## Package 'SimTools'

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## Description

Boxplots with simultaenous error bars around all quantiles for iid data.

## Usage

boxplot.Siid

#### **Arguments**

x : a 'Siid' class object

... : arguments sent to boxplot

alpha : confidence level of simultaneous confidence intervals

thresh : numeric typically less than .005 for the accuracy of the simulteaneous proce-

dure

mean.col : color for the mean confidence interval

quan.col : color for the quantile confidence intervals

opaq : opacity of mean.col and quan.col. A value of 0 is transparent and 1 is

completely opaque.

range : as defined for base boxplot
width : as defined for base boxplot
varwidth : as defined for base boxplot
outline : as defined for base boxplot

plot : logical indicating whether the plot is to be constructed

border : as defined for base boxplot

col : as defined for base boxplot

ann : as defined for base boxplot

horizontal : as defined for base boxplot

add : as defined for base boxplot

#### Value

returns the base boxplot with simultaneous confidence intervals around all quantiles

#### References

Robertson, N., Flegal, J. M., Vats, D., and Jones, G. L., "Assessing and Visualizing Simultaneous Simulation Error", Journal of Computational and Graphical Statistics, 2020.

#### **Examples**

```
# Generating iid data
chain <- matrix(rnorm(3*1e3), nrow = 1e3, ncol = 3)
siid.obj <- Siid(chain)
boxplot(chain)</pre>
```

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| plot.Smcmc Plot Smcmc |
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#### **Description**

Density plots with simultaenous error bars around means and quantiles for MCMC data. The error bars account for the correlated nature of the process.

#### Usage

#### **Arguments**

x : a 'Smcmc' class object
Q : vector of quantiles

alpha : confidence level of simultaneous confidence intervals

thresh : numeric typically less than .005 for the accuracy of the simulteaneous proce-

dure

iid : logical argument for constructing density plot for iid samples. Defaults to

FALSE

plot : logical argument for is plots are to be returned
mean : logical argument whether the mean is to be plotted

border : whether a border is required for the simultaneous confidence intervals

mean.col : color for the mean confidence interval quan.col : color for the quantile confidence intervals

opaq : opacity of mean.col and quan.col. A value of 0 is transparent and 1 is

completely opaque.

auto.layout : logical argument for an automatic layout of plots

ask : activating interactive plots

... : arguments passed on to the density plot in base R

## Value

returns a plot of the univariate density estimates with simultaneous confidence intervals wherever asked. If plot == FALSE a list of estimates and simultaneous confidence intervals.

#### References

Robertson, N., Flegal, J. M., Vats, D., and Jones, G. L., "Assessing and Visualizing Simultaneous Simulation Error", Journal of Computational and Graphical Statistics, 2020.

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#### **Examples**

```
# Producing Markov chain
chain <- numeric(length = 1e3)
chain[1] <- 0
err <- rnorm(1e3)
for(i in 2:1e3)
{
   chain[i] <- .3*chain[i-1] + err[i]
}
chain <- Smcmc(chain)
plot(chain)</pre>
```

Siid

Siid class

## Description

Class for independent and identically distributed (iid) samples

## Usage

```
Siid(data, varnames = colnames(data))
```

## **Arguments**

data : an iid output matrix with nsim rows and p columns

varnames : a character string equal to the number of columns in data

## Value

an Siid class object

## **Examples**

```
# Generating iid data
chain <- matrix(rnorm(3*1e3), nrow = 1e3, ncol = 3)
siid.obj <- Siid(chain)</pre>
```

 $\operatorname{Smcmc}$ 

Smcmc class

## Description

Smcmc class for simulated data using Markov chain Monte Carlo

## Usage

```
Smcmc(data, batch.size = FALSE, varnames = colnames(data))
```

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#### **Arguments**

data : an MCMC output matrix with nsim rows and p columns

batch.size : logical vector, if true, calculates the batch size appropriate for this Markov

chain. Setting to TRUE saves time in future steps.

varnames : a character string equal to the number of columns in data

#### Value

an Smcmc class object

## **Examples**

```
# Producing Markov chain
chain <- numeric(length = 1e3)
chain[1] <- 0
err <- rnorm(1e3)
for(i in 2:1e3)
{
   chain[i] <- .3*chain[i-1] + err[i]
}
smcmc.obj <- Smcmc(chain)</pre>
```

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