Package ‘subcopem2D’

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Title Bivariate Empirical Subcopula
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Description Calculate empirical subcopula and dependence measures from a given bivariate sample, and Bernstein copula approximations.
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R topics documented:

| Bcopula | Bernstein Copula Approximation |

Description

Bernstein copula approximation from the empirical subcopula of given bivariate data.

Usage

```
Bcopula(mat.xy, m, both.cont = FALSE, tolimit = 1e-05)
```
**Arguments**

- `mat.xy` 2-column matrix with bivariate observations of a random vector \((X, Y)\).
- `m` integer value of approximation order, where \(m = 2, \ldots, n\) with \(n\) equal to sample size. A recommended value for \(m\) would be the minimum between \(\sqrt{n}\) and 50.
- `both.cont` logical value, if TRUE then \((X, Y)\) are considered (both) as continuous random variables, and jittering will be applied to repeated values (if any).
- `tolimit` tolerance limit in numerical approximation of the inverse of the first partial derivatives of the estimated Bernstein copula.

**Details**

Each of the random variables \(X\) and \(Y\) may be of any kind (discrete, continuous, or mixed). NA values are not allowed.

**Value**

A list containing the following components:

- `copula` bivariate Bernstein Copula function (BC) of order \(m\).
- `du` bivariate function \(\partial BC(u, v)/\partial u\).
- `dv` bivariate function \(\partial BC(u, v)/\partial v\).
- `du.inv` inverse of `du` with respect to \(v\), given \(u\) and alpha (numerical approx).
- `dv.inv` inverse of `dv` with respect to \(u\), given \(v\) and alpha (numerical approx).
- `density` bivariate Bernstein copula density function of order \(m\).
- `bilinearCopula` bivariate function of bilinear approximation of copula.
- `bilinearSubcopula` \((m + 1) \times (m + 1)\) matrix with empirical subcopula values.
- `sample.size` sample size of bivariate observations.
- `order` approximation order \(m\) used.
- `both.cont` logical value, TRUE if both variables considered as continuous.
- `tolimit` tolerance limit in numerical approximation of \(du.inv\) and \(dv.inv\).
- `subcopemObject` list object with the output from `subcopem` if `both.cont = FALSE` or from `subcopemc` if `both.cont = TRUE`.

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**Note**

If both \(X\) and \(Y\) are continuous random variables it is faster and better to set `both.cont = TRUE`.

**Author(s)**

Arturo Erdely [https://sites.google.com/site/arturoerdely](https://sites.google.com/site/arturoerdely)
dependence

References


See Also

subcopem, subcopemc

Examples

## (X,Y) continuous random variables with copula FGM(param = 1)

# Theoretical formulas
FGMcopula <- function(u, v) u*v*(1 + (1 - u)*(1 - v))
dFGM.du <- function(u, v) (2*u - 1)*(v^2) + 2*v*(1 - u)
dFGM.dv <- function(u, v) (2*v - 1)*(u^2) + 2*u*(1 - v)
A1 <- function(u) 2*(1 - u)
A2 <- function(u, z) sqrt(A1(u)^2 - 4*(A1(u) - 1)*z)
dFGM.du.inv <- function(u, z) 2*z/(A1(u) + A2(u, z))
FGMdensity <- function(u, v) 2*(1 - u - v + 2*u*v)

# Simulating FGM observations
n <- 3000
U <- runif(n)
Z <- runif(n)
V <- mapply(dFGM.du.inv, U, Z)

# Applying Bcopula to FGM simulated values
B <- Bcopula(cbind(U, V), 50, TRUE)
str(B)

# Comparing theoretical values versus Bernstein and Bilinear approximations
u <- 0.70; v <- 0.55
FGMcopula(u, v); B["copula"](u, v); B["bilinearCopula"](u, v)
dFGM.du(u, v); B["du"](u, v)
dFGM.dv(u, v); B["dv"](u, v)
dFGM.du.inv(u, 0.8); B["du.inv"](u, 0.8)
FGMdensity(u, v); B["density"](u, v)

dependence

Dependence Measures

Description

Calculation of pairwise monotone and supremum dependence, monotone/supremum dependence ratio, and proportion of pairwise NAs.
dependence(mat, cont = NULL, sc.order = 0)

Arguments

mat
  k-column matrix with n observations of a k-dimensional random vector (NA values are allowed).
cont
  vector of column numbers to consider/coerce as continuous random variables (optional).
sc.order
  order of subcopula approximation (continuous random variables). If 0 (default) then maximum order m = n is used. Often m = 50 is a good recommended value, higher values demand more computing time.

Details

Each of the random variables in the k-dimensional random vector under consideration may be of any kind (discrete, continuous, or mixed). NA values are allowed.

Value

A 3-dimensional array k x k x 4 with pairwise monotone and supremum dependence, monotone/supremum dependence ratio, and proportion of pairwise NAs.

Note

NA values are allowed.

Author(s)

Arturo Erdely https://sites.google.com/site/arturoerdely

References


See Also

subcopem, subcopemc

Examples

V <- runif(300)  # Continuous Uniform(0,1)
W <- V*(1-V) # Continuous transform of V
# X given V=v as continuous Uniform(0,v)
X <- mapply(runif, rep(1, length(V)), rep(0, length(V)), V)
Y <- 1*(0.2 < X)*(X < 0.6) # Discrete transform of X
Z <- X*(0.1 < X)*(X < 0.9) + 1*(X >= 0.9) # Mixed transform of X
V[1:10] <- NA # Introducing some NAs
W[3:12] <- NA # Introducing some NAs
Y[5:25] <- NA # Introducing some NAs
vector5D <- cbind(V, W, X, Y, Z) # Matrix of 5-variate observations
# Monotone and supremum dependence, ratio and proportion of NAs:
(deparray <- dependence(vector5D, cont = c(1, 2, 3), 30))
# Pearson's correlations:
cor(vector5D, method = "pearson", use = "pairwise.complete.obs")
# Spearman's correlations:
cor(vector5D, method = "spearman", use = "pairwise.complete.obs")
# Kendall's correlations:
cor(vector5D, method = "kendall", use = "pairwise.complete.obs")
pairs(vector5D) # Matrix of pairwise scatterplots

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subcopem

*Bivariate Empirical Subcopula*

**Description**

Calculation of bivariate empirical subcopula matrix, induced partitions, standardized bivariate sample, and dependence measures for a given bivariate sample.

**Usage**

subcopem(mat.xy, display = FALSE)

**Arguments**

- `mat.xy` 2-column matrix with bivariate observations of a random vector \((X, Y)\).
- `display` logical value indicating if graphs and dependence measures should be displayed.

**Details**

Each of the random variables \(X\) and \(Y\) may be of any kind (discrete, continuous, or mixed). NA values are not allowed.

**Value**

A list containing the following components:

- `depMon` monotone standardized supremum distance in \([-1, 1]\).
- `depMonNonSTD` monotone non-standardized supremum distance \([\text{min}, \text{value}, \text{max}]\).
- `depSup` standardized supremum distance in \([0, 1]\).
- `depSupNonSTD` non-standardized supremum distance \([\text{min}, \text{value}, \text{max}]\).
- `matrix` matrix with empirical subcopula values.
- `part1` vector with partition induced by first variable \(X\).
- `part2` vector with partition induced by second variable \(Y\).
sample.size numeric value of sample size.
std.sample 2-column matrix with the standardized bivariate sample.
sample 2-column matrix with the original bivariate sample of \((X, Y)\).

If display = TRUE then the values of depMon, depMonNonSTD, depSup, and depSupNonSTD will be displayed, and the following graphs will be generated: marginal histograms of \(X\) and \(Y\), scatterplots of the original and the standardized bivariate sample, contour and image bivariate graphs of the empirical subcopula.

Note

If both \(X\) and \(Y\) are continuous random variables it is faster and better to use subcopemc.

Author(s)

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References


See Also

subcopemc

Examples

```r
## Example 1: Discrete-discrete Poisson positive dependence
n <- 1000  # sample size
X <- rpois(n, 5)  # Poisson(parameter = 5)
p <- 2  # another parameter
Y <- mapply(rpois, rep(1, n), 1 + p*X)  # creating dependence
XY <- cbind(X, Y)  # 2-column matrix with bivariate sample
cor(XY, method = "pearson")[1, 2]  # Pearson's correlation
cor(XY, method = "spearman")[1, 2]  # Spearman's correlation
cor(XY, method = "kendall")[1, 2]  # Kendall's correlation
SC <- subcopem(XY, display = TRUE)
str(SC)

## Example 2: Continuous-discrete non-monotone dependence
n <- 1000  # sample size
X <- rnorm(n)  # Normal(0,1)
Y <- 2*(X > 1) - 1*(X > -1)  # Discrete({-1, 0, 1})
XY <- cbind(X, Y)  # 2-column matrix with bivariate sample
cor(XY, method = "pearson")[1, 2]  # Pearson's correlation
cor(XY, method = "spearman")[1, 2]  # Spearman's correlation
```
cor(XY, method = "kendall")[1, 2] # Kendall's correlation
SC <- subcopem(XY, display = TRUE)
str(SC)

---

**subcopemc**  
*Bivariate Empirical Subcopula of Given Approximation Order*

**Description**

Calculation of bivariate empirical subcopula matrix of given approximation order, induced partitions, standardized bivariate sample, and dependence measures for a given continuous bivariate sample.

**Usage**

`subcopemc(mat.xy, m = nrow(mat.xy), display = FALSE)`

**Arguments**

- **mat.xy**
  
  2-column matrix with bivariate observations of a continuous random vector \((X, Y)\).

- **m**
  
  integer value of approximation order, where \(m = 2, \ldots, n\) with \(n\) equal to sample size.

- **display**
  
  logical value indicating if graphs and dependence values should be displayed.

**Details**

Both random variables \(X\) and \(Y\) must be continuous, and therefore repeated values in the sample are not expected. If found, jitter will be applied to break ties. NA values are not allowed.

**Value**

A list containing the following components:

- **depMon**
  
  monotone standardized supremum distance in \([-1, 1]\).

- **depMonNonSTD**
  
  monotone non-standardized supremum distance \([min, value, max]\).

- **depSup**
  
  standardized supremum distance in \([0, 1]\).

- **depSupNonSTD**
  
  non-standardized supremum distance \([min, value, max]\).

- **matrix**
  
  matrix with empirical subcopula values.

- **part1**
  
  vector with partition induced by first variable \(X\).

- **part2**
  
  vector with partition induced by second variable \(Y\).

- **sample.size**
  
  numeric value of sample size.

- **order**
  
  numeric value of approximation order.

- **std.sample**
  
  2-column matrix with the standardized bivariate sample.
sample 2-column matrix with the original bivariate sample of \((X, Y)\).

If \texttt{display = TRUE} then the values of \texttt{depMon}, \texttt{depMonNonSTD}, \texttt{depSup}, and \texttt{depSupNonSTD} will be displayed, and the following graphs will be generated: marginal histograms of \(X\) and \(Y\), scatterplots of the original and the standardized bivariate sample, contour and image bivariate graphs of the empirical subcopula.

**Note**

If approximation order \(m > 2000\) calculation may take more than 2 minutes. Usually \(m = 50\) would be enough for an acceptable approximation.

**Author(s)**

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**References**


**See Also**

\texttt{subcopem}

**Examples**

```r
## Example 1: Independent Normal and Gamma

n <- 300  # sample size
X <- rnorm(n)  # Normal(0,1)
Y <- rgamma(n, 2, 3)  # Gamma(2,3)
XY <- cbind(X, Y)  # 2-column matrix with bivariate sample
cor(XY, method = "pearson")[1, 2]  # Pearson's correlation
cor(XY, method = "spearman")[1, 2]  # Spearman's correlation
cor(XY, method = "kendall")[1, 2]  # Kendall's correlation
SC <- subcopemc(XY, display = TRUE)
str(SC)

## Example 2: Non-monotone dependence

n <- 300  # sample size
Theta <- runif(n, 0, 2*pi)  # Uniform circular distribution
X <- cos(Theta)
Y <- sin(Theta)
```

XY <- cbind(X, Y)  # 2-column matrix with bivariate sample
cor(XY, method = "pearson")[1, 2]  # Pearson's correlation
cor(XY, method = "spearman")[1, 2]  # Spearman's correlation
cor(XY, method = "kendall")[1, 2]  # Kendall's correlation
SC <- subcopemc(XY, display = TRUE)
str(SC)
## Approximation of order m = 15
SCm15 <- subcopemc(XY, 15, display = TRUE)
str(SCm15)
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