements calculation of probability density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for the following multivariate distributions: Lomax (Pareto Type II), generalized Lomax, Mardia’s Pareto of Type I, Logistic, Burr, Cook-Johnson’s uniform, F and Inverted Beta. See Tapan Nayak (1987) <doi:10.2307/3214068>.

Depends R (>= 3.6.0)

License GPL-3

Encoding UTF-8

LazyData true

Imports stats, cubature

RoxxygenNote 7.0.2

NeedsCompilation no

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Repository CRAN

Date/Publication 2020-03-23 15:40:02 UTC

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MvtBurr*Multivariate Burr Distribution*

Description

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for multivariate Burr distribution with a scalar parameter `parm1` and vectors of parameters `parm2` and `parm3`.

Usage

```
dmvburr(x, parm1 = 1, parm2 = rep(1, k), parm3 = rep(1, k), log = FALSE)

pmvburr(q, parm1 = 1, parm2 = rep(1, k), parm3 = rep(1, k))

qmvburr(
  p,
  parm1 = 1,
  parm2 = rep(1, k),
  parm3 = rep(1, k),
  interval = c(0, 1e+08)
)

rmvburr(n, parm1 = 1, parm2 = rep(1, k), parm3 = rep(1, k))

smvburr(q, parm1 = 1, parm2 = rep(1, k), parm3 = rep(1, k))
```

Arguments

- `x` vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
- `parm1` a scalar parameter, see parameter a in **Details**.
- `parm2` a vector of parameters, see parameters d_i in **Details**.
- `parm3` a vector of parameters, see parameters c_i in **Details**.
- `log` logical; if TRUE, probability densities f are given as $\log(f)$.
- `q` a vector of quantiles.
- `p` a scalar value corresponding to probability.
- `interval` a vector containing the end-points of the interval to be searched. Default value is set as `c(0, 1e8)`.
- `n` number of observations.
- `k` dimension of data or number of variates.

Details

Multivariate Burr distribution (Johnson and Kotz, 1972) is a joint distribution of positive random variables X_1, \dots, X_k . Its probability density is given as

$$f(x_1, \dots, x_k) = \frac{[\prod_{i=1}^k c_i d_i] a(a+1) \cdots (a+k-1) [\prod_{i=1}^k x_i^{c_i-1}]}{(1 + \sum_{i=1}^k d_i x_i^{c_i})^{a+k}},$$

where $x_i > 0, a, c_i, d_i > 0, i = 1, \dots, k$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is obtained by the following formula related to survival function $\bar{F}(x_1, \dots, x_k)$ (Joe, 1997)

$$F(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} \bar{F}_S(x_j, j \in S),$$

where the survival function is given by

$$\bar{F}(x_1, \dots, x_k) = \left(1 + \sum_{i=1}^k d_i x_i^{c_i} \right)^{-a}.$$

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function [uniroot](#):

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

Random numbers X_1, \dots, X_k from multivariate Burr distribution can be generated through transformation of multivariate Lomax random variables Y_1, \dots, Y_k by letting $X_i = (\theta_i Y_i / d_i)^{1/c_i}, i = 1, \dots, k$; see Nayak (1987).

Value

`dmvburr` gives the numerical values of the probability density.

`pmvburr` gives the cumulative probability.

`qmvburr` gives the equicoordinate quantile.

`rmvburr` generates random numbers.

`smvburr` gives the value of survival function.

References

Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.

Johnson, N. L. and Kotz, S. (1972). *Distribution in Statistics: Continuous Multivariate Distributions*. New York: John Wiley & Sons, INC.

Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the multivariate Burr with parameters:
# a = 3, d1 = 1, d2 = 3, d3 = 5, c1 = 2, c2 = 4, c3 = 6
# Vector of quantiles: c(3, 2, 1)

dmvburr(x = c(3, 2, 1), parm1 = 3, parm2 = c(1, 3, 5), parm3 = c(2, 4, 6)) # Density

pmvburr(q = c(3, 2, 1), parm1 = 3, parm2 = c(1, 3, 5), parm3 = c(2, 4, 6)) # Cumulative Probability

# Equicoordinate quantile of cumulative probability 0.5
qmvburr(p = 0.5, parm1 = 3, parm2 = c(1, 3, 5), parm3 = c(2, 4, 6))

# Random numbers generation with sample size 100
rmvburr(n = 100, parm1 = 3, parm2 = c(1, 3, 5), parm3 = c(2, 4, 6))

smvburr(q = c(3, 2, 1), parm1 = 3, parm2 = c(1, 3, 5), parm3 = c(2, 4, 6)) # Survival function
```

Description

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for multivariate *F* distribution with degrees of freedom df.

Usage

```
dmvf(x, df = rep(1, k + 1), log = FALSE)

pmvf(q, df = rep(1, k + 1), algorithm = c("numerical", "MC"), nsim = 1e+07)

qmvf(
  p,
  df = rep(1, k + 1),
  interval = c(1e-08, 1e+08),
  algorithm = c("numerical", "MC"),
  nsim = 1e+06
)

rmvf(n, df = rep(1, k + 1))

smvf(q, df = rep(1, k + 1), algorithm = c("numerical", "MC"), nsim = 1e+07)
```

Arguments

<code>x</code>	vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
<code>df</code>	a vector of $k+1$ degrees of freedom, see parameter $(2a, 2l_1, \dots, 2l_k)$ in Details .
<code>log</code>	logical; if TRUE, probability densities f are given as $\log(f)$.
<code>q</code>	a vector of quantiles.
<code>algorithm</code>	method to be used for calculating cumulative probability. Two options are provided as (i) numerical using adaptive multivariate integral and (ii) MC using Monte Carlo method. Recommend algorithm numerical for ($k \leq 4$) dimension and MC for ($k > 4$) dimension based on running time consumption. Default option is set as numerical.
<code>nsim</code>	number of simulations used in algorithm MC.
<code>p</code>	a scalar value corresponding to probability.
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(1e-8, 1e8)</code> .
<code>n</code>	number of observations.
<code>k</code>	dimension of data or number of variates.

Details

Multivariate F distribution (Johnson and Kotz, 1972) is a joint probability distribution of positive random variables and its probability density is given as

$$f(x_1, \dots, x_k) = \frac{[\prod_{i=1}^k (l_i/a)^{l_i}] \Gamma(\sum_{i=1}^k l_i + a) \prod_{i=1}^k x_i^{l_i-1}}{\Gamma(a)[\prod_{i=1}^k \Gamma(l_i)](1 + \sum_{i=1}^k \frac{l_i}{a} x_i)^{\sum_{i=1}^k l_i + a}},$$

where $x_i > 0, a > 0, l_i > 0, i = 1, \dots, k$. The degrees of freedom are $(2a, 2l_1, \dots, 2l_k)$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is obtained by multiple integral

$$F(x_1, \dots, x_k) = \int_0^{x_1} \cdots \int_0^{x_k} f(y_1, \dots, y_k) dy_k \cdots dy_1.$$

This multiple integral is calculated by either adaptive multivariate integration using `hcubature` in package `cubature` (Narasimhan et al., 2018) or via Monte Carlo method.

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function `uniroot`:

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

The survival function $\bar{F}(x_1, \dots, x_k)$ is obtained either by the following formula related to cumulative distribution function $F(x_1, \dots, x_k)$ (Joe, 1997)

$$\bar{F}(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} F_S(x_j, j \in S),$$

or via Monte Carlo method.

Random numbers X_1, \dots, X_k from multivariate F distribution can be generated through parameter substitutions in simulation of generalized multivariate Lomax distribution by letting $\theta_i = l_i/a$, $i = 1, \dots, k$; see Nayak (1987).

Value

`dmvf` gives the numerical values of the probability density.

`pmvf` gives a list of two items:

value cumulative probability

`error` the estimated relative error by `algorithm = "numerical"` or the estimated standard error by `algorithm = "MC"`

`qmvf` gives the equicoordinate quantile. `NaN` is returned for no solution found in the given interval. The result is seed dependent if Monte Carlo algorithm is chosen (`algorithm = "MC"`).

`rmvf` generates random numbers.

`smvf` gives a list of two items:

value the value of survival function

`error` the estimated relative error by `algorithm = "numerical"` or the estimated standard error by `algorithm = "MC"`

References

Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.

Johnson, N. L. and Kotz, S. (1972). *Distribution in Statistics: Continuous Multivariate Distributions*. New York: John Wiley & Sons, INC.

Narasimhan, B., Koller, M., Johnson, S. G., Hahn, T., Bouvier, A., Kiêu, K. and Gaure, S. (2018). cubature: Adaptive Multivariate Integration over Hypercubes. R package version 2.0.3.

Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the multivariate F with degrees of freedom:
# df = c(2, 4, 6)
# Vector of quantiles: c(1, 2)

dmvf(x = c(1, 2), df = c(2, 4, 6)) # Density

# Cumulative Probability using adaptive multivariate integral
pmvf(q = c(1, 2), df = c(2, 4, 6), algorithm = "numerical")
```

```
# Cumulative Probability using Monte Carlo method
pmvf(q = c(1, 2), df = c(2, 4, 6), algorithm = "MC")

# Equicoordinate quantile of cumulative probability 0.5
qmvf(p = 0.5, df = c(2, 4, 6))

# Random numbers generation with sample size 100
rmvf(n = 100, df = c(2, 4, 6))

smvf(q = c(1, 2), df = c(2, 4, 6)) # Survival function
```

Description

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for generalized multivariate Lomax distribution with a scalar parameter `parm1` and vectors of parameters `parm2` and `parm3`.

Usage

```
dmvglomax(x, parm1 = 1, parm2 = rep(1, k), parm3 = rep(1, k), log = FALSE)

pmvglomax(
  q,
  parm1 = 1,
  parm2 = rep(1, k),
  parm3 = rep(1, k),
  algorithm = c("numerical", "MC"),
  nsim = 1e+07
)

qmvglomax(
  p,
  parm1 = 1,
  parm2 = rep(1, k),
  parm3 = rep(1, k),
  interval = c(1e-08, 1e+08),
  algorithm = c("numerical", "MC"),
  nsim = 1e+06
)

rmvglomax(n, parm1 = 1, parm2 = rep(1, k), parm3 = rep(1, k))

smvglomax(
```

```

q,
parm1 = 1,
parm2 = rep(1, k),
parm3 = rep(1, k),
algorithm = c("numerical", "MC"),
nsim = 1e+07
)

```

Arguments

<code>x</code>	vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
<code>parm1</code>	a scalar parameter, see parameter a in Details .
<code>parm2</code>	a vector of parameters, see parameters θ_i in Details .
<code>parm3</code>	a vector of parameters, see parameters l_i in Details .
<code>log</code>	logical; if TRUE, probability densities f are given as $\log(f)$.
<code>q</code>	a vector of quantiles.
<code>algorithm</code>	method to be used for calculating cumulative probability. Two options are provided as (i) numerical using adaptive multivariate integral and (ii) MC using Monte Carlo method. Recommend algorithm numerical for ($k \leq 4$) dimension and MC for ($k > 4$) dimension based on running time consumption. Default option is set as numerical.
<code>nsim</code>	number of simulations used in algorithm MC.
<code>p</code>	a scalar value corresponding to probability.
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(1e-8, 1e8)</code> .
<code>n</code>	number of observations.
<code>k</code>	dimension of data or number of variates.

Details

Generalized multivariate Lomax (Pareto type II) distribution was introduced by Nayak (1987) as a joint probability distribution of several skewed nonnegative random variables X_1, X_2, \dots, X_k . Its probability density function is given by

$$f(x_1, \dots, x_k) = \frac{[\prod_{i=1}^k \theta_i^{l_i}] \Gamma(\sum_{i=1}^k l_i + a) \prod_{i=1}^k x_i^{l_i-1}}{\Gamma(a)[\prod_{i=1}^k \Gamma(l_i)](1 + \sum_{i=1}^k \theta_i x_i)^{\sum_{i=1}^k l_i + a}},$$

where $x_i > 0, a, \theta_i, l_i > 0, i = 1, \dots, k$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is obtained by multiple integral

$$F(x_1, \dots, x_k) = \int_0^{x_1} \cdots \int_0^{x_k} f(y_1, \dots, y_k) dy_k \cdots dy_1.$$

This multiple integral is calculated by either adaptive multivariate integration using [hcubature](#) in package [cubature](#) (Narasimhan et al., 2018) or via Monte Carlo method.

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function [uniroot](#):

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

The survival function $\bar{F}(x_1, \dots, x_k)$ is obtained either by the following formula related to cumulative distribution function $F(x_1, \dots, x_k)$ (Joe, 1997)

$$\bar{F}(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} F_S(x_j, j \in S),$$

or via Monte Carlo method.

Random numbers from generalized multivariate Lomax distribution can be generated by simulating k independent gamma random variables having a common parameter following gamma distribution with shape parameter a and scale parameter 1; see Nayak (1987).

Value

`dmvglomax` gives the numerical values of the probability density.

`pmvglomax` gives a list of two items:

value cumulative probability

`error` the estimated relative error by `algorithm = "numerical"` or the estimated standard error by `algorithm = "MC"`

`qmvglomax` gives the equicoordinate quantile. `NaN` is returned for no solution found in the given interval. The result is seed dependent if Monte Carlo algorithm is chosen (`algorithm = "MC"`).

`rmvglomax` generates random numbers.

`smvglomax` gives a list of two items:

value the value of survival function

`error` the estimated relative error by `algorithm = "numerical"` or the estimated standard error by `algorithm = "MC"`

References

Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.

Narasimhan, B., Koller, M., Johnson, S. G., Hahn, T., Bouvier, A., Kiêu, K. and Gaure, S. (2018). cubature: Adaptive Multivariate Integration over Hypercubes. R package version 2.0.3.

Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the generalized multivariate Lomax with parameters:  
# a = 5, theta1 = 1, theta2 = 2, l1 = 4, l2 = 5  
# Vector of quantiles: c(5, 6)  
  
dmvglomax(x = c(5, 6), parm1 = 5, parm2 = c(1, 2), parm3 = c(4, 5)) # Density  
  
# Cumulative Probability using adaptive multivariate integral  
pmvglomax(q = c(5, 6), parm1 = 5, parm2 = c(1, 2), parm3 = c(4, 5))  
  
# Cumulative Probability using Monte Carlo method  
pmvglomax(q = c(5, 6), parm1 = 5, parm2 = c(1, 2), parm3 = c(4, 5), algorithm = "MC")  
  
# Equicoordinate quantile of cumulative probability 0.5  
qmvglomax(p = 0.5, parm1 = 5, parm2 = c(1, 2), parm3 = c(4, 5))  
  
# Random numbers generation with sample size 100  
rmvglomax(n = 100, parm1 = 5, parm2 = c(1, 2), parm3 = c(4, 5))  
  
smvglomax(q = c(5, 6), parm1 = 5, parm2 = c(1, 2), parm3 = c(4, 5)) # Survival function
```

Description

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for multivariate inverted beta distribution with a scalar parameter `parm1` and a vector of parameters `parm2`.

Usage

```
dmvinvbeta(x, parm1 = 1, parm2 = rep(1, k), log = FALSE)  
  
pmvinvbeta(  
  q,  
  parm1 = 1,  
  parm2 = rep(1, k),  
  algorithm = c("numerical", "MC"),  
  nsim = 1e+07  
)  
  
qmvinvbeta(  
  p,
```

```

parm1 = 1,
parm2 = rep(1, k),
interval = c(1e-08, 1e+08),
algorithm = c("numerical", "MC"),
nsim = 1e+06
)

rmvinvbeta(n, parm1 = 1, parm2 = rep(1, k))

smvinvbeta(
  q,
  parm1 = 1,
  parm2 = rep(1, k),
  algorithm = c("numerical", "MC"),
  nsim = 1e+07
)

```

Arguments

<code>x</code>	vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
<code>parm1</code>	a scalar parameter, see parameter a in Details .
<code>parm2</code>	a vector of parameters, see parameter l_i in Details .
<code>log</code>	logical; if TRUE, probability densities f are given as $\log(f)$.
<code>q</code>	a vector of quantiles.
<code>algorithm</code>	method to be used for calculating cumulative probability. Two options are provided as (i) numerical using adaptive multivariate integral and (ii) MC using Monte Carlo method. Recommend algorithm numerical for ($k \leq 4$) dimension and MC for ($k > 4$) dimension based on running time consumption. Default option is set as <code>numerical</code> .
<code>nsim</code>	number of simulations used in algorithm MC.
<code>p</code>	a scalar value corresponding to probability.
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(1e-8, 1e8)</code> .
<code>n</code>	number of observations.
<code>k</code>	dimension of data or number of variates.

Details

Multivariate inverted beta distribution is an alternative expression of multivariate F distribution and is a special case of multivariate Lomax distribution (Balakrishnan and Lai, 2009). Its probability density is given as

$$f(x_1, \dots, x_p) = \frac{\Gamma(\sum_{i=1}^p l_i + a) \prod_{i=1}^p x_i^{l_i - 1}}{\Gamma(a)[\prod_{i=1}^p \Gamma(l_i)](1 + \sum_{i=1}^p x_i)^{\sum_{i=1}^p l_i + a}},$$

where $x_i > 0, a > 0, l_i > 0, i = 1, \dots, p$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is obtained by multiple integral

$$F(x_1, \dots, x_k) = \int_0^{x_1} \cdots \int_0^{x_k} f(y_1, \dots, y_k) dy_k \cdots dy_1.$$

This multiple integral is calculated by either adaptive multivariate integration using `hcubature` in package `cubature` (Narasimhan et al., 2018) or via Monte Carlo method.

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function `uniroot`:

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

The survival function $\bar{F}(x_1, \dots, x_k)$ is obtained either by the following formula related to cumulative distribution function $F(x_1, \dots, x_k)$ (Joe, 1997)

$$\bar{F}(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} F_S(x_j, j \in S),$$

or via Monte Carlo method.

Random numbers X_1, \dots, X_k from multivariate inverted beta distribution can be generated through parameter substitutions in simulation of generalized multivariate Lomax distribution by letting $\theta_i = 1, i = 1, \dots, k$.

Value

`dmvinvbeta` gives the numerical values of the probability density.

`pmvinvbeta` gives a list of two items:

value cumulative probability

`error` the estimated relative error by `algorithm = "numerical"` or the estimated standard error by `algorithm = "MC"`

`qmvinvbeta` gives the equicoordinate quantile. NaN is returned for no solution found in the given interval. The result is seed dependent if Monte Carlo algorithm is chosen (`algorithm = "MC"`).

`rmvinvbeta` generates random numbers.

`smvinvbeta` gives a list of two items:

value the value of survival function

`error` the estimated relative error by `algorithm = "numerical"` or the estimated standard error by `algorithm = "MC"`

References

Balakrishnan, N. and Lai, C. (2009). *Continuous Bivariate Distributions. 2nd Edition*. New York: Springer.

Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.

Narasimhan, B., Koller, M., Johnson, S. G., Hahn, T., Bouvier, A., Kiêu, K. and Gaure, S. (2018). cubature: Adaptive Multivariate Integration over Hypercubes. R package version 2.0.3.

Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the multivariate inverted beta with parameters:  
# a = 7, l1 = 1, l2 = 3  
# Vector of quantiles: c(2, 4)  
  
dmvinvbeta(x = c(2, 4), parm1 = 7, parm2 = c(1, 3)) # Density  
  
# Cumulative Probability using adaptive multivariate integral  
pmvinvbeta(q = c(2, 4), parm1 = 7, parm2 = c(1, 3))  
  
# Cumulative Probability using Monte Carlo method  
pmvinvbeta(q = c(2, 4), parm1 = 7, parm2 = c(1, 3), algorithm = "MC")  
  
# Equicoordinate quantile of cumulative probability 0.5  
qmvinvbeta(p = 0.5, parm1 = 7, parm2 = c(1, 3))  
  
# Random numbers generation with sample size 100  
rmvinvbeta(n = 100, parm1 = 7, parm2 = c(1, 3))  
  
smvinvbeta(q = c(2, 4), parm1 = 7, parm2 = c(1, 3)) # Survival function
```

Description

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for multivariate logistic distribution with vector parameter `parm1` and vector parameter `parm2`.

Usage

```
dmvlogis(x, parm1 = rep(1, k), parm2 = rep(1, k), log = FALSE)  
  
pmvlogis(q, parm1 = rep(1, k), parm2 = rep(1, k))
```

```
qmvlogis(p, parm1 = rep(1, k), parm2 = rep(1, k), interval = c(0, 1e+08))

rmvlogis(n, parm1 = rep(1, k), parm2 = rep(1, k))

smvlogis(q, parm1 = rep(1, k), parm2 = rep(1, k))
```

Arguments

<code>x</code>	vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
<code>parm1</code>	a vector of location parameters, see parameter μ_i in Details .
<code>parm2</code>	a vector of scale parameters, see parameters σ_i in Details .
<code>log</code>	logical; if TRUE, probability densities f are given as $\log(f)$.
<code>q</code>	a vector of quantiles.
<code>p</code>	a scalar value corresponding to probability.
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(0, 1e8)</code> .
<code>n</code>	number of observations.
<code>k</code>	dimension of data or number of variates.

Details

Bivariate logistic distribution was introduced by Gumbel (1961) and its multivariate generalization was given by Malik and Abraham (1973) as

$$f(x_1, \dots, x_k) = \frac{k! \exp\left(-\sum_{i=1}^k \frac{x_i - \mu_i}{\sigma_i}\right)}{[\prod_{i=1}^k \sigma_i][1 + \sum_{i=1}^k \exp\left(-\frac{x_i - \mu_i}{\sigma_i}\right)]^{1+k}},$$

where $-\infty < x_i, \mu_i < \infty, \sigma_i > 0, i = 1, \dots, k$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is given as

$$F(x_1, \dots, x_k) = \left[1 + \sum_{i=1}^k \exp\left(-\frac{x_i - \mu_i}{\sigma_i}\right) \right]^{-1}.$$

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function [uniroot](#):

$$\int_{-\infty}^q \cdots \int_{-\infty}^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

The survival function $\bar{F}(x_1, \dots, x_k)$ is obtained by the following formula related to cumulative distribution function $F(x_1, \dots, x_k)$ (Joe, 1997)

$$\bar{F}(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} F_S(x_j, j \in S).$$

Random numbers X_1, \dots, X_k from multivariate logistic distribution can be generated through transformation of multivariate Lomax random variables Y_1, \dots, Y_k by letting $X_i = \mu_i - \sigma_i \ln(\theta_i Y_i)$, $i = 1, \dots, k$; see Nayak (1987).

Value

`dmvlogis` gives the numerical values of the probability density.
`pmvlogis` gives the cumulative probability.
`qmvlogis` gives the equicoordinate quantile.
`rmvlogis` generates random numbers.
`smvlogis` gives the value of survival function

References

- Gumbel, E.J. (1961). Bivariate logistic distribution. *J. Am. Stat. Assoc.*, 56, 335-349.
- Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.
- Malik, H. J. and Abraham, B. (1973). Multivariate logistic distributions. *Ann. Statist.* 3, 588-590.
- Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the multivariate logistic distribution with parameters:  

# mu1 = 0.5, mu2 = 1, mu3 = 2, sigma1 = 1, sigma2 = 2 and sigma3 = 3  

# Vector of quantiles: c(3, 2, 1)  

  

dmvlogis(x = c(3, 2, 1), parm1 = c(0.5, 1, 2), parm2 = c(1, 2, 3)) # Density  

  

pmvlogis(q = c(3, 2, 1), parm1 = c(0.5, 1, 2), parm2 = c(1, 2, 3)) # Cumulative Probability  

  

# Equicoordinate quantile of cumulative probability 0.5  

qmvlogis(p = 0.5, parm1 = c(0.5, 1, 2), parm2 = c(1, 2, 3))  

  

# Random numbers generation with sample size 100  

rmvlogis(n = 100, parm1 = c(0.5, 1, 2), parm2 = c(1, 2, 3))  

  

smvlogis(q = c(3, 2, 1), parm1 = c(0.5, 1, 2), parm2 = c(1, 2, 3)) # Survival function
```

MvtLomax*Multivariate Lomax (Pareto Type II) Distribution***Description**

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for multivariate Lomax (Pareto Type II) distribution with a scalar parameter `parm1` and vector parameter `parm2`.

Usage

```
dmvlomax(x, parm1 = 1, parm2 = rep(1, k), log = FALSE)

pmvlomax(q, parm1 = 1, parm2 = rep(1, k))

qmvvlomax(p, parm1 = 1, parm2 = rep(1, k), interval = c(0, 1e+08))

rmvlomax(n, parm1 = 1, parm2 = rep(1, k))

smvlomax(q, parm1 = 1, parm2 = rep(1, k))
```

Arguments

<code>x</code>	vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
<code>parm1</code>	a scalar parameter, see parameter a in Details .
<code>parm2</code>	a vector of parameters, see parameters θ_i in Details .
<code>log</code>	logical; if TRUE, probability densities f are given as $\log(f)$.
<code>q</code>	a vector of quantiles.
<code>p</code>	a scalar value corresponding to probability.
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(0, 1e8)</code> .
<code>n</code>	number of observations.
<code>k</code>	dimension of data or number of variates.

Details

Multivariate Lomax (Pareto type II) distribution was introduced by Nayak (1987) as a joint probability distribution of several skewed positive random variables X_1, X_2, \dots, X_k . Its probability density function is given by

$$f(x_1, x_2, \dots, x_k) = \frac{[\prod_{i=1}^k \theta_i] a(a+1) \cdots (a+k-1)}{(1 + \sum_{i=1}^k \theta_i x_i)^{a+k}},$$

where $x_i > 0, a > 0, \theta_i > 0, i = 1, \dots, k$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is obtained by the following formula related to survival function $\bar{F}(x_1, \dots, x_k)$ (Joe, 1997)

$$F(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} \bar{F}_S(x_j, j \in S),$$

where the survival function is given by

$$\bar{F}(x_1, \dots, x_k) = (1 + \sum_{i=1}^k \theta_i x_i)^{-a}.$$

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function [uniroot](#):

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

Random numbers from multivariate Lomax distribution can be generated by simulating k independent exponential random variables having a common environment parameter following gamma distribution with shape parameter a and scale parameter 1; see Nayak (1987).

Value

`dmvlomax` gives the numerical values of the probability density.

`pmvlomax` gives the cumulative probability.

`qmvlomax` gives the equicoordinate quantile.

`rmvlomax` generates random numbers.

`smvlomax` gives the value of survival function.

References

Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.

Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the multivariate Lomax with parameters:
# a = 5, theta1 = 1, theta2 = 2 and theta3 = 3.
# Vector of quantiles: c(3, 2, 1)

dmvlomax(x = c(3, 2, 1), parm1 = 5, parm2 = c(1, 2, 3)) # Density
```

```

pmvlpmax(q = c(3, 2, 1), parm1 = 5, parm2 = c(1, 2, 3)) # Cumulative Probability

# Equicoordinate quantile of cumulative probability 0.5
qmvlpmax(p = 0.5, parm1 = 5, parm2 = c(1, 2, 3))

# Random numbers generation with sample size 100
rmvlpmax(n = 100, parm1 = 5, parm2 = c(1, 2, 3))

smvlpmax(q = c(3, 2, 1), parm1 = 5, parm2 = c(1, 2, 3)) # Survival function

```

MvtMardiaPareto1*Mardia's Multivariate Pareto Type I Distribution***Description**

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for Mardia's multivariate Pareto Type I distribution with a scalar parameter `parm1` and a vector of parameters `parm2`.

Usage

```

dmvmpareto1(x, parm1 = 1, parm2 = rep(1, k), log = FALSE)

pmvmpareto1(q, parm1 = 1, parm2 = rep(1, k))

qmvmpareto1(
  p,
  parm1 = 1,
  parm2 = rep(1, k),
  interval = c(max(1/parm2) + 1e-08, 1e+08)
)

rmvmpareto1(n, parm1 = 1, parm2 = rep(1, k))

smvmpareto1(q, parm1 = 1, parm2 = rep(1, k))

```

Arguments

- `x` vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
- `parm1` a scalar parameter, see parameter a in **Details**.
- `parm2` a vector of parameters, see parameters θ_i in **Details**.
- `log` logical; if TRUE, probability densities f are given as $\log(f)$.
- `q` a vector of quantiles.

<code>p</code>	a scalar value corresponding to probability.
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(max(1 / param2) + 1e-8, 1e8)</code> according to $x_i > 1/\theta_i, \theta_i > 0, i = 1, \dots, k$.
<code>n</code>	number of observations.
<code>k</code>	dimension of data or number of variates.

Details

Multivariate Pareto type I distribution was introduced by Mardia (1962) as a joint probability distribution of several nonnegative random variables X_1, \dots, X_k . Its probability density function is given by

$$f(x_1, \dots, x_k) = \frac{[\prod_{i=1}^k \theta_i] a(a+1) \cdots (a+k-1)}{(\sum_{i=1}^k \theta_i x_i - k+1)^{a+k}},$$

where $x_i > 1/\theta_i, a > 0, \theta_i > 0, i = 1, \dots, k$.

Cumulative distribution function $F(x_1, \dots, x_k)$ is obtained by the following formula related to survival function $\bar{F}(x_1, \dots, x_k)$ (Joe, 1997)

$$F(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} \bar{F}_S(x_j, j \in S),$$

where the survival function is given by

$$\bar{F}(x_1, \dots, x_k) = \left(\sum_{i=1}^k \theta_i x_i - k + 1 \right)^{-a}.$$

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function `uniroot`:

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

Random numbers X_1, \dots, X_k from Mardia's multivariate Pareto type I distribution can be generated through linear transformation of multivariate Lomax random variables Y_1, \dots, Y_k by letting $X_i = Y_i + 1/\theta_i, i = 1, \dots, k$; see Nayak (1987).

Value

`dmvmpareto1` gives the numerical values of the probability density.

`pmvmpareto1` gives the cumulative probability.

`qmvmpareto1` gives the equicoordinate quantile.

`rmvmpareto1` generates random numbers.

`smvmpareto1` gives the value of survival function.

References

- Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.
- Mardia, K. V. (1962). Multivariate Pareto distributions. *Ann. Math. Statist.* 33, 1008-1015.
- Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the Mardia's multivariate Pareto Type I with parameters:
# a = 5, theta1 = 1, theta2 = 2, theta3 = 3
# Vector of quantiles: c(2, 1, 1)

dmvmpareto1(x = c(2, 1, 1), parm1 = 5, parm2 = c(1, 2, 3)) # Density

pmvmpareto1(q = c(2, 1, 1), parm1 = 5, parm2 = c(1, 2, 3)) # Cumulative Probability

# Equicoordinate quantile of cumulative probability 0.5
qmvmpareto1(p = 0.5, parm1 = 5, parm2 = c(1, 2, 3))

# Random numbers generation with sample size 100
rmvmpareto1(n = 100, parm1 = 5, parm2 = c(1, 2, 3))

smvmpareto1(q = c(2, 1, 1), parm1 = 5, parm2 = c(1, 2, 3)) # Survival function
```

Description

Calculation of density function, cumulative distribution function, equicoordinate quantile function and survival function, and random numbers generation for Cook-Johnson's multivariate uniform distribution with a scalar parameter `parm`.

Usage

```
dmvunif(x, parm = 1, log = FALSE)

pmvunif(q, parm = 1)

qmvunif(p, parm = 1, dim = k, interval = c(0, 1))

rmvunif(n, parm = 1, dim = 1)

smvunif(q, parm = 1)
```

Arguments

<code>x</code>	vector or matrix of quantiles. If x is a matrix, each row vector constitutes a vector of quantiles for which the density $f(x)$ is calculated (for i -th row x_i , $f(x_i)$ is reported).
<code>parm</code>	a scalar parameter, see parameter a in Details .
<code>log</code>	logical; if TRUE, probability densities f are given as $\log(f)$.
<code>q</code>	a vector of quantiles.
<code>p</code>	a scalar value corresponding to probability.
<code>dim</code>	dimension of data or number of variates (k).
<code>interval</code>	a vector containing the end-points of the interval to be searched. Default value is set as <code>c(0, 1)</code> .
<code>n</code>	number of observations.

Details

Multivariate uniform distribution of Cook and Johnson (1981) is a joint distribution of uniform variables over $(0, 1]$ and its probability density is given as

$$f(x_1, \dots, x_k) = \frac{\Gamma(a+k)}{\Gamma(a)a^k} \prod_{i=1}^k x_i^{(-1/a)-1} \left[\sum_{i=1}^k x_i^{-1/a} - k + 1 \right]^{-(a+k)},$$

where $0 < x_i \leq 1, a > 0, i = 1, \dots, k$. In fact, Cook-Johnson's uniform distribution is also called Clayton copula (Nelsen, 2006).

Cumulative distribution function $F(x_1, \dots, x_k)$ is given as

$$F(x_1, \dots, x_k) = \left[\sum_{i=1}^k x_i^{-1/a} - k + 1 \right]^{-a}.$$

Equicoordinate quantile is obtained by solving the following equation for q through the built-in one dimension root finding function `uniroot`:

$$\int_0^q \cdots \int_0^q f(x_1, \dots, x_k) dx_k \cdots dx_1 = p,$$

where p is the given cumulative probability.

The survival function $\bar{F}(x_1, \dots, x_k)$ is obtained by the following formula related to cumulative distribution function $F(x_1, \dots, x_k)$ (Joe, 1997)

$$\bar{F}(x_1, \dots, x_k) = 1 + \sum_{S \in \mathcal{S}} (-1)^{|S|} F_S(x_j, j \in S).$$

Random numbers X_1, \dots, X_k from Cook-Johnson's multivariate uniform distribution can be generated through transformation of multivariate Lomax random variables Y_1, \dots, Y_k by letting $X_i = (1 + \theta_i Y_i)^{-a}, i = 1, \dots, k$; see Nayak (1987).

Value

`dmvunif` gives the numerical values of the probability density.
`pmvunif` gives the cumulative probability.
`qmvunif` gives the equicoordinate quantile.
`rmvunif` generates random numbers.
`smvunif` gives the value of survival function.

References

- Cook, R. E. and Johnson, M. E. (1981). A family of distributions for modeling non-elliptically symmetric multivariate data. *J.R. Statist. Soc. B* 43, No. 2, 210-218.
- Joe, H. (1997). *Multivariate Models and Dependence Concepts*. London: Chapman & Hall.
- Nayak, T. K. (1987). Multivariate Lomax Distribution: Properties and Usefulness in Reliability Theory. *Journal of Applied Probability*, Vol. 24, No. 1, 170-177.
- Nelsen, R. B. (2006). *An Introduction to Copulas, Second Edition*. New York: Springer.

See Also

[uniroot](#) for one dimensional root (zero) finding.

Examples

```
# Calculations for the Cook-Johnson's multivariate uniform distribution with parameters:  

# a = 2, dim = 3  

# Vector of quantiles: c(0.8, 0.5, 0.2)  
  

dmvunif(x = c(0.8, 0.5, 0.2), parm = 2) # Density  
  

pmvunif(q = c(0.8, 0.5, 0.2), parm = 2) # Cumulative Probability  
  

# Equicoordinate quantile of cumulative probability 0.5  

qmvunif(p = 0.5, parm = 2, dim = 3)  
  

# Random numbers generation with sample size 100  

rmvunif(n = 100, parm = 2, dim = 3)  
  

smvunif(q = c(0.8, 0.5, 0.2), parm = 3) # Survival function
```

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