Package 'dbnR'

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Type Package

Title Dynamic Bayesian Network Learning and Inference

Version 0.8.0

Description Learning and inference over dynamic Bayesian networks of arbitrary Markovian order. Extends some of the functionality offered by the 'bnlearn' package to learn the networks from data and perform exact inference. It offers three structure learning algorithms for dynamic Bayesian networks: Trabelsi G. (2013) <doi:10.1007/978-3-642-41398-8_34>, Santos F.P. and Maciel C.D. (2014) <doi:10.1109/BRC.2014.6880957>, Quesada D., Bielza C. and Larrañaga P. (2021) <doi:10.1007/978-3-030-86271-8_14>. It also offers the possibility to perform forecasts of arbitrary length. A tool for visualizing the structure of the net is also provided via the 'visNetwork' package. Further detailed information and examples can be found in our Journal of Statistical Software paper Quesada D., Larrañaga P. and Bielza C. (2025) <doi:10.18637/jss.v115.i06>.

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AIC.dbn — *Calculate the AIC of a dynamic Bayesian network*

Description

Generic method for calculating the Akaike information criterion (AIC) of a "dbn" S3 object given some data. Calls bnlearn's AIC underneath.

Usage

```
## S3 method for class 'dbn'
AIC(object, ..., k)
```

Arguments

object	the structure of the network
...	additional parameters for the network scoring
k	the penalty parameter

Value

the AIC score of the network

AIC.dbn.fit — *Calculate the AIC of a dynamic Bayesian network*

Description

Generic method for calculating the Akaike information criterion (AIC) of a "dbn.fit" S3 object given some data. Calls bnlearn's AIC underneath.

Usage

```
## S3 method for class 'dbn.fit'
AIC(object, ..., k)
```

Arguments

object	the fitted network
...	additional parameters for the network scoring
k	the penalty parameter

Value

the AIC score of the network

all.equal.dbn — *Check if two network structures are equal to each other*

Description

Generic method for checking the equality of two "dbn" S3 objects. Calls bnlearn's all.equal underneath.

Usage

```
## S3 method for class 'equal.dbn'
all(target, current, ...)
```

Arguments

target	"dbn" object
current	the other "dbn" object
...	additional parameters

Value

boolean result of the comparison

all.equal.dbn.fit — *Check if two fitted networks are equal to each other*

Description

Generic method for checking the equality of two "dbn.fit" S3 objects. Calls bnlearn's all.equal underneath.

Usage

```
## S3 method for class 'equal.dbn.fit'
all(target, current, ...)
```

Arguments

target	"dbn.fit" object
current	the other "dbn.fit" object
...	additional parameters

Value

boolean result of the comparison

as.character.dbn *Convert a network structure into a model string*

Description

Generic method for converting a "dbn" S3 object into a string. Calls bnlearn's `as.character` underneath.

Usage

```
## S3 method for class 'dbn'
as.character(x, ...)
```

Arguments

`x`	a "dbn" object
`...`	additional parameters

Value

string representing the DBN model

BIC.dbn *Calculate the BIC of a dynamic Bayesian network*

Description

Generic method for calculating the Bayesian information criterion (BIC) of a "dbn" S3 object given some data. Calls bnlearn's `BIC` underneath.

Usage

```
## S3 method for class 'dbn'
BIC(object, ...)
```

Arguments

`object`	the structure of the network
`...`	additional parameters for the network scoring

Value

the BIC score of the network

BIC.dbn.fit *Calculate the BIC of a dynamic Bayesian network*

Description

Generic method for calculating the Bayesian information criterion (BIC) of a "dbn.fit" S3 object given some data. Calls bnlearn's BIC underneath.

Usage

```
## S3 method for class 'dbn.fit'
BIC(object, ...)
```

Arguments

object	the fitted network
...	additional parameters for the network scoring

Value

the BIC score of the network

calc_mu *Calculate the mu vector from a fitted BN or DBN*

Description

Given a "bn.fit" or a "dbn.fit" object, calculate the mu vector of the equivalent multivariate Gaussian distribution. Front end of a C++ function.

Usage

```
calc_mu(fit)
```

Arguments

fit	a bn.fit or dbn.fit object

Value

a named numeric vector of the means of each variable

Examples

```
dt_train <- dbnR::motor[200:2500]
net <- bnlearn::mmhc(dt_train)
fit <- bnlearn::bn.fit(net, dt_train, method = "mle-g")
mu <- dbnR::calc_mu(fit)

f_dt_train <- dbnR::fold_dt(dt_train, size = 2)
net <- dbnR::learn_dbn_struc(dt_train, size = 2)
fit <- dbnR::fit_dbn_params(net, f_dt_train)
mu <- dbnR::calc_mu(fit)
```

---

calc_sigma *Calculate the sigma covariance matrix from a fitted BN or DBN*

Description

Given a "bn.fit" or a "dbn.fit" object, calculate the sigma covariance matrix of the equivalent multivariate Gaussian distribution. Front end of a C++ function.

Usage

```
calc_sigma(fit)
```

Arguments

fit	a bn.fit or dbn.fit object

Value

a named numeric covariance matrix of the nodes

Examples

```
dt_train <- dbnR::motor[200:2500]
net <- bnlearn::mmhc(dt_train)
fit <- bnlearn::bn.fit(net, dt_train, method = "mle-g")
sigma <- dbnR::calc_sigma(fit)

f_dt_train <- dbnR::fold_dt(dt_train, size = 2)
net <- dbnR::learn_dbn_struc(dt_train, size = 2)
fit <- dbnR::fit_dbn_params(net, f_dt_train)
sigma <- dbnR::calc_sigma(fit)
```

coef.dbn.fit *Extracts the coefficients of a DBN*

Description

Generic method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 method for class 'dbn.fit'
coef(object, ...)
```

Arguments

object	the fitted network
...	additional parameters

Value

the coefficients of the network

degree *Calculates the degree of a list of nodes*

Description

#' Generic method for calculating the degree of a list of nodes in a BN or a DBN. Calls bnlearn's degree underneath. I have to redefine the generic and mask the original for it to work on both bn and dbn objects without the user having to import bnlearn.

Usage

```
degree(object, Nodes, ...)
```

Arguments

object	a "bn", "dbn", "bn.fit" or "dbn.fit" object
Nodes	which nodes to check
...	additional parameters

Value

the degree of the nodes

filtered_fold_dt *Fold a dataset avoiding overlapping of different time series*

Description

If the dataset that is going to be folded contains several different time series instances of the same process, folding it could introduce false rows with data from different time series. Given an id variable that labels the different instances of a time series inside a dataset and a desired size, this function folds the dataset and avoids mixing data from different origins in the same instance.

Usage

```
filtered_fold_dt(dt, size, id_var, clear_id_var = TRUE)
```

Arguments

dt	data.table to be folded
size	the size of the data.table
id_var	the variable that labels each individual instance of the time series
clear_id_var	boolean that decides whether or not the id_var column is deleted

Value

the filtered data.table

Examples

```
dt <- dbnR::motor[201:2500]
dt[, n_sec := rep(seq(46), each = 50)] # I'll create secuences of 50 instances each
f_dt <- dbnR::fold_dt(dt, size = 2)
dim(f_dt)
f_dt <- dbnR::filtered_fold_dt(dt, size = 2, id_var = "n_sec")
dim(f_dt)  # The filtered folded dt has a row less for each independent secuence
```

filter_same_cycle *Filter the instances in a data.table with different ids in each row*

Description

Given an id variable that labels the different instances of a time series inside a dataset, discard the rows that have values from more than 1 id.

Usage

```
filter_same_cycle(f_dt, size, id_var)
```

Arguments

f_dt	folded data.table
size	the size of the data.table
id_var	the variable that labels each individual instance of the time series

Value

the filtered data.table

Examples

```
dt <- dbnR::motor[201:2500]
dt[, n_sec := rep(seq(46), each = 50)] # I'll create secuences of 50 instances each
f_dt <- dbnR::fold_dt(dt, size = 2)
f_dt[50, .SD, .SDcols = c("n_sec_t_0", "n_sec_t_1")]
f_dt <- dbnR::filter_same_cycle(f_dt, size = 2, id_var = "n_sec")
f_dt[50, .SD, .SDcols = c("n_sec_t_0", "n_sec_t_1")]
```

---

fitted.dbn.fit *Extracts the fitted values of a DBN*

Description

Generic method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 method for class 'dbn.fit'
fitted(object, ...)
```

Arguments

object	the fitted network
...	additional parameters

Value

the fitted values of the network

fit_dbn_params *Fits a markovian n DBN model*

Description

Fits the parameters of the DBN via MLE. The "mu" vector of means and the "sigma" covariance matrix are set as attributes of the dbn.fit object for future exact inference.

Usage

```
fit_dbn_params(net, f_dt, ...)
```

Arguments

net	the structure of the DBN
f_dt	a folded data.table
...	additional parameters for the bn.fit function

Value

a "dbn.fit" S3 object with the fitted net

Examples

```
size = 3
dt_train <- dbnR::motor[200:2500]
net <- learn_dbn_struc(dt_train, size)
f_dt_train <- fold_dt(dt_train, size)
fit <- fit_dbn_params(net, f_dt_train, method = "mle-g")
```

fold_dt *Widens the dataset to take into account the t previous time slices*

Description

This function will widen the dataset to put the t previous time slices in each row, so that it can be used to learn temporal arcs in the second phase of the dmmhc.

Usage

```
fold_dt(dt, size)
```

Arguments

dt	the data.table to be treated
size	number of time slices to unroll. Markovian 1 would be size 2

Value

the extended data.table

Examples

```
data(motor)
size <-  3
f_dt <- fold_dt(motor, size)
```

`forecast_ts` *Performs forecasting with the GDBN over a dataset*

Description

Given a dbn.fit object, the size of the net and a folded dataset, performs a forecast over the initial evidence taken from the dataset.

Usage

```
forecast_ts(
  dt,
  fit,
  size = NULL,
  obj_vars,
  ini = 1,
  len = dim(dt)[1] - ini,
  rep = 1,
  num_p = 50,
  print_res = TRUE,
  plot_res = TRUE,
  mode = "exact",
  prov_ev = NULL
)
```

Arguments

`dt`	data.table object with the TS data
`fit`	dbn.fit object
`size`	number of time slices of the net. Deprecated, will be removed in the future
`obj_vars`	variables to be predicted
`ini`	starting point in the dataset to forecast.
`len`	length of the forecast
`rep`	number of times to repeat the approximate forecasting
`num_p`	number of particles in the approximate forecasting
`print_res`	if TRUE prints the mae and sd metrics of the forecast

plot_res	if TRUE plots the results of the forecast
mode	"exact" for exact inference, "approx" for approximate
prov_ev	variables to be provided as evidence in each forecasting step

Value

a list with the original time series values and the results of the forecast

Examples

```
size = 3
data(motor)
dt_train <- motor[200:900]
dt_val <- motor[901:1000]
obj <- c("pm_t_0")
net <- learn_dbn_struc(dt_train, size)
f_dt_train <- fold_dt(dt_train, size)
f_dt_val <- fold_dt(dt_val, size)
fit <- fit_dbn_params(net, f_dt_train, method = "mle-g")
res <- suppressWarnings(forecast_ts(f_dt_val, fit,
        obj_vars = obj, len = 10, print_res = FALSE, plot_res = FALSE))
```

---

generate_random_network_exp

Generate a random DBN and a sampled dataset

Description

This function generates both a random DBN and a dataset that can be used to learn its structure from data. It's intended for experimental use.

Usage

```
generate_random_network_exp(
  n_vars,
  size,
  min_mu,
  max_mu,
  min_sd,
  max_sd,
  min_coef,
  max_coef,
  seed = NULL
)
```

Arguments

n_vars	number of desired variables per time-slice
size	desired size of the networks
min_mu	minimum mean allowed for the variables
max_mu	maximum mean allowed for the variables
min_sd	minimum standard deviation allowed for the variables
max_sd	maximum standard deviation allowed for the variables
min_coef	minimum coefficient allowed for the parent nodes
max_coef	maximum coefficient allowed for the parent nodes
seed	the seed of the experiment

Value

a list with the original network structure and the sampled dataset

---

learn_dbn_struc *Learns the structure of a markovian n DBN model from data*

Description

Learns a gaussian dynamic Bayesian network from a dataset. It allows the creation of markovian n nets rather than only markov 1.

Usage

```
learn_dbn_struc(dt, size = 2, method = "dmmhc", f_dt = NULL, ...)
```

Arguments

dt	the data.frame or data.table to be used
size	number of time slices of the net. Markovian 1 would be size 2
method	the structure learning method of choice to use
f_dt	previously folded dataset, in case some specific rows have to be removed after the folding
...	additional parameters for rsmax2 function

Value

a "dbn" S3 object with the structure of the network

Examples

```
data("motor")
net <- learn_dbn_struc(motor, size = 3)
```

logLik.dbn *Calculate the log-likelihood of a dynamic Bayesian network*

Description

Generic method for calculating the log-likelihood of a "dbn" S3 object given some data. Calls bnlearn's logLik underneath.

Usage

```
## S3 method for class 'dbn'
logLik(object, dt, ...)
```

Arguments

object	the structure of the network
dt	the dataset to calculate the score of the network
...	additional parameters for the network scoring

Value

the log-likelihood score of the network

logLik.dbn.fit *Calculate the log-likelihood of a dynamic Bayesian network*

Description

Generic method for calculating the log-likelihood of a "dbn.fit" S3 object given some data. Calls bnlearn's logLik underneath.

Usage

```
## S3 method for class 'dbn.fit'
logLik(object, dt, ...)
```

Arguments

object	the fitted network
dt	the dataset to calculate the score of the network
...	additional parameters for the network scoring

Value

the log-likelihood score of the network

mean.dbn.fit *Average the parameters of multiple dbn.fit objects with identical structures*

Description

Generic method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 method for class 'dbn.fit'
mean(x, ...)
```

Arguments

x	the fitted network
...	additional parameters

Value

the averaged parameters

motor *Multivariate time series dataset on the temperature of an electric motor*

Description

Data from several sensors on an electric motor that records different benchmark sessions of measurements at 2 Hz. The dataset is reduced to 3000 instances from the 60th session in order to include it in the package for testing purposes. For the complete dataset, refer to the source.

Usage

```
data(motor)
```

Format

An object of class `data.table` (inherits from `data.frame`) with 3000 rows and 11 columns.

Source

Kaggle, <https://www.kaggle.com/wkirgsn/electric-motor-temperature>

`mvn_inference` *Performs inference over a multivariate normal distribution*

Description

Given some evidence, this function performs inference over a multivariate normal distribution. After converting a Gaussian linear network to its MVN form, this kind of inference can be performed. It's recommended to use `predict_dt` functions instead unless you need a more flexible inference method.

Usage

```
mvn_inference(mu, sigma, evidence)
```

Arguments

`mu`	the mean vector
`sigma`	the covariance matrix
`evidence`	a single row data.table or a named vector with the values and names of the variables given as evidence

Value

a list with the posterior mean and covariance matrix

Examples

```
size = 3
data(motor)
dt_train <- motor[200:2500]
dt_val <- motor[2501:3000]
obj <- c("pm_t_0")

net <- learn_dbn_struc(dt_train, size)
f_dt_train <- fold_dt(dt_train, size)
f_dt_val <- fold_dt(dt_val, size)
ev <- f_dt_val[1, .SD, .SDcols = obj]
fit <- fit_dbn_params(net, f_dt_train, method = "mle-g")

pred <- mvn_inference(calc_mu(fit), calc_sigma(fit), ev)
```

nodes — *Returns a list with the names of the nodes of a BN or a DBN*

Description

Generic method for obtaining the names of the nodes in a BN or a DBN. Calls bnlearn's nodes underneath. I have to redefine the generic and mask the original for it to work on both bn and dbn objects without the user having to import bnlearn.

Usage

```
nodes(object, ...)
```

Arguments

object	a "bn", "dbn", "bn.fit" or "dbn.fit" object
...	additional parameters

Value

the names of the nodes

nodes<- — *Relabel the names of the nodes of a BN or a DBN*

Description

Generic method for renaming the nodes in a BN or a DBN. Calls bnlearn's nodes<- underneath. I have to redefine the generic and mask the original for it to work on both bn and dbn objects without the user having to import bnlearn.

Usage

```
nodes(object) <- value
```

Arguments

object	a "bn", "dbn", "bn.fit" or "dbn.fit" object
value	a list with the new names

Value

the modified object

plot.dbn *Plots a dynamic Bayesian network*

Description

Generic method for plotting the "dbn" S3 objects. Calls `plot_dynamic_network` underneath.

Usage

```
## S3 method for class 'dbn'
plot(x, ...)
```

Arguments

x	the structure of the network.
...	additional parameters for the visualization of a DBN

plot.dbn.fit *Plots a fitted dynamic Bayesian network*

Description

Generic method for plotting the "dbn.fit" S3 objects. Calls `plot_dynamic_network` underneath.

Usage

```
## S3 method for class 'dbn.fit'
plot(x, ...)
```

Arguments

x	the structure of the network.
...	additional parameters for the visualization of a DBN

`plot_dynamic_network` *Plots a dynamic Bayesian network in a hierarchical way*

Description

To plot the DBN, this method first computes a hierarchical structure for a time slice and replicates it for each slice. Then, it calculates the relative position of each node with respect to his equivalent in the first slice. The result is a net where each time slice is ordered and separated from one another, where the leftmost slice is the oldest and the rightmost represents the present time. This function is also called by the generic plot function of "dbn" and "dbn.fit" S3 objects.

Usage

```
plot_dynamic_network(
  structure,
  offset = 200,
  subset_nodes = NULL,
  reverse = FALSE
)
```

Arguments

`structure`	the structure or fit of the network.
`offset`	the blank space between time slices
`subset_nodes`	a vector containing the names of the subset of nodes to plot
`reverse`	reverse to the classic naming convention of the nodes. The oldest time-slice will now be t_0 and the most recent one t_n. Only for visualization purposes, the network is unmodified underneath. If using subset_nodes, remember that t_0 is now the oldest time-slice.

Value

the visualization of the DBN

Examples

```
size = 3
dt_train <- dbnR::motor[200:2500]
net <- learn_dbn_struc(dt_train, size)
plot_dynamic_network(net)
```

plot_static_network *Plots a Bayesian network in a hierarchical way*

Description

This function calculates the levels of each node and then plots them in a hierarchical layout in visNetwork. Can be used in place of the generic plot function offered by bnlearn for "bn" and "bn.fit" S3 objects.

Usage

```
plot_static_network(structure)
```

Arguments

structure	the structure or fit of the network.

Examples

```
dt_train <- dbnR::motor[200:2500]
net <- bnlearn::mmhc(dt_train)
plot_static_network(net)
fit <- bnlearn::bn.fit(net, dt_train, method = "mle-g")
plot_static_network(fit) # Works for both the structure and the fitted net
```

predict.dbn.fit *Performs inference in every row of a dataset with a DBN*

Description

Generic method for predicting a dataset with a "dbn.fit" S3 objects. Calls predict_dt underneath.

Usage

```
## S3 method for class 'dbn.fit'
predict(object, ...)
```

Arguments

object	a "dbn.fit" object
...	additional parameters for the inference process

Value

a data.table with the prediction results

predict_bn *Performs inference over a fitted GBN*

Description

Performs inference over a Gaussian BN. It's thought to be used in a map for a data.table, to use as evidence each separate row. If not specifically needed, it's recommended to use the function predict_dt instead. This function is deprecated and will be removed in a future version.

Usage

```
predict_bn(fit, evidence)
```

Arguments

fit	the fitted bn
evidence	values of the variables used as evidence for the net

Value

a data.table with the predictions

Examples

```
size = 3
data(motor)
dt_train <- motor[200:2500]
dt_val <- motor[2501:3000]
net <- learn_dbn_struc(dt_train, size)
f_dt_train <- fold_dt(dt_train, size)
f_dt_val <- fold_dt(dt_val, size)
fit <- fit_dbn_params(net, f_dt_train, method = "mle-g")
res <- f_dt_val[, predict_bn(fit, .SD), .SDcols = c("pm_t_0", "coolant_t_0"), by = 1:nrow(f_dt_val)]
```

predict_dt *Performs inference over a test dataset with a GBN*

Description

This function performs inference over each row of a folded data.table, plots the results and gives metrics of the accuracy of the predictions. Given that only a single row is predicted, the horizon of the prediction is at most 1. This function is also called by the generic predict method for "dbn.fit" objects. For long term forecasting, please refer to the forecast_ts function.

Usage

```
predict_dt(fit, dt, obj_nodes, verbose = T, look_ahead = F)
```

Arguments

fit	the fitted bn
dt	the test dataset
obj_nodes	the nodes that are going to be predicted. They are all predicted at the same time
verbose	if TRUE, displays the metrics and plots the real values against the predictions
look_ahead	boolean that defines whether or not the values of the variables in t_0 should be used when predicting, even if they are not present in obj_nodes. This decides if look-ahead bias is introduced or not.

Value

a data.table with the prediction results for each row

Examples

```
size = 3
data(motor)
dt_train <- motor[200:900]
dt_val <- motor[901:1000]

# With a DBN
obj <- c("pm_t_0")
net <- learn_dbn_struc(dt_train, size)
f_dt_train <- fold_dt(dt_train, size)
f_dt_val <- fold_dt(dt_val, size)
fit <- fit_dbn_params(net, f_dt_train, method = "mle-g")
res <- suppressWarnings(predict_dt(fit, f_dt_val, obj_nodes = obj, verbose = FALSE))

# With a Gaussian BN directly from bnlearn
obj <- c("pm")
net <- bnlearn::mmhc(dt_train)
fit <- bnlearn::bn.fit(net, dt_train, method = "mle-g")
res <- suppressWarnings(predict_dt(fit, dt_val, obj_nodes = obj, verbose = FALSE))
```

---

print.dbn *Print method for "dbn" objects*

Description

Generic print method for "dbn" S3 objects. Calls bnlearn's print underneath

Usage

```
## S3 method for class 'dbn'
print(x, ...)
```

Arguments

x	the "dbn" object
...	additional parameters

`print.dbn.fit` *Print method for "dbn.fit" objects*

Description

Generic print method for "dbn.fit" S3 objects. Calls bnlearn's print underneath

Usage

```
## S3 method for class 'dbn.fit'
print(x, ...)
```

Arguments

x	the "dbn.fit" object
...	additional parameters

`rbn.dbn.fit` *Simulates random samples from a fitted DBN*

Description

Generic method for "dbn.fit" S3 objects. Calls bnlearn's rbn underneath.

Usage

```
rbn.dbn.fit(x, n, ...)
```

Arguments

x	the fitted network
n	number of samples
...	additional parameters

Value

the sampled dataset

reduce_freq *Reduce the frequency of the time series data in a data.table*

Description

In a time series dataset, there is a time difference between one row and the next one. This function reduces the number of rows from its current frequency to the desired one by averaging batches of rows. Instead of the frequency in Hz, the number of seconds between rows is asked (Hz = 1/s).

Usage

```
reduce_freq(dt, obj_freq, curr_freq, id_var = NULL)
```

Arguments

dt	the original data.table
obj_freq	the desired number of seconds between rows
curr_freq	the number of seconds between rows in the original dataset
id_var	optional variable that labels different time series in a dataset, to avoid averaging values from different processes

Value

the data.table with the desired frequency

Examples

```
# Let's assume that the dataset has a frequency of 4Hz, 0.25 seconds between rows
dt <- dbnR::motor
dim(dt)
# Let's change the frequency to 2Hz, 0.5 seconds between rows
dt <- reduce_freq(dt, obj_freq = 0.5, curr_freq = 0.2)
dim(dt)
```

residuals.dbn.fit *Returns the residuals from fitting a DBN*

Description

Generic method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 method for class 'dbn.fit'
residuals(object, ...)
```

Arguments

object	the fitted network
...	additional parameters

Value

the residuals of fitting the network

---

score *Computes the score of a BN or a DBN*

Description

Generic method for computing the score of a BN or a DBN. Calls bnlearn's nodes underneath. I have to redefine the generic and mask the original for it to work on both bn and dbn objects without the user having to import bnlearn.

Usage

```
score(object, ...)
```

Arguments

object	a "bn" or "dbn" object
...	additional parameters

Value

the score of the network

---

shift_values *Move the window of values backwards in a folded dataset row*

Description

This function moves the values in t_0, t_1, ..., t_n-1 in a folded dataset row to t_1, t_2, ..., t_n. All the variables in t_0 will be inputed with NAs and the obtained row can be used to forecast up to any desired point.

Usage

```
shift_values(f_dt, row)
```

Arguments

f_dt	a folded dataset
row	the index of the row that is going to be processed

Value

a one row data.table the shifted values

Examples

```
dt <- dbnR::motor
f_dt <- dbnR::fold_dt(dt, size = 2)
s_row <- dbnR::shift_values(f_dt, row = 500)
```

---

sigma.dbn.fit *Returns the standard deviation of the residuals from fitting a DBN*

Description

Generic method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 method for class 'dbn.fit'
sigma(object, ...)
```

Arguments

object	the fitted network
...	additional parameters

Value

the standard deviation residuals of fitting the network

`smooth_ts` *Performs smoothing with the GDBN over a dataset*

Description

Given a dbn.fit object, the size of the net and a folded dataset, performs a smoothing of a trajectory. Smoothing is the opposite of forecasting: given a starting point, predict backwards in time to obtain the time series that generated that point.

Usage

```
smooth_ts(
  dt,
  fit,
  size = NULL,
  obj_vars,
  ini = dim(dt)[1],
  len = ini - 1,
  print_res = TRUE,
  plot_res = TRUE,
  prov_ev = NULL
)
```

Arguments

`dt`	data.table object with the TS data
`fit`	dbn.fit object
`size`	number of time slices of the net. Deprecated, will be removed in the future
`obj_vars`	variables to be predicted. Should be in the oldest time step
`ini`	starting point in the dataset to smooth
`len`	length of the smoothing
`print_res`	if TRUE prints the mae and sd metrics of the smoothing
`plot_res`	if TRUE plots the results of the smoothing
`prov_ev`	variables to be provided as evidence in each smoothing step. Should be in the oldest time step

Value

a list with the original values and the results of the smoothing

Examples

```
size = 3
data(motor)
dt_train <- motor[200:900]
dt_val <- motor[901:1000]
obj <- c("pm_t_2")
net <- learn_dbn_struc(dt_train, size)
f_dt_train <- fold_dt(dt_train, size)
f_dt_val <- fold_dt(dt_val, size)
fit <- fit_dbn_params(net, f_dt_train, method = "mle-g")
res <- suppressWarnings(smooth_ts(f_dt_val, fit,
        obj_vars = obj, len = 10, print_res = FALSE, plot_res = FALSE))
```

---

time_rename *Renames the columns in a data.table so that they end in '_t_0'*

Description

This will rename the columns in a data.table so that they end in '_t_0', which will be needed when folding the data.table. If any of the columns already ends in '_t_0', a warning will be issued and no further operation will be done. There is no need to use this function to learn a DBN unless some operation with the variable names wants to be done prior to folding a dataset.

Usage

```
time_rename(dt)
```

Arguments

dt	the data.table to be treated

Value

the renamed data.table

Examples

```
data("motor")
dt <- time_rename(motor)
```

[[<-.dbn.fit *Replacement function for parameters inside DBNs*

Description

Generic parameter replacement method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 replacement method for class 'dbn.fit'
x[[name]] <- value
```

Arguments

x	the fitted network
name	name of the node to replace its parameters
value	the new parameters

Value

the modified network

$<-.dbn.fit *Replacement function for parameters inside DBNs*

Description

Generic parameter replacement method for "dbn.fit" S3 objects. Calls bnlearn underneath.

Usage

```
## S3 replacement method for class 'dbn.fit'
x$name <- value
```

Arguments

x	the fitted network
name	name of the node to replace its parameters
value	the new parameters

Value

the modified network

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