

Package ‘gratis’

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Type Package

Title Generating Time Series with Diverse and Controllable Characteristics

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Description Generates time series based on mixture autoregressive models. Kang, Y., Hyndman, R., Li, F. (2020) <doi:10.1002/sam.11461>.

LazyLoad yes

Repository CRAN

URL <https://github.com/ykang/gratis>

BugReports <https://github.com/ykang/gratis/issues/>

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NeedsCompilation no

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| | |
|-------|---|
| arinf | <i>Compute pi coefficients from ARIMA model</i> |
|-------|---|

Description

Compute pi coefficients from ARIMA model

Usage

```
arinf(object)
```

Arguments

object An object of class "Arima"

Value

A vector of AR coefficients

Author(s)

Rob J Hyndman

Examples

```
# Not Run
```

`fitness_ts`*Fitness function for time series generation.*

Description

Fitness function for time series generation.

Usage

```
fitness_ts(  
  pars,  
  features,  
  seasonal,  
  n = 120,  
  freq = 12,  
  target,  
  nComp,  
  selected.features  
)
```

Arguments

| | |
|--------------------------------|---|
| <code>pars</code> | Parameters |
| <code>features</code> | Time series features. |
| <code>seasonal</code> | Seasonal effects. |
| <code>n</code> | Length of time series |
| <code>freq</code> | Frequency of time series |
| <code>target</code> | Target time series features |
| <code>nComp</code> | No. of components used in mixture models. |
| <code>selected.features</code> | Selected features. |

Value

NA

Examples

```
# Not Run
```

| | |
|-------|--|
| ga_ts | <i>A revised version of genetic algorithms (R package 'GA') to allow for time series generation.</i> |
|-------|--|

Description

A revised version of genetic algorithms (R package 'GA') to allow for time series generation.

Usage

```
ga_ts(
  type = c("binary", "real-valued", "permutation"),
  fitness,
  ...,
  n,
  min,
  max,
  nBits,
  population = gaControl(type)$population,
  selection = gaControl(type)$selection,
  crossover = gaControl(type)$crossover,
  mutation = gaControl(type)$mutation,
  popSize = 50,
  pcrossover = 0.8,
  pmutation = 0.1,
  elitism = base::max(1, round(popSize * 0.05)),
  updatePop = FALSE,
  postFitness = NULL,
  maxiter = 100,
  run = maxiter,
  maxFitness = Inf,
  names = NULL,
  suggestions = NULL,
  optim = FALSE,
  optimArgs = list(method = "L-BFGS-B", poptim = 0.05, pressel = 0.5, control =
    list(fnscale = -1, maxit = 100)),
  keepBest = FALSE,
  parallel = FALSE,
  monitor = if (interactive()) { if (shiny::is.RStudio()) gaMonitor
    else FALSE } else { FALSE },
  seed = NULL
)
```

Arguments

| | |
|------|--|
| type | the type of genetic algorithm to be run depending on the nature of decision variables. |
|------|--|

| | |
|-------------|---|
| fitness | the fitness function, any allowable R function which takes as input an individual string representing a potential solution, and returns a numerical value describing its “fitness“ |
| ... | additional arguments to be passed to the fitness function. |
| n | Length of the time series to be generated. |
| min | a vector of length equal to the decision variables providing the lower bounds of the search space in case of real-valued or permutation encoded optimizations. |
| max | a vector of length equal to the decision variables providing the upper bounds of the search space in case of real-valued or permutation encoded optimizations. |
| nBits | a value specifying the number of bits to be used in binary encoded optimizations. |
| population | an R function for randomly generating an initial population. |
| selection | an R function performing selection, i.e. a function which generates a new population of individuals from the current population probabilistically according to individual fitness. |
| crossover | an R function performing crossover, i.e. a function which forms offsprings by combining part of the genetic information from their parents. |
| mutation | an R function performing mutation, i.e. a function which randomly alters the values of some genes in a parent chromosome. |
| popSize | the population size. |
| pcrossover | the probability of crossover between pairs of chromosomes. |
| pmutation | the probability of mutation in a parent chromosome. |
| elitism | the number of best fitness individuals to survive at each generation. |
| updatePop | If set at TRUE the first attribute attached to the value returned by the user-defined fitness function is used to update the population. |
| postFitness | a user-defined function which, if provided, receives the current ga-class object as input, performs post fitness-evaluation steps, then returns an updated version of the object which is used to update the GA search. |
| maxiter | the maximum number of iterations to run before the GA search is halted. |
| run | the number of consecutive generations without any improvement in the best fitness value before the GA is stopped. |
| maxFitness | the upper bound on the fitness function after that the GA search is interrupted. |
| names | a vector of character strings providing the names of decision variables. |
| suggestions | a matrix of solutions strings to be included in the initial population. |
| optim | a logical defaulting to FALSE determining whether or not a local search using general-purpose optimisation algorithms should be used. |
| optimArgs | a list controlling the local search algorithm. |
| keepBest | a logical argument specifying if best solutions at each iteration should be saved in a slot called bestSol. |
| parallel | An optional argument which allows to specify if the Genetic Algorithm should be run sequentially or in parallel. |
| monitor | a logical or an R function which takes as input the current state of the ga-class object and show the evolution of the search. |
| seed | an integer value containing the random number generator state. |

Value

An object of class 'ga-class'.

Examples

```
# Not Run
```

| | |
|---------------|---|
| generate_msts | <i>Generate mutiple seasonal time series from random parameter spaces of the mixture autoregressive (MAR) models.</i> |
|---------------|---|

Description

Generate mutiple seasonal time series from random parameter spaces of the mixture autoregressive (MAR) models.

Usage

```
generate_msts(  
  seasonal.periods = c(7, 365),  
  n = 800,  
  nComp = NULL,  
  output_format = "list"  
)
```

Arguments

| | |
|------------------|--|
| seasonal.periods | a vector of seasonal periods of the time series to be generated. |
| n | length of the generated time series. |
| nComp | number of mixing components when simulating time series using MAR models. |
| output_format | An optional argument which allows to choose output format between "list" and "tsibble" |

Value

a time series with multiple seasonal periods.

Examples

```
x <- generate_msts(seasonal.periods = c(7, 365), n = 800, nComp = 2, output_format= "list")  
forecast::autoplot(x)
```

| | |
|-------------|--|
| generate_ts | <i>Generate time series from random parameter spaces of the mixture autoregressive (MAR) models.</i> |
|-------------|--|

Description

Generate time series from random parameter spaces of the mixture autoregressive (MAR) models.

Usage

```
generate_ts(n.ts = 1, freq = 1, nComp = NULL, n = 120, output_format = "list")
```

Arguments

| | |
|---------------|--|
| n.ts | number of time series to be generated. |
| freq | seasonal period of the time series to be generated. |
| nComp | number of mixing components when simulating time series using MAR models. |
| n | length of the generated time series. |
| output_format | An optional argument which allows to choose output format between "list" and "tsibble" |

Value

A list of time series together with the SARIMA coefficients used in each mixing component and the corresponding mixing weights.

Author(s)

Yanfei Kang and Feng Li

References

Wong, CS & WK Li (2000).

Examples

```
x <- generate_ts(n.ts = 2, freq = 12, nComp = 2, n = 120)
x$N1$pars
forecast::autoplot(x$N1$x)
```

`generate_ts_with_target`*Generating time series with controllable features.*

Description

Generating time series with controllable features.

Usage

```
generate_ts_with_target(  
  n,  
  ts.length,  
  freq,  
  seasonal,  
  features,  
  selected.features,  
  target,  
  parallel = TRUE,  
  output_format = "list"  
)
```

Arguments

| | |
|--------------------------------|--|
| <code>n</code> | number of time series to be generated. |
| <code>ts.length</code> | length of the time series to be generated. |
| <code>freq</code> | frequency of the time series to be generated. |
| <code>seasonal</code> | 0 for non-seasonal data, 1 for single-seasonal data, and 2 for multiple seasonal data. |
| <code>features</code> | a vector of function names. |
| <code>selected.features</code> | selected features to be controlled. |
| <code>target</code> | target feature values. |
| <code>parallel</code> | An optional argument which allows to specify if the Genetic Algorithm should be run sequentially or in parallel. |
| <code>output_format</code> | An optional argument which allows to choose output format between 'list' and 'tsibble' |

Value

A time-series object of class "ts" or "msts".

Author(s)

Yanfei Kang

Examples

```
library(tsfeatures)
x <- generate_ts_with_target(n = 1, ts.length = 60, freq = 1, seasonal = 0,
  features = c('entropy', 'stl_features'), selected.features = c('entropy', 'trend'),
  target=c(0.6, 0.9), parallel=FALSE)
forecast::autoplot(x)
```

nsdiffs1 *Set the number of seasonal differences for yearly data to be -1.*

Description

Set the number of seasonal differences for yearly data to be -1.

Usage

```
nsdiffs1(x)
```

Arguments

x Univariate time series or numerical vector

Value

A numerical scalar value

Examples

```
# Not Run
```

pi_coefficients *Compute pi coefficients of an AR process from SARIMA coefficients.*

Description

Convert SARIMA coefficients to pi coefficients of an AR process.

Usage

```
pi_coefficients(
  ar = 0,
  d = 0L,
  ma = 0,
  sar = 0,
  D = 0L,
  sma = 0,
  m = 1L,
  tol = 1e-07
)
```

Arguments

| | |
|-----|--|
| ar | AR coefficients in the SARIMA model. |
| d | number of differences in the SARIMA model. |
| ma | MA coefficients in the SARIMA model. |
| sar | seasonal AR coefficients in the SARIMA model. |
| D | number of seasonal differences in the SARIMA model. |
| sma | seasonal MA coefficients in the SARIMA model. |
| m | seasonal period in the SARIMA model. |
| tol | tolerance value used. Only return up to last element greater than tolerance. |

Value

A vector of AR coefficients.

Author(s)

Rob J Hyndman

Examples

```
# Not Run
```

rlnorm2

Log-normal distribution with alternative parametrization.

Description

Alternative parametrization of log normal distribution.

Usage

```
rlnorm2(n, mean, sd)
```

Arguments

| | |
|------|---|
| n | number of observations. |
| mean | "vector" the mean value of the log-normal distribution. |
| sd | "vector" the variance of the log-normal distribution. |

Details

See help("rlnorm") for the details for the log-normal distribution.

Value

See the corresponding help for the usual log-normal functions.

Author(s)

Feng Li, Central University of Finance and Economics .

References

Feng Li, Mattias Villani, and Robert Kohn. (2010). Flexible Modeling of Conditional Distributions using Smooth Mixtures of Asymmetric Student T Densities, *Journal of Statistical Planning and Inference*, 140(12), pp. 3638-3654

 rmixnorm

Generate random variables from mixture normal distribution.

Description

Random variables from mixture of normals.

Usage

```
rmixnorm(n, means, sigmas, weights)
```

Arguments

| | |
|---------|---|
| n | "integer", numbers of samples to be generated. |
| means | "q-by-k matrix" mean value within each component, total k components. |
| sigmas | "q-by-q-by-k" variance covariance matrix with in each component. |
| weights | "k-length vector" weights in each component. |

Value

"matrix".

Author(s)

Feng Li, Central University of Finance and Economics.

References

Villani et al 2009.

Examples

```
n <- 1000
means <- matrix(c(-5, 0, 5), 1)
sigmas <- array(c(1, 1, 1), c(1, 1, 3))
weights <- c(0.3, 0.4, 0.3)
out <- rmixnorm(n, means, sigmas, weights)
hist(out, breaks = 100, freq = FALSE)
```

 rmixnorm_ts

Simulate AR type random variables from mixture of normal

Description

This function simulates random samples from a finite mixture of Gaussian distribution where the mean from each components are AR(p) process.

Usage

```
rmixnorm_ts(n, means.ar.par.list, sigmas.list, weights, yinit = 0)
```

Arguments

| | |
|-------------------|---|
| n | number of samples. |
| means.ar.par.list | parameters in AR(p) within each mixing component. |
| sigmas.list | variance list. |
| weights | weight in each list. |
| yinit | initial values. |

Value

vector of length n follows a mixture distribution.

Author(s)

Feng Li, Central University of Finance and Economics.

References

Feng Li, Mattias Villani, and Robert Kohn. (2010). Flexible Modeling of Conditional Distributions using Smooth Mixtures of Asymmetric Student T Densities, *Journal of Statistical Planning and Inference*, 140(12), pp. 3638-3654.

Examples

```
n = 1000
means.ar.par.list = list(c(0, 0.8), c(0, 0.6, 0.3))
require("fGarch")
sigmas.spec <- list(garchSpec(model = list(alpha = c(0.05, 0.06)), cond.dist = "norm"),
                   garchSpec(model = list(alpha = c(0.05, 0.05)), cond.dist = "norm"))
sigmas.list <- lapply(lapply(sigmas.spec, garchSim, extended = TRUE, n = n),
                     function(x) x$sigma)
weights <- c(0.8, 0.2)
y = rmixnorm_ts(n = n, means.ar.par.list = means.ar.par.list, sigmas.list = sigmas.list,
               weights = weights)
plot(y)
```

| | |
|----------------|--|
| run_gratis_app | <i>Web Application to generate time series with controllable features.</i> |
|----------------|--|

Description

Web Application to generate time series with controllable features.

Usage

```
run_gratis_app()
```

```
app_gratis()
```

Examples

```
# Not Run
```

| | |
|--------------|-------------------------------|
| tsgeneration | <i>Time Series Generation</i> |
|--------------|-------------------------------|

Description

The tsgeneration package generates time series data based on MAR models.

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