Package 'markovchain'

July 2, 2014

Type Package

Title A package to handle and analyse discrete Markov chains

Version 0.0.9.5

Date 2014-06-21

Author Giorgio Alfredo Spedicato

Maintainer Giorgio Alfredo Spedicato <spedicato_giorgio@yahoo.it>

Description A package for easily handling discrete Markov chains

License GPL-2

Depends R (>= 2.14), methods

Imports igraph, Matrix, matlab, expm, stats4, parallel

LazyLoad yes

ByteCompile yes

BugReports Giorgio Alfredo Spedicato <spedicato_giorgio@yahoo.it>

NeedsCompilation no

Repository CRAN

Date/Publication 2014-06-21 01:12:08

R topics documented:

`markovchain-package` *A package for easily handling discrete Markov chains*

Description

It contains S4 classes and methods to create and operates with Markov chains

Details

Package:	markovchain
Type:	Package
Version:	0.0.9.5
Date:	2014-06-21
License:	GPL-2
Depends:	R (>= 2.14), methods, expm, matlab, igraph, Matrix

Author(s)

Giorgio Alfredo Spedicato Maintainer: Giorgio Alfredo Spedicato <spedicato_giorgio@yahoo.it>

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

Examples

```
#create some markov chains
statesNames=c("a","b")
 mcA<-new("markovchain", transitionMatrix=matrix(c(0.7,0.3,0.1,0.9),byrow=TRUE,
 nrow=2, dimnames=list(statesNames,statesNames)))

statesNames=c("a","b","c")
```

```
mcB<-new("markovchain", states=statesNames, transitionMatrix=
          matrix(c(0.2,0.5,0.3,
                   0,1,0,
                   0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,
   statesNames)
                ))

statesNames=c("a","b","c","d")
matrice<-matrix(c(0.25,0.75,0,0,0.4,0.6,0,0,0,0,0.1,0.9,0,0,0.7,0.3),
nrow=4, byrow=TRUE)
mcC<-new("markovchain", states=statesNames, transitionMatrix=matrice)
mcD<-new("markovchain", transitionMatrix=matrix(c(0,1,0,1), nrow=2,byrow=TRUE))



#operations with S4 methods

mcA^2
steadyStates(mcB)
absorbingStates(mcB)
markovchainSequence(n=20, markovchain=mcC, include=TRUE)
```

---

absorbingStates *Various function to perform statistical and probabilistic analysis*

Description

These functions return absorbing and transient states of the markovchain objects.

Usage

```
absorbingStates(object)
transientStates(object)
canonicForm(object)
```

Arguments

object	A markovchain object.

Value

A matrix

Author(s)

Giorgio Spedicato

References

Feres, Matlab listing for markov chain.

See Also

markovchain

Examples

```
statesNames=c("a","b","c")
markovB<-new("markovchain", states=statesNames, transitionMatrix=
          matrix(c(0.2,0.5,0.3,
                   0,1,0,
                 0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)
                 ))
absorbingStates(markovB)
transientStates(markovB)
canonicForm(markovB)
#periodicity analysis
E=matrix(0,nrow=4,ncol=4)
E[1,2]=1
#E[2,c(1, 3)]=1/2;
E[2,1]=1/3;E[2,3]=2/3
#E[3,c(2, 4)]=1/2;
E[3,2]=1/4;E[3,4]=3/4
E[4,3]=1
mcE <- new("markovchain", states = c("a", "b", "c", "d"),
                 transitionMatrix = E,
                 name = "E")

is.irreducible(mcE) #true
period(mcE) #2
```

---

blanden *Mobility between income quartiles*

Description

This table show mobility between income quartiles for father and sons for the 1970 cohort born

Usage

```
data(blanden)
```

Format

The format is: table [1:4, 1:4] 0.38 0.25 0.21 0.16 0.29 0.28 0.26 0.17 0.22 0.26 ... - attr(*, "dimnames")=List of 2 ..$: chr [1:4] "Bottom" "2nd" "3rd" "Top" ..$: chr [1:4] "Bottom" "2nd" "3rd" "Top"

Details

The rows represent father's income quartile when the son is aged 16, whilst the columns represent sons' income quartiles when he is aged 30 (in 2000).

Source

Giorgio Spedicato from references

References

Jo Blanden, Paul Gregg and Stephen Machin, Intergenerational Mobility in Europe and North America, Center for Economic Performances (2005)

Examples

```
data(blanden)
mobilityMc<-as(blanden, "markovchain")
```

`conditionalDistribution` `conditionalDistribution` *of a Markov Chain*

Description

It extracts the conditional distribution of the subsequent state, given current state.

Usage

```
conditionalDistribution(object,state)
```

Arguments

`object`	A `markovchain` object.
`state`	Subsequent state.

Value

A named probability vector

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

markovchain

Examples

```
#define a markov chain
statesNames=c("a","b","c")
markovB<-new("markovchain", states=statesNames, transitionMatrix=matrix(c(0.2,0.5,0.3,
0,1,0,0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)))
conditionalDistribution(markovB,"b")
```

---

craigsendi *CD4 cells counts on HIV Infects between zero and six month*

Description

This is the table shown in Craig and Sendi paper showing zero and six month CD4 cells count in six brakets

Usage

```
data(craigsendi)
```

Format

The format is: table [1:3, 1:3] 682 154 19 33 64 19 25 47 43 - attr(*, "dimnames")=List of 2 ..$: chr [1:3] "0-49" "50-74" "75-UP" ..$: chr [1:3] "0-49" "50-74" "75-UP"

Details

Rows represent counts at the beginning, cols represent counts after six months.

Source

Estimation of the transition matrix of a discrete time Markov chain, Bruce A. Craig and Peter P. Sendi, Health Economics 11, 2002.

References

See source

Examples

```
data(craigsendi)
csMc<-as(craigsendi, "markovchain")
steadyStates(csMc)
```

`firstPassage` *First passage across states*

Description

This function compute the first passage probability in states

Usage

```
firstPassage(object, state, n)
```

Arguments

`object`	A `markovchain` object
`state`	Initial state
`n`	Number of rows on which compute the distribution

Details

Based on Feres' Matlab listings

Value

A matrix of size 1:n x number of states showing the probability of the first time of passage in states to be exactly the number in the row.

Author(s)

Giorgio Spedicato

References

Renaldo Feres, Notes for Math 450 Matlab listings for Markov chains

See Also

`conditionalDistribution`

Examples

```
#create a simple Markov chain
simpleMc<-new("markovchain", states=c("a","b"),
transitionMatrix=matrix(c(0.4,0.6,.3,.7),nrow=2,byrow=TRUE))
firstPassage(simpleMc,"b",20)
```

`is.accessible` *Verify if a state j is reachable from state i.*

Description

This function verifies if a state is reachable from another, i.e., if exists a path that leads to state j leaving from state i with positive probability

Usage

```
is.accessible(object, from, to)
```

Arguments

`object`	A `markovchain` object.
`from`	The name of state "i" (beginning state).
`to`	The name of state "j" (ending state).

Details

If wraps and internal function named `.commStatesFinder`.

Value

A boolean value.

Author(s)

Giorgio Spedicato

References

James Montgomery, University of Madison

See Also

`is.irreducible`

Examples

```
statesNames<-c("a","b","c")
markovB<-new("markovchain", states=statesNames, transitionMatrix=
          matrix(c(0.2,0.5,0.3,
                   0,1,0,
                 0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)
                 ))
is.accessible(markovB,"a","c")
```

`is.irreducible` *Function to check if a Markov chain is irreducible*

Description

This function verifies whether a `markovchain` object transition matrix is composed by only one communicating class.

Usage

```
is.irreducible(object)
```

Arguments

`object`	A `markovchain` object

Details

It is based on `.communicatingClasses` internal function.

Value

A boolean values.

Author(s)

Giorgio Spedicato

References

Feres, Matlab listings for Markov Chains.

See Also

`summary`

Examples

```
statesNames=c("a","b")
mcA<-new("markovchain", transitionMatrix=matrix(c(0.7,0.3,0.1,0.9),byrow=TRUE, nrow=2,
                                                dimnames=list(statesNames,statesNames)
                                                ))
is.irreducible(mcA)
```

markovchain-class *Class* "markovchain"

Description

The S4 class that describes markovchain objects.

Objects from the Class

Objects can be created by calls of the form new("markovchain", states, byrow, transitionMatrix, ...).

Slots

states: Name of the states. Must be the same of colnames and rownames of the transition matrix

byrow: Binary flag.

transitionMatrix: Square transition matrix

name: Optional character name of the Markov chain

Methods

* signature(e1 = "markovchain", e2 = "markovchain"): multiply two markovchain objects

* signature(e1 = "markovchain", e2 = "matrix"): markovchain by matrix multiplication

* signature(e1 = "markovchain", e2 = "numeric"): markovchain by numeric vector multiplication

* signature(e1 = "matrix", e2 = "markovchain"): matrix by markov chain

* signature(e1 = "numeric", e2 = "markovchain"): numeric vector by markovchain multiplication

[signature(x = "markovchain", i = "ANY", j = "ANY", drop = "ANY"): ...

^ signature(e1 = "markovchain", e2 = "numeric"): power of a markovchain object

== signature(e1 = "markovchain", e2 = "markovchain"): equality of two markovchain object

absorbingStates signature(object = "markovchain"): method to get absorbing states

canonicForm signature(object = "markovchain"): return a markovchain object into canonic form

coerce signature(from = "markovchain", to = "data.frame"): coerce method from markovchain to data.frame

conditionalDistribution signature(object = "markovchain"): returns the conditional probability of subsequent states given a state

coerce signature(from = "data.frame", to = "markovchain"): coerce method from data.frame to markovchain

coerce signature(from = "table", to = "markovchain"): coerce method from table to markovchain

coerce `signature(from = "markovchain", to = "igraph")`: coercing to `igraph` objects

coerce `signature(from = "markovchain", to = "matrix")`: coercing to `matrix` objects

coerce `signature(from = "matrix", to = "markovchain")`: coercing to `markovchain` objects from `matrix` one

dim `signature(x = "markovchain")`: method to get the size

initialize `signature(.Object = "markovchain")`: initialize method

plot `signature(x = "markovchain", y = "missing")`: plot method for `markovchain` objects

predict `signature(object = "markovchain")`: predict method

print `signature(x = "markovchain")`: print method.

show `signature(object = "markovchain")`: show method.

states `signature(object = "markovchain")`: states method.

steadyStates `signature(object = "markovchain")`: method to get the steady vector.

summary `signature(object = "markovchain")`: method to summarize structure of the markov chain

transientStates `signature(object = "markovchain")`: method to get the transient states.

t `signature(x = "markovchain")`: transpose matrix

transitionProbability `signature(object = "markovchain")`: transition probability

Warning

Validation method is used to assess whether either columns or rows totals to one. Rounding is used up to 5th decimal. If state names are not properly defined for a probability `matrix`, coercing to `markovhcain` object leads to overriding states name with artificial "s1", "s2", ... sequence

Note

`markovchain` object are written in S4 Classes.

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

`markovchainSequence`,`markovchainFit`

Examples

```
#show markovchain definition
showClass("markovchain")
#create a simple Markov chain
transMatr<-matrix(c(0.4,0.6,.3,.7),nrow=2,byrow=TRUE)
simpleMc<-new("markovchain", states=c("a","b"),
transitionMatrix=transMatr,
name="simpleMc")
#power
simpleMc^4
#some methods
steadyStates(simpleMc)
absorbingStates(simpleMc)
simpleMc[2,1]
t(simpleMc)
is.irreducible(simpleMc)
#conditional distributions
conditionalDistribution(simpleMc, "b")
#example for predict method
sequence<-c("a", "b", "a", "a", "a", "a", "b", "a", "b", "a", "b", "a", "a", "b", "b", "b", "a")
mcFit<-markovchainFit(data=sequence)
predict(mcFit$estimate, newdata="b",n.ahead=3)
#direct conversion
myMc<-as(transMatr, "markovchain")

#example of summary
summary(simpleMc)
## Not run: plot(simpleMc)
```

`markovchainFit` *Function to fit a discrete Markov chain*

Description

Given a sequence of states arising from a stationary state, it fits the underlying Markov chain distribution using either MLE (also using a Laplacian smoother) or bootstrap.

Usage

```
markovchainFit(data, method = "mle", byrow = TRUE, nboot = 10,laplacian=0,name,
parallel=FALSE)
createSequenceMatrix(stringchar, toRowProbs = FALSE, sanitize = TRUE)
```

Arguments

`data`	A character list.
`method`	Method used to estimate the Markov chain. Either "mle" or "bootstrap" or "laplace"

byrow	it tells whether the output Markov chain should show the transition probabilities by row.
nboot	Number of bootstrap replicates in case "bootstrap" is used.
laplacian	Laplacian smoothing parameter, default zero. It is only used when "laplace" method is chosen.
name	Optional character for name slot.
parallel	Use parallel processing when performing Boostrap estimates.
stringchar	Equivalent to data
toRowProbs	converts a sequence matrix into a probability matrix
sanitize	put 1 in all rows having rowSum equal to zero

Value

A list containing an estimate and, when "bootstrap" method is used, a matrix of standards deviations and the bootstrap samples.

Warning

"mle" method calls `createSequenceMatrix` function using `sanitize` parameter set to TRUE.

Note

When MLE method is called, the lists contains one entry: estimate. Bootstrap algorithm has been defined "euristically". In addition, parallel facility is not complete, involving only a part of the bootstrap process.

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

markovchainSequence

Examples

```
sequence<-c("a", "b", "a", "a", "a", "a", "b", "a", "b", "a", "b", "a", "a",
"b", "b", "b", "a")
sequenceMatr<-createSequenceMatrix(sequence,sanitize=FALSE)
mcFitMLE<-markovchainFit(data=sequence)
mcFitBSP<-markovchainFit(data=sequence,method="bootstrap",nboot=5, name="Bootstrap Mc")
```

`markovchainList-class` *Class* `"markovchainList"`

Description

A class to handle non - homogeneous Markov chains

Objects from the Class

Objects can be created by calls of the form `new("markovchainList", ...)`. Each item in the list is a `markovchain` object.

Slots

`markovchains`**:** Object of class `"list"`: a list of markovchains

`name`**:** Object of class `"character"`: optional name of the class

Methods

[[`signature(x = "markovchainList")`: extract the i-th `markovchain`

dim `signature(x = "markovchainList")`: number of `markovchain` underlying the matrix

predict `signature(object = "markovchainList")`: predict from a `markovchainList`

print `signature(x = "markovchainList")`: prints the list of markovchains

show `signature(object = "markovchainList")`: same as print

Note

The class consists in a list of `markovchain` objects. It can help to deal with non - homogeneous Markov chains.

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

`markovchain`

Examples

```
showClass("markovchainList")
#define a markovchainList
statesNames=c("a","b")

mcA<-new("markovchain",name="MCA", transitionMatrix=matrix(c(0.7,0.3,0.1,0.9),
byrow=TRUE, nrow=2, dimnames=list(statesNames,statesNames)))

mcB<-new("markovchain", states=c("a","b","c"), name="MCB",
transitionMatrix=matrix(c(0.2,0.5,0.3,0,1,0,0.1,0.8,0.1),
nrow=3, byrow=TRUE))

mcC<-new("markovchain", states=c("a","b","c","d"), name="MCC",
   transitionMatrix=matrix(c(0.25,0.75,0,0,0.4,0.6,
    0,0,0,0,0.1,0.9,0,0,0.7,0.3), nrow=4, byrow=TRUE)
)
mcList<-new("markovchainList",markovchains=list(mcA, mcB, mcC),
name="Non - homogeneous Markov Chain")
```

---

preproglucacon *Preprogluccacon DNA protein bases sequences*

Description

Sequence of bases for preproglucacon DNA protein

Usage

```
data(preproglucacon)
```

Format

A data frame with 1572 observations on the following 2 variables.

`V1` a numeric vector, showing original coding

`preproglucacon` a character vector, showing initial of DNA bases (Adenine, Cytosine, Guanine, Thymine)

Source

Avery Henderson

References

Averuy Henderson, Fitting markov chain models on discrete time series such as DNA sequences

Examples

```
data(preproglucacon)
preproglucaconMc<-markovchainFit(data=preproglucacon$preproglucacon)
```

rain *Alofi island daily rainfall*

Description

Rainfall measured in Alofi Island

Usage

```
data(rain)
```

Format

A data frame with 1096 observations on the following 2 variables.

`V1` a numeric vector, showing original coding

`rain` a character vector, showing daily rainfall millilitres brackets

Source

Avery Henderson

References

Avery Henderson, Fitting markov chain models on discrete time series such as DNA sequences

Examples

```
data(rain)
rainMc<-markovchainFit(data=rain$rain)
```

rmarkovchain *Function to generate a sequence of states from homogeneous or non-homogeneous Markov chains.*

Description

Provided any `markovchain` or `markovchainList` objects, it returns a sequence of states coming from the underlying stationary distribution.

Usage

```
rmarkovchain(n, object, ...)
markovchainSequence(n, markovchain, t0 = sample(markovchain@states, 1),
include.t0 = FALSE)
```

Arguments

n	Sample size
object	Either a markovchain or a markovchainList object.
...	additional parameters passed to the internal sampler
markovchain	The markovchain object
t0	The initial state.
include.t0	Specify if the initial state shall be used.

Details

When an homogeneous process is assumed (markovchain object) a sequence is sampled of size n. When an non - homogeneous process is assumed, n samples are taken but the process is assumed to last from the begin to the end of the non-homogeneous markov process.

Value

Either a character vector or a data frame

Note

Check the type of input

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

markovchainFit

Examples

```
#define the Markov chain
statesNames=c("a","b","c")
mcB<-new("markovchain", states=statesNames, transitionMatrix=matrix(c(0.2,0.5,0.3,
0,0.2,0.8,0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)
                ))
#show the sequence
outs<-markovchainSequence(n=100,markovchain=mcB, t0="a")
outs2<-rmarkovchain(n=20, object=mcB)
```

`states` *Defined states of a transition matrix*

Description

This method returns the states of a transition matrix.

Usage

```
states(object)
```

Arguments

`object`	A discrete `markovchain` object

Value

The character vector corresponding to states slot.

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

`markovchain`

Examples

```
statesNames=c("a","b","c")
markovB<-new("markovchain", states=statesNames, transitionMatrix=
          matrix(c(0.2,0.5,0.3,
                   0,1,0,
                 0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)
                 ))
states(markovB)
```

steadyStates *Stationary states of a* markovchain *objeect*

Description

This method returns the stationary vector in matricial form of a markovchain object.

Usage

```
steadyStates(object)
```

Arguments

object	A discrete markovchain object

Value

A matrix corresponding to the stationary states

Note

The steady states are identified starting from which eigenvectors correspond to identity eigenvalues and then normalizing them to sum up to unity.

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

markovchain

Examples

```
statesNames=c("a","b","c")
markovB<-new("markovchain", states=statesNames, transitionMatrix=
          matrix(c(0.2,0.5,0.3,
                   0,1,0,
                 0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)
                 ))
steadyStates(markovB)
```

`transitionProbability` *Function to get the transition probabilities from initial to subsequent status.*

Description

This is a convenience function to get transition probabilities.

Usage

```
transitionProbability(object,t0,t1)
```

Arguments

`object`	A `markovchain` object.
`t0`	Initial state.
`t1`	Subsequent state.

Value

A matrix

Author(s)

Giorgio Spedicato

References

A First Course in Probability (8th Edition), Sheldon Ross, Prentice Hall 2010

See Also

`markovchain`

Examples

```
#define a markov chain
statesNames=c("a","b","c")
markovB<-new("markovchain", states=statesNames, transitionMatrix=
          matrix(c(0.2,0.5,0.3,
                   0,1,0,
                 0.1,0.8,0.1),nrow=3, byrow=TRUE, dimnames=list(statesNames,statesNames)
                 ))
transitionProbability(markovB,"b", "c")
```

Index