

Package ‘HMMcopula’

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Type Package

Title Markov Regime Switching Copula Models Estimation and Goodness of Fit

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Author Mamadou Yamar Thioub <mamadou-yamar.thioub@hec.ca>,
Bouchra Nasri <bouchra.nasri@umontreal.ca>,
Romanic Pieugueu <romanic.pieugueu@gerad.ca>, and
Bruno Remillard <bruno.remillard@hec.ca>

Maintainer Mamadou Yamar Thioub <mamadou-yamar.thioub@hec.ca>

Description R functions to estimate and perform goodness of fit test for several Markov regime switching and mixture bivariate copula models. The goodness of fit test is based on a Cramer von Mises statistic and uses the Rosenblatt transform and parametric bootstrap to estimate the p-value. The estimation of the copula parameters are based on the pseudo-maximum likelihood method using pseudo-observations defined as normalized ranks.

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Imports stats

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dilog

Dilogarithm function

Description

This function computes the dilogarithm of a number.

Usage

dilog(x)

Arguments

x a real number

Value

out dilogarithm

EstHMMCop	<i>Estimation of bivariate Markov regime switching bivariate copula model</i>
-----------	---

Description

This function estimates parameters from a bivariate Markov regime switching bivariate copula model

Usage

```
EstHMMCop(y, reg, family, max_iter, eps)
```

Arguments

y	(nx2) data matrix (observations or residuals) that will be transformed to pseudo-observations
reg	number of regimes
family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
max_iter	maximum number of iterations of the EM algorithm
eps	precision (stopping criteria); suggestion 0.0001.

Value

theta	(1 x reg) estimated parameter of the copula according to CRAN copula package (except for Frank copula, where theta = log(theta_R_Package)) for each regime (except for degrees of freedom)
dof	estimated degree of freedom, only for the Student copula
Q	(reg x reg) estimated transition matrix
eta	(n x reg) conditional probabilities of being in regime k at time t given observations up to time t
tau	estimated Kendall tau for each regime
U	(n x 2) matrix of Rosenblatt transforms
cvm	Cramer-von-Mises statistic for goodness-of-fit
W	regime probabilities for the conditional distribution given the past Kendall's tau

Examples

```
Q <- matrix(c(0.8, 0.3, 0.2, 0.7),2,2) ; kendallTau <- c(0.3 ,0.7) ;
data <- SimHMMCop(Q, 'clayton', kendallTau, 10)$SimData;
estimations <- EstHMMCop(data,2,'clayton',10000,0.0001)
```

EstKendallTau	<i>Sample Kendall's tau Estimation</i>
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Description

This function estimates the sample Kendall's tau of a bivariate data matrix

Usage

```
EstKendallTau(X)
```

Arguments

X	(n x 2) matrix
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Value

KendallTau	estimated sample Kendall's tau of the data
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EstMixtureCop	<i>Estimation of bivariate mixture bivariate copula model</i>
---------------	---

Description

This function estimates parameters from a mixture bivariate copula model

Usage

```
EstMixtureCop(y, reg, family, max_iter, eps)
```

Arguments

y	(nx2) data matrix (observations or residuals) that will be transformed to pseudo-observations
reg	number of regimes
family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
max_iter	maximum number of iterations of the EM algorithm
eps	precision (stopping criteria); suggestion 0.0001.

Value

theta	(1 x reg) estimated parameter of the copula according to CRAN copula package (except for Frank copula, where $\theta = \log(\theta_{R_Package})$) for each component (except for degrees of freedom)
dof	estimated degree of freedom, only for the Student copula
Q	(1 x reg) estimated weights vector
eta	(n x reg) conditional probabilities of being in regime k at time t given observations up to time t
tau	estimated Kendall tau for each regime
U	(n x 2) matrix of Rosenblatt transforms
cvm	Cramer-von-Mises statistic for goodness-of-fit

GofHMMCop

*Goodness-of-fit of Markov regime switching bivariate copula model***Description**

This function performs goodness-of-fit test of a Markov regime switching bivariate copula model

Usage

```
GofHMMCop(R, reg, family, max_iter, eps, n_sample, n_cores)
```

Arguments

R	(n x 2) data matrix that will be transformed to pseudo-observations
reg	number of regimes
family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
max_iter	maximum number of iterations of the EM algorithm
eps	precision (stopping criteria); suggestion 0.0001
n_sample	number of bootstrap; suggestion 1000
n_cores	number of cores to use in the parallel computing

Value

pvalue	pvalue (significant when the result is greater than 5)
theta	(1 x reg) estimated parameter of the copula according to CRAN copula package (except for Frank copula, where $\theta = \log(\theta_{R_Package})$) for each regime (except for degrees of freedom)
dof	estimated degree of freedom, only for the Student copula
Q	(reg x reg) estimated transition matrix

eta	(n x reg) conditional probabilities of being in regime k at time t given observations up to time t
tau	estimated Kendall tau for each regime
U	(n x 2) matrix of Rosenblatt transforms
cvm	Cramer-von-Mises statistic for goodness-of-fit
W	regime probabilities for the conditional distribution given the past Kendall's tau

GofMixtureCop	<i>Goodness-of-fit of mixture bivariate copula model</i>
---------------	--

Description

This function performs goodness-of-fit test of a mixture bivariate copula model

Usage

```
GofMixtureCop(R, reg, family, max_iter, eps, n_sample, n_cores)
```

Arguments

R	(nx2) data matrix (observations or residuals) that will be transformed to pseudo-observations
reg	number of regimes
family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
max_iter	maximum number of iterations of the EM algorithm
eps	precision (stopping criteria); suggestion 0.0001
n_sample	number of bootstrap; suggestion 1000
n_cores	number of cores to use in the parallel computing

Value

pvalue	pvalue (significant when the result is greater than 5)
theta	(1 x reg) estimated parameter of the copula according to CRAN copula package (except for Frank copula, where theta = log(theta_R_Package)) for each component (except for degrees of freedom)
dof	estimated degree of freedom, only for the Student copula
Q	(1 x reg) estimated weights vector
eta	(n x reg) conditional probabilities of being in regime k at time t given observations up to time t
tau	estimated Kendall tau for each regime
U	(n x 2) matrix of Rosenblatt transforms
cvm	Cramer-von-Mises statistic for goodness-of-fit

KendallTau	<i>Kendall's tau of a copula</i>
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Description

This function computes the Kendall's tau of a copula family with a unconstrained parameter alpha.

Usage

```
KendallTau(family, alpha)
```

Arguments

family	"gaussian" , "t" , "clayton" , "frank" , "gumbel"
alpha	unconstrained parameters of the copula family

Value

tau	estimated Kendall's tau
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ParamCop	<i>Theta estimation</i>
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Description

This function computes the parameter of the copula according to CRAN copula package (except for Frank copula, where $\theta = \log(\theta_{R_Package})$), corresponding to the unconstrained parameters alpha.

Usage

```
ParamCop(family, alpha)
```

Arguments

family	"gaussian" , "t" , "clayton" , "frank" , "gumbel"
alpha	unconstrained parameters of the copula family

Value

theta	matlab parameters
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ParamTau	<i>Alpha estimation</i>
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Description

This function computes the unconstrained parameter alpha for given Kendall's tau value

Usage

```
ParamTau(family, tau)
```

Arguments

family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
tau	Kendall's tau of the copula family

Value

alpha	estimated unconstrained parameter
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RosenblattClayton	<i>Rosenblatt transform for Clayton copula</i>
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Description

This function computes the Rosenblatt transform for the Clayton copula

Usage

```
RosenblattClayton(u, theta)
```

Arguments

u	(n x d) matrix of pseudos-observations (normalized ranks)
theta	parameter of the Clayton copula

Value

R	Rosenblatt transform
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RosenblattFrank	<i>Rosenblatt transform for Frank copula</i>
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Description

This function computes the Rosenblatt transform for the Frank copula

Usage

```
RosenblattFrank(U, theta)
```

Arguments

U	(n x d) matrix of pseudos-observations (normalized ranks)
theta	parameter of the Frank copula

Value

R	Rosenblatt transform
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RosenblattGaussian	<i>Rosenblatt transform for Gaussian copula</i>
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Description

This function computes the Rosenblatt transform for the Gaussian copula

Usage

```
RosenblattGaussian(u, rho)
```

Arguments

u	(n x d) matrix of pseudos-observations (normalized ranks)
rho	(d x d) correlation matrix, or the correlation coefficient (if, d = 2)

Value

R	Rosenblatt transform
---	----------------------

RosenblattGumbel *Rosenblatt transform for Gumbel copula*

Description

This function computes the Rosenblatt transform for the Gumbel copula

Usage

```
RosenblattGumbel(U, theta)
```

Arguments

U (n x d) matrix of pseudos-observations (normalized ranks)
theta parameter of the Gumbel copula

Value

R Rosenblatt transform

RosenblattStudent *Rosenblatt transform for Student copula*

Description

This function computes the Rosenblatt transform for the Student copula

Usage

```
RosenblattStudent(u, rho, nu)
```

Arguments

u (n x d) matrix of pseudos-observations (normalized ranks)
rho (d x d) correlation matrix
nu degrees of freedom

Value

R Rosenblatt transform

SimHMMCop	<i>Simulation of bivariate Markov regime switching copula model</i>
-----------	---

Description

This function simulates observation from a bivariate Markov regime switching copula model

Usage

```
SimHMMCop(Q, family, KendallTau, n, DoF)
```

Arguments

Q	Transition probability matrix (d x d);
family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
KendallTau	Kendall's rank correlation
n	number of simulated vectors
DoF	degree of freedom only for the Student copula

Value

SimData	Simulated Data
MC	Markov chain regimes
alpha	parameters alpha

Examples

```
Q <- matrix(c(0.8, 0.3, 0.2, 0.7),2,2) ; kendallTau <- c(0.3 ,0.7) ;
simulations <- SimHMMCop(Q, 'gumbel', kendallTau, 300)
```

SimMarkovChain	<i>Markov chain simulation</i>
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Description

This function generates a Markov chain $X(1), \dots, X(n)$ with transition matrix Q , starting from a state η_0 or the uniform distribution on $1, \dots, r$

Usage

```
SimMarkovChain(Q, n, eta0)
```

Arguments

Q	Transition probability matrix (d x d)
n	number of simulated vectors
eta0	variable eta

SimMixtureCop	<i>Simulation of bivariate mixture copula model</i>
---------------	---

Description

This function simulates observation from a bivariate mixture copula model

Usage

```
SimMixtureCop(Q, family, KendallTau, n, DoF)
```

Arguments

Q	Weights vector (1 x component);
family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
KendallTau	Kendall's rank correlation
n	number of simulated vectors
DoF	vector of degree of freedom only for the Student copula

Value

SimData	Simulated Data
MC	Markov chain regimes
alpha	parameters alpha

Examples

```
Q <- matrix(c(0.8, 0.2),1,2) ; kendallTau <- c(0.3 ,0.7) ;
simulations <- SimMixtureCop(Q, 'gaussian', kendallTau, 300)
```

SnB	<i>Cramer-von Mises statistic SnB for GOF based on the Rosenblatt transform</i>
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Description

This function computes the Cramer-von Mises statistic SnB for GOF based on the Rosenblatt transform

Usage

```
SnB(E)
```

Arguments

E	(n x d) matrix of pseudos-observations (normalized ranks)
---	---

Value

Sn	Cramer-von Mises statistic
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Tau2Rho	<i>Spearman's rho</i>
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Description

This function estimates the Spearman's rho corresponding to a constrained (matlab) parameter theta for a copula family.

Usage

```
Tau2Rho(family, theta)
```

Arguments

family	'gaussian', 't', 'clayton', 'frank', 'gumbel'
theta	parameter of the copula according to CRAN copula package (except for Frank copula, where theta = log(theta_R_Package))

Value

rho	estimated Spearman's rho
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