

Package ‘Copula.Markov’

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Type Package

Title Copula-Based Estimation and Statistical Process Control for
Serially Correlated Time Series

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Description Estimation and statistical process control are performed under
copula-based time-series models.

Available are statistical methods in Long and Emura (2014 JCSA),
Emura et al. (2017 Commun Stat-Simul) <DOI:10.1080/03610918.2015.1073303>,
Huang and Emura (2021 Commun Stat-Simul) <DOI:10.1080/03610918.2019.1602647>,
Lin et al. (2021 Comm Stat-Simul) <DOI:10.1080/03610918.2019.1652318>,
Sun et al. (2020 JSS Series in Statistics)<DOI:10.1007/978-981-15-4998-4>,
and Huang and Emura (2021, in revision).

License GPL-2

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Copula.Markov-package *Copula-Based Estimation and Statistical Process Control for Serially Correlated Time Series*

Description

Copulas are applied to model a Markov dependence for serially correlated time series. The Clayton and Joe copulas are available to specify the dependence structure. The normal and binomial distributions are available for the marginal model. Maximum likelihood estimation is implemented for estimating parameters, and a Shewhart control chart is drawn for performing statistical process control.

Details

Package:	Copula.Markov
Type:	Package
Version:	2.9
Date:	2021-11-29
License:	GPL-2

Author(s)

Emura T, Huang XW, Chen WR, Long TH, Sun LH. Maintainer: Takeshi Emura <takeshiemura@gmail.com>

References

- Chen W (2018) Copula-based Markov chain model with binomial data, NCU Library
- Huang XW, Emura T (2021-), Computational methods for a copula-based Markov chain model with a binomial time series, in review
- Emura T, Long TH, Sun LH (2017), R routines for performing estimation and statistical process control under copula-based time series models, Communications in Statistics - Simulation and Computation, 46(4):3067-87
- Long TH and Emura T (2014), A control chart using copula-based Markov chain models, Journal of the Chinese Statistical Association, 52(4):466-96

Lin WC, Emura T, Sun LH (2021), Estimation under copula-based Markov normal mixture models for serially correlated data, Communications in Statistics - Simulation and Computation, 50(12):4483-515

Huang XW, Emura T (2021), Model diagnostic procedures for copula-based Markov chain models for statistical process control, Communications in Statistics - Simulation and Computation, doi: 50(8):2345-67

Clayton.Markov.DATA *Generating Time Series Data Under a Copula-Based Markov Chain Model with the Clayton Copula*

Description

Time-series data are generated under a copula-based Markov chain model with the Clayton copula. See Long et al. (2014) and Emura et al. (2017) for the details.

Usage

```
Clayton.Markov.DATA(n, mu, sigma, alpha)
```

Arguments

n	sample size
mu	mean
sigma	standard deviation
alpha	association parameter

Details

-1<alpha<0 for negative association; alpha>0 for positive association

Value

Time series data of size n.

Author(s)

Takeshi Emura

References

Emura T, Long TH, Sun LH (2017), R routines for performing estimation and statistical process control under copula-based time series models, Communications in Statistics - Simulation and Computation, 46 (4): 3067-87

Long TH and Emura T (2014), A control chart using copula-based Markov chain models, Journal of the Chinese Statistical Association 52 (No.4): 466-96

Examples

```
set.seed(1)
Y=Clayton.Markov.DATA(n=1000, mu=0, sigma=1, alpha=8)
Clayton.Markov.MLE(Y, plot=TRUE)
```

Clayton.Markov.DATA.binom

Generating Time Series Data Under a Copula-Based Markov Chain Model with the Clayton Copula and Binomial Margin.

Description

Time-series data are generated under a copula-based Markov chain model with the Clayton copula and binomial margin.

Usage

```
Clayton.Markov.DATA.binom(n, size, prob, alpha)
```

Arguments

n	number of observations
size	number of binomial trials
prob	binomial probability; $0 < p < 1$
alpha	association parameter

Details

$-1 < \text{alpha} < 0$ for negative association; $\text{alpha} > 0$ for positive association

Value

Time series data of size n (this is different from the number of binomial trials = "size").

Author(s)

Huang XW, Chen W, Emura T

References

- Chen W (2018) Copula-based Markov chain model with binomial data, NCU Library
- Huang XW, Emura T (2021-), Computational methods for a copula-based Markov chain model with a binomial time series, in review

Examples

```
size=50
prob=0.5
alpha=2
set.seed(1)
Y=Clayton.Markov.DATA.binom(n=500,size,prob,alpha)
### sample mean and SD ####
mean(Y)
sd(Y)
### true mean and SD ####
size*prob
sqrt(size*prob*(1-prob))
```

Clayton.Markov.GOF *A goodness-of-fit test for the marginal normal distribution.*

Description

Perform a parametric bootstrap test based on the Cramer-von Mises and Kolmogorov-Smirnov statistics as proposed by Huang and Emura (2019).

Usage

```
Clayton.Markov.GOF(Y, k = 3, D = 1, B = 200, GOF.plot=FALSE, method = "Newton")
```

Arguments

Y	vector of datasets
k	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
D	diameter for U(-D, D) used in randomized Newton-Raphson
B	the number of Bootstrap replications
GOF.plot	if TRUE, show the model diagnostic plots for B bootstrap replications
method	Newton-Raphson method or nlm can be chosen

Value

CM	The Cramer-von Mises statistic and its P-value
KS	The Kolmogorov-Smirnov statistic and its P-value
CM.boot	Bootstrap values of the Cramer-von Mises statistics
KS.boot	Bootstrap values of the Kolmogorov-Smirnov statistics

Author(s)

Takeshi Emura

References

- Emura T, Long TH, Sun LH (2017), R routines for performing estimation and statistical process control under copula-based time series models, Communications in Statistics - Simulation and Computation, 46 (4): 3067-87
- Long TH and Emura T (2014), A control chart using copula-based Markov chain models, Journal of the Chinese Statistical Association 52 (No.4): 466-96
- Huang XW, Emura T (2021), Model diagnostic procedures for copula-based Markov chain models for statistical process control, Communications in Statistics - Simulation and Computation, doi: 50(8):2345-67

Examples

```
set.seed(1)
Y=Clayton.Markov.DATA(n=1000, mu=0, sigma=1, alpha=2)
Clayton.Markov.GOF(Y, B=5, GOF.plot=TRUE)
```

Clayton.Markov.GOF.binom

A goodness-of-fit test for the marginal binomial distribution.

Description

Perform a parametric bootstrap test based on the Cramer-von Mises and Kolmogorov-Smirnov statistics as proposed by Huang and Emura (2019) and Huang et al. (2019-).

Usage

```
Clayton.Markov.GOF.binom(Y, k = 3, size, B = 200, GOF.plot=FALSE, method = "Newton")
```

Arguments

Y	vector of datasets
k	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
size	number of binomial trials
B	the number of Bootstrap replications
GOF.plot	if TRUE, show the model diagnostic plots for B bootstrap replications
method	Newton-Raphson method or nlm can be chosen

Value

CM	The Cramer-von Mises statistic and its P-value
KS	The Kolmogorov-Smirnov statistic and its P-value
CM.boot	Bootstrap values of the Cramer-von Mises statistics
KS.boot	Bootstrap values of the Kolmogorov-Smirnov statistics

Author(s)

Huang XW, Emura T

References

Huang XW, Emura T (2021), Model diagnostic procedures for copula-based Markov chain models for statistical process control, Communications in Statistics - Simulation and Computation, doi: 50(8):2345-67

Huang XW, Emura T (2021-), Computational methods for a copula-based Markov chain model with a binomial time series, in review

Examples

```
size=50
prob=0.5
alpha=2
set.seed(1)
Y=Clayton.Markov.DATA.binom(n=500,size,prob,alpha)
Clayton.Markov.GOF.binom(Y,size=size,B=5,k=3,GOF.plot=TRUE) ## B=5 to save time
```

Clayton.Markov.MLE

Maximum Likelihood Estimation and Statistical Process Control Under the Clayton Copula

Description

The maximum likelihood estimates are produced and the Shewhart control chart is drawn with k-sigma control limits (e.g., 3-sigma). The dependence model follows the Clayton copula and the marginal (stationary) distribution follows the normal distribution.

Usage

```
Clayton.Markov.MLE(Y, k = 3, D = 1, plot = TRUE, GOF=FALSE, method = "nlm")
```

Arguments

Y	vector of datasets
k	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
D	diameter for U(-D, D) used in randomized Newton-Raphson
plot	show the control chart if TRUE
GOF	show the model diagnostic plot if TRUE
method	apply "nlm" or "Newton" method

Value

<code>mu</code>	estimate, SE, and 95 percent CI
<code>sigma</code>	estimate, SE, and 95 percent CI
<code>alpha</code>	estimate, SE, and 95 percent CI
<code>Control_Limit</code>	Center = mu, LCL = mu - k*sigma, UCL = mu + k*sigma
<code>out_of_control</code>	IDs for out-of-control points
<code>Gradient</code>	gradients (must be zero)
<code>Hessian</code>	Hessian matrix
<code>Eigenvalue_Hessian</code>	Eigenvalues for the Hessian matrix
<code>KS.test</code>	KS statistics
<code>CM.test</code>	CM statistics
<code>log.likelihood</code>	Log-likelihood value for the estimation

Author(s)

Long TH, Huang XW and Emura T

References

- Emura T, Long TH, Sun LH (2017), R routines for performing estimation and statistical process control under copula-based time series models, Communications in Statistics - Simulation and Computation, 46 (4): 3067-87
- Long TH and Emura T (2014), A control chart using copula-based Markov chain models, Journal of the Chinese Statistical Association 52 (No.4): 466-96

Examples

```
set.seed(1)
Y=Clayton.Markov.DATA(n=1000,mu=0,sigma=1,alpha=2)
Clayton.Markov.MLE(Y,plot=TRUE)
```

Clayton.Markov.MLE.binom

Maximum Likelihood Estimation and Statistical Process Control Under the Clayton Copula

Description

The maximum likelihood estimates are produced and the Shewhart control chart is drawn with k-sigma control limits (e.g., 3-sigma). The dependence model follows the Clayton copula and the marginal (stationary) distribution follows the normal distribution.

Usage

```
Clayton.Markov.MLE.binom(Y, size, k = 3, method="nlm", plot = TRUE, GOF=FALSE)
```

Arguments

<code>Y</code>	vector of observations
<code>size</code>	numbe of binomial trials
<code>method</code>	nlm or Newton
<code>k</code>	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
<code>plot</code>	show the control chart if TRUE
<code>GOF</code>	show the model diagnostic plot if TRUE

Value

<code>p</code>	estimate, SE, and 95 percent CI
<code>alpha</code>	estimate, SE, and 95 percent CI
<code>Control_Limit</code>	Center = $n*p$, LCL = $\mu - k\sigma$, UCL = $\mu + k\sigma$
<code>out_of_control</code>	IDs for out-of-control points
<code>Gradient</code>	gradients (must be zero)
<code>Hessian</code>	Hessian matrix
<code>Eigenvalue_Hessian</code>	Eigenvalues for the Hessian matrix
<code>KS.test</code>	KS statistics
<code>CM.test</code>	CM statistics
<code>log_likelihood</code>	Log-likelihood value for the estimation

Author(s)

Huang XW, Emura T

References

- Chen W (2018) Copula-based Markov chain model with binomial data, NCU Library
 Huang XW, Emura T (2021-), Computational methods for a copula-based Markov chain model with a binomial time series, in review

Examples

```
size=50
prob=0.5
alpha=2
set.seed(1)
Y=Clayton.Markov.DATA.binom(n=500,size,prob,alpha)
Clayton.Markov.MLE.binom(Y,size=size,k=3,plot=TRUE)
```

Clayton.Markov2.DATA *Generating Time Series Data Under a Copula-Based 2nd-order Markov Chain Model with the Clayton Copula*

Description

Time-series data are generated under a copula-based 2nd order Markov chain model with the Clayton copula.

Usage

```
Clayton.Markov2.DATA(n, mu, sigma, alpha)
```

Arguments

n	sample size
mu	mean
sigma	standard deviation
alpha	association parameter

Details

-1<alpha<0 for negative association; alpha>0 for positive association

Value

Time series data of size n

Author(s)

Xinwei Huang and Takeshi Emura

References

Huang XW, Emura T (2021), Model diagnostic procedures for copula-based Markov chain models for statistical process control, Communications in Statistics - Simulation and Computation, doi: 50(8):2345-67

Examples

```
Clayton.Markov2.DATA(n = 100, mu = 0, sigma = 1, alpha = 2)
```

`Clayton.Markov2.MLE` *Maximum Likelihood Estimation and Statistical Process Control Under the Clayton Copula with a 2nd order Markov chain.*

Description

The maximum likelihood estimates are produced and the Shewhart control chart is drawn with k-sigma control limits (e.g., 3-sigma). The dependence model follows the Clayton copula and the marginal (stationary) distribution follows the normal distribution. The model diagnostic plot is also given (by the option "GOF=TRUE"). See Huang and Emura (2019) for the methodological details.

Usage

```
Clayton.Markov2.MLE(Y, k = 3, D = 1, plot = TRUE, GOF=FALSE)
```

Arguments

<code>Y</code>	vector of datasets
<code>k</code>	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
<code>D</code>	diameter for U(-D, D) used in randomized Newton-Raphson
<code>plot</code>	show the control chart if TRUE
<code>GOF</code>	show the model diagnostic plot if TRUE

Value

<code>mu</code>	estimate, SE, and 95 percent CI
<code>sigma</code>	estimate, SE, and 95 percent CI
<code>alpha</code>	estimate, SE, and 95 percent CI
<code>Control_Limit</code>	Center = mu, LCL = mu - k*sigma, UCL = mu + k*sigma
<code>out_of_control</code>	IDs for out-of-control points
<code>Gradient</code>	gradients (must be zero)
<code>Hessian</code>	Hessian matrix
<code>Eigenvalue_Hessian</code>	Eigenvalues for the Hessian matrix
<code>KS.test</code>	KS statistics
<code>CM.test</code>	CM statistics
<code>log.likelihood</code>	Log-likelihood value for the estimation

Author(s)

Xinwei Huang and Takeshi Emura

References

Huang XW, Emura T (2021), Model diagnostic procedures for copula-based Markov chain models for statistical process control, Communications in Statistics - Simulation and Computation, doi: 50(8):2345-67

Examples

```

Y = c(0.265, 0.256, 0.261, 0.261, 0.260, 0.257, 0.258, 0.263, 0.254, 0.254,
     0.258, 0.256, 0.256, 0.265, 0.270, 0.267, 0.270, 0.267, 0.266, 0.271,
     0.270, 0.264, 0.261, 0.264, 0.266, 0.264, 0.269, 0.268, 0.264, 0.262,
     0.257, 0.255, 0.255, 0.253, 0.251, 0.254, 0.255)
Clayton.Markov2.MLE(Y, k = 1, D = 1, plot = TRUE)

Y=Clayton.Markov2.DATA(n=1000,mu=0,sigma=1,alpha=8)
Clayton.Markov2.MLE(Y, plot=TRUE)

```

Clayton.MixNormal.Markov.MLE

Maximum Likelihood Estimation using Newton-Raphson Method Under the Clayton Copula and the Mix-Normal distribution

Description

The maximum likelihood estimates are produced. The dependence model follows the Clayton copula and the marginal distribution follows the Mix-Normal distribution.

Usage

```
Clayton.MixNormal.Markov.MLE(y)
```

Arguments

y	vector of datasets
---	--------------------

Value

alpha	estimate, SE, and 95 percent CI
mu1	estimate, SE, and 95 percent CI
mu2	estimate, SE, and 95 percent CI
sigma1	estimate, SE, and 95 percent CI
sigma2	estimate, SE, and 95 percent CI
p	estimate, SE, and 95 percent CI
Gradient	gradients (must be zero)
Hessian	Hessian matrix
Eigenvalue_Hessian	Eigenvalues for the Hessian matrix
log.likelihood	Log-likelihood value for the estimation

Author(s)

Sun LH, Huang XW

References

Lin WC, Emura T, Sun LH (2021), Estimation under copula-based Markov normal mixture models for serially correlated data, Communications in Statistics - Simulation and Computation, 50(12):4483-515

Examples

```
data(DowJones)
Y=as.vector(DowJones$log_return)
Clayton.MixNormal.Markov.MLE(y=Y)
```

DowJones

Dow Jones Industrial Average

Description

The log return of weekly stock price of Dow Jones Industrial Average from 2008/1/1 to 2012/1/1.

Usage

```
data("DowJones")
```

Format

A data frame with 754 observations on the following 1 variables.

`log_return` a numeric vector

References

Lin WC, Emura T, Sun LH (2021), Estimation under copula-based Markov normal mixture models for serially correlated data, Communications in Statistics - Simulation and Computation, 50(12):4483-515

Examples

```
data(DowJones)
DowJones
```

Joe.Markov.DATA

Generating Time Series Data Under a Copula-Based Markov Chain Model with the Joe Copula

Description

Time-series data are generated under a copula-based Markov chain model with the Joe copula.

Usage

```
Joe.Markov.DATA(n, mu, sigma, alpha)
```

Arguments

n	sample size
mu	mean
sigma	standard deviation
alpha	association parameter

Details

alpha>=1 for positive association

Value

Time series data of size n

Author(s)

Takeshi Emura

References

Emura T, Long TH, Sun LH (2017), R routines for performing estimation and statistical process control under copula-based time series models, Communications in Statistics - Simulation and Computation, 46 (4): 3067-87

Long TS and Emura T (2014), A control chart using copula-based Markov chain models, Journal of the Chinese Statistical Association 52 (No.4): 466-96

Examples

```
n=1000
alpha=2.856 ### Kendall's tau =0.5 ####
mu=2
sigma=1
Y=Joe.Markov.DATA(n,mu,sigma,alpha)
mean(Y)
```

```

sd(Y)
cor(Y[-1], Y[-n], method="kendall")

Joe.Markov.MLE(Y, k=2)

```

Joe.Markov.DATA.binom *Generating Time Series Data Under a Copula-Based Markov Chain Model with the Joe Copula and Binomial Margin.*

Description

Time-series data are generated under a copula-based Markov chain model with the Joe copula and binomial margin.

Usage

```
Joe.Markov.DATA.binom(n, size, prob, alpha)
```

Arguments

n	number of observations
size	number of binomial trials
prob	binomial probability; 0<p<1
alpha	association parameter

Details

alpha>=1 for positive association

Value

time series data

Author(s)

Huang X, Emura T

References

- Chen W (2018) Copula-based Markov chain model with binomial data, NCU Library
Huang XW, Emura T (2021-), Computational methods for a copula-based Markov chain model with a binomial time series, in review

Examples

```

size=50
prob=0.5
alpha=2
set.seed(1)
Y=Joe.Markov.DATA.binom(n=500,size,prob,alpha)
### sample mean and SD ####
mean(Y)
sd(Y)
### true mean and SD ####
size*prob
sqrt(size*prob*(1-prob))

```

Joe.Markov.GOF.binom *A goodness-of-fit test for the marginal binomial distribution.*

Description

Perform a parametric bootstrap test based on the Cramer-von Mises and Kolmogorov-Smirnov statistics as proposed by Huang and Emura (2019) and Huang et al. (2019-).

Usage

```
Joe.Markov.GOF.binom(Y, k = 3, size, B = 200,GOF.plot=FALSE)
```

Arguments

Y	vector of datasets
k	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
size	number of binomial trials
B	the number of Bootstrap replications
GOF.plot	if TRUE, show the model diagnostic plots for B bootstrap replications

Value

CM	The Cramer-von Mises statistic and its P-value
KS	The Kolmogorov-Smirnov statistic and its P-value
CM.boot	Bootstrap values of the Cramer-von Mises statistics
KS.boot	Bootstrap values of the Kolmogorov-Smirnov statistics

Author(s)

Huang XW, Emura T

References

- Huang XW, Emura T (2021), Model diagnostic procedures for copula-based Markov chain models for statistical process control, Communications in Statistics - Simulation and Computation, doi: 50(8):2345-67
- Huang XW, Emura T (2021-), Computational methods for a copula-based Markov chain model with a binomial time series, in review

Examples

```
size=50
prob=0.5
alpha=2
set.seed(1)
Y=Joe.Markov.DATA.binom(n=500,size,prob,alpha)
Joe.Markov.GOF.binom(Y,size=size,B=5,k=3,GOF.plot=TRUE) ## B=5 to save time
```

Joe.Markov.MLE

Maximum Likelihood Estimation and Statistical Process Control Under the Joe Copula

Description

The maximum likelihood estimates are produced and the Shewhart control chart is drawn with k-sigma control limits (e.g., 3-sigma). The dependence model follows the Joe copula and the marginal (stationary) distribution follows the normal distribution.

Usage

```
Joe.Markov.MLE(Y, k = 3, D = 1, plot = TRUE, GOF=FALSE,method = "nlm")
```

Arguments

Y	vector of datasets
k	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
D	diameter for U(-D, D) used in randomized Newton-Raphson
plot	show the control chart if TRUE
GOF	show the model diagnostic plot if TRUE
method	apply "nlm" or "Newton" method

Value

<code>mu</code>	estimate, SE, and 95 percent CI
<code>sigma</code>	estimate, SE, and 95 percent CI
<code>alpha</code>	estimate, SE, and 95 percent CI
<code>Control_Limit</code>	Center = mu, LCL = mu - k*sigma, UCL = mu + k*sigma
<code>out_of_control</code>	IDs for out-of-control points
<code>Gradient</code>	gradients (must be zero)
<code>Hessian</code>	Hessian matrix
<code>Eigenvalue_Hessian</code>	Eigenvalues for the Hessian matrix
<code>KS.test</code>	KS statistics
<code>CM.test</code>	CM statistics
<code>log.likelihood</code>	Log-likelihood value for the estimation

Author(s)

Long TH, Huang XW and Takeshi Emura

References

- Emura T, Long TH, Sun LH (2017), R routines for performing estimation and statistical process control under copula-based time series models, Communications in Statistics - Simulation and Computation, 46 (4): 3067-87
- Long TH and Emura T (2014), A control chart using copula-based Markov chain models, Journal of the Chinese Statistical Association 52 (No.4): 466-96

Examples

```

n=1000
alpha=2.856 ### Kendall's tau =0.5 ####
mu=2
sigma=1
Y=Joe.Markov.DATA(n,mu,sigma,alpha)
mean(Y)
sd(Y)
cor(Y[-1],Y[-n],method="kendall")

Joe.Markov.MLE(Y,k=2)

```

Joe.Markov.MLE.binom *Maximum Likelihood Estimation and Statistical Process Control Under the Joe Copula*

Description

The maximum likelihood estimates are produced and the Shewhart control chart is drawn with k-sigma control limits (e.g., 3-sigma). The dependence model follows the Joe copula and the marginal (stationary) distribution follows the binomial distribution.

Usage

```
Joe.Markov.MLE.binom(Y, size, k = 3, plot = TRUE, GOF=FALSE)
```

Arguments

<code>Y</code>	vector of observations
<code>size</code>	number of binomial trials
<code>k</code>	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
<code>plot</code>	show the control chart if TRUE
<code>GOF</code>	show the model diagnostic plot if TRUE

Value

<code>p</code>	estimate, SE, and 95 percent CI
<code>alpha</code>	estimate, SE, and 95 percent CI
<code>Control_Limit</code>	Center = $n*p$, LCL = $\mu - k\sigma$, UCL = $\mu + k\sigma$
<code>out_of_control</code>	IDs for out-of-control points
<code>Gradient</code>	gradients (must be zero)
<code>Hessian</code>	Hessian matrix
<code>Eigenvalue_Hessian</code>	Eigenvalues for the Hessian matrix
<code>KS.test</code>	KS statistics
<code>CM.test</code>	CM statistics
<code>log_likelihood</code>	Log-likelihood value for the estimation

Author(s)

Huang XW, Emura T

References

- Chen W (2018) Copula-based Markov chain model with binomial data, NCU Library
Huang XW, Emura T (2021+), Computational methods for a copula-based Markov chain model with a binomial time series, under review

Examples

```
size=50
prob=0.5
alpha=2
set.seed(1)
Y=Joe.Markov.DATA.binom(n=500,size,prob,alpha)
Joe.Markov.MLE.binom(Y,size=size,k=3,plot=TRUE)
```

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