# Package 'Copula.Markov'

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<b>Title</b> 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).</doi:10.1007></doi:10.1080></doi:10.1080></doi:10.1080>
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# 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 implmented for estimating parameters, and a Shewhart control chart is drawn for performing statistical process control.

#### **Details**

Package: Copula.Markov
Type: Package
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## Author(s)

Emura T, Huang XW, Chen WR, Long TH, Sun LH. Maintainer: Takeshi Emura <takeshi emura@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

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<alpha<0 for negative association; 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

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## **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**

Υ	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
В	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**

Υ	vector of datasets
k	constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
size	number of binomial trials
В	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

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#### 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**

Υ	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

mu estimate, SE, and 95 percent CI sigma estimate, SE, and 95 percent CI alpha estimate, SE, and 95 percent CI

Control\_Limit Center = mu, LCL = mu - k\*sigma, UCL = mu + k\*sigma

out\_of\_control

IDs for out-of-control points

Gradient gradients (must be zero)

Hessian Hessian matrix

Eigenvalue\_Hessian

Eigenvalues for the Hessian matrix

KS.test KS statistics
CM.test CM statistics

log.likelihood

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**

Y vector of observations size numbe of binomial trials

method nlm or Newton

k constant determining the length between LCL and UCL (k=3 corresponds to

3-sigma limit)

plot show the control chart if TRUE

GOF show the model diagnostic plot if TRUE

#### Value

p estimate, SE, and 95 percent CI alpha estimate, SE, and 95 percent CI

Control\_Limit Center = n\*p, LCL = mu - k\*sigma, UCL = mu + k\*sigma

out\_of\_control

IDs for out-of-control points

Gradient gradients (must be zero)

Hessian Hessian matrix

Eigenvalue\_Hessian

Eigenvalues for the Hessian matrix

KS.test KS statistics
CM.test CM statistics

log\_likelihood

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

```
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

```
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

Υ	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

## Value

mu	estimate, SE, and 95 percent CI
sigma	estimate, SE, and 95 percent CI
alpha	estimate, SE, and 95 percent CI
Control_Limit	Center = $mu$ , $LCL = mu - k*sigma$ , $UCL = mu + k*sigma$
out_of_control	
	IDs for out-of-control points
Gradient	gradients (must be zero)
Hessian	Hessian matrix
Eigenvalue_Hessian	
	Eigenvalues for the Hessian matrix
KS.test	KS statistics
CM.test	CM statistics
log.likelihood	
	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, 0.254, 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

```
estimate, SE, and 95 percent CI
alpha
mu1
                  estimate, SE, and 95 percent CI
mu2
                  estimate, SE, and 95 percent CI
                  estimate, SE, and 95 percent CI
sigma1
                  estimate, SE, and 95 percent CI
sigma2
                  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

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#### 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

```
data(DowJones)
DowJones
```

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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

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

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```
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< td=""></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

Joe.Markov.GOF.binom

## **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**

Υ	vector of datasets
k	constant determining the length between LCL and UCL ( $k=3$ corresponds to 3-sigma limit)
size	number of binomial trials
В	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

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#### 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**

Υ	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

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#### Value

estimate, SE, and 95 percent CI mu estimate, SE, and 95 percent CI sigma alpha estimate, SE, and 95 percent CI Control\_Limit Center = mu, LCL = mu - k\*sigma, UCL = mu + k\*sigmaout\_of\_control IDs for out-of-control points Gradient gradients (must be zero) Hessian Hessian matrix Eigenvalue\_Hessian Eigenvalues for the Hessian matrix KS.test KS statistics CM.test CM statistics log.likelihood

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

```
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

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**

y vector of observations
size number of binomial trials
k constant determining the length between LCL and UCL (k=3 corresponds to 3-sigma limit)
plot show the control chart if TRUE

GOF show the model diagnostic plot if TRUE

#### Value

IDs for out-of-control points

-

Gradient gradients (must be zero)

Hessian Hessian matrix

Eigenvalue\_Hessian

Eigenvalues for the Hessian matrix

KS.test KS statistics
CM.test CM statistics
log\_likelihood

Log-likelihood value for the estimation

## Author(s)

Huang XW, Emura T

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## 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

```
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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