

Package ‘dbnR’

June 11, 2020

Type Package

Title Dynamic Bayesian Network Learning and Inference

Version 0.4.5

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 a modification of Trabelsi (2013) <doi:10.1007/978-3-642-41398-8_34> dynamic max-min hill climbing algorithm for structure learning and the possibility to perform forecasts of arbitrary length. A tool for visualizing the structure of the net is also provided via the 'visNetwork' package.

Depends R (>= 3.5.0)

Imports bnlearn (>= 4.5), data.table (>= 1.12.4), Rcpp (>= 1.0.2), magrittr (>= 1.5)

Suggests visNetwork (>= 2.0.8), grDevices (>= 3.6.0), utils (>= 3.6.0), graphics (>= 3.6.0), stats (>= 3.6.0), testthat (>= 2.1.0)

LinkingTo Rcpp

URL <https://github.com/dkesada/dbnR>

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 7.1.0

NeedsCompilation yes

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Repository CRAN

Date/Publication 2020-06-11 08:40:02 UTC

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acc_successions	<i>Returns a vector with the number of consecutive nodes in each level</i>
-----------------	--

Description

This method processes the vector of node levels to get the position of each node inside the level. E.g. `c(1,1,1,2,2,3,4,4,5,5)` turns into `c(1,2,3,1,2,1,1,2,1,2)`

Usage

```
acc_successions(nodes, res = NULL, prev = 0, acc = 0)
```

Arguments

nodes	a vector with the level of each node
res	the accumulative results of the sub successions
prev	the level of the previous node processed
acc	the accumulator of the index in the current sub successions

Value

the vector of sub successions in each level

add_attr_to_fit	<i>Adds the mu vector and sigma matrix as attributes to the bn.fit or dbn.fit object</i>
-----------------	--

Description

Adds the mu vector and sigma matrix as attributes to the bn.fit or dbn.fit object to allow performing exact MVN inference on both cases.

Usage

```
add_attr_to_fit(fit)
```

Arguments

fit	a fitted bn or dbn
-----	--------------------

Value

the fitted net with attributes

approximate_inference	<i>Performs approximate inference forecasting with the GDBN over a data set</i>
-----------------------	---

Description

Given a bn.fit object, the size of the net and a data.set, performs approximate forecasting with bnlearns cpdist function over the initial evidence taken from the data set.

Usage

```
approximate_inference(dt, fit, size, obj_vars, ini, rep, len, num_p)
```

Arguments

dt	data.table object with the TS data
fit	bn.fit object
size	number of time slices of the net
obj_vars	variables to be predicted
ini	starting point in the data set to forecast.
rep	number of repetitions to be performed of the approximate inference
len	length of the forecast
num_p	number of particles to be used by bnlearn

Value

the results of the forecast

approx_prediction_step

Performs approximate inference in a time slice of the dbn

Description

Given a bn.fit object and some variables, performs particle inference over such variables in the net for a given time slice.

Usage

```
approx_prediction_step(fit, variables, particles, n = 50)
```

Arguments

fit	bn.fit object
variables	variables to be predicted
particles	a list with the provided evidence
n	the number of particles to be used by bnlearn

Value

the inferred particles

calc_mu	<i>Calculate the mu vector of means of a Gaussian linear network. Front end of a C++ function.</i>
---------	--

Description

Calculate the mu vector of means of a Gaussian linear network. 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")
mu <- calc_mu(fit)
```

calc_mu_cpp	<i>Calculate the mu vector of means of a Gaussian linear network. This is the C++ backend of the function.</i>
-------------	--

Description

Calculate the mu vector of means of a Gaussian linear network. This is the C++ backend of the function.

Usage

```
calc_mu_cpp(fit, order)
```

Arguments

fit a bn.fit object as a Rcpp::List
order a topological ordering of the nodes as a vector of strings

Value

the map with the nodes and their mu. Returns as a named numeric vector

calc_sigma	<i>Calculate the sigma covariance matrix of a Gaussian linear network. Front end of a C++ function.</i>
------------	---

Description

Calculate the sigma covariance matrix of a Gaussian linear network. Front end of a C++ function.

Usage

```
calc_sigma(fit)
```

Arguments

fit	a bn.fit or dbn.fit object
-----	----------------------------

Value

a 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")
sigma <- calc_sigma(fit)
```

calc_sigma_cpp	<i>Calculate the sigma covariance matrix of a Gaussian linear network. This is the C++ backend of the function.</i>
----------------	---

Description

Calculate the sigma covariance matrix of a Gaussian linear network. This is the C++ backend of the function.

Usage

```
calc_sigma_cpp(fit, order)
```

Arguments

fit	a bn.fit object as a Rcpp::List
order	a topological ordering of the nodes as a vector of strings

Value

the covariance matrix

check_time0_formatted *Checks if the vector of names are time formatted to t0*

Description

This will check if the names are properly time formatted in `t_0` to be folded into more time slices. A vector is well formatted in `t_0` when all of its column names end in `'_t_0'`.

Usage

```
check_time0_formatted(obj)
```

Arguments

`obj` the vector of names

Value

TRUE if it is well formatted. FALSE in other case.

create_blacklist *Creates the blacklist of arcs from a folded data.table*

Description

This will create the blacklist of arcs that are not to be learned in the second phase of the dmmhc. This includes arcs backwards in time or inside time-slices.

Usage

```
create_blacklist(name, size, acc = NULL, slice = 1)
```

Arguments

`name` the names of the first time slice, ended in `_t_0`
`size` the number of time slices of the net. Markovian 1 would be size 2
`acc` accumulator of the results in the recursion
`slice` current time slice that is being processed

Value

the two column matrix with the blacklisted arcs

dynamic_ordering	<i>Gets the ordering of a single time slice in a DBN</i>
------------------	--

Description

This method gets the structure of a DBN, isolates the nodes of a single time slice and then gives a topological ordering of them.

Usage

```
dynamic_ordering(structure)
```

Arguments

structure	the structure of the network.
-----------	-------------------------------

Value

the ordered nodes of t_0

exact_inference	<i>Performs exact inference forecasting with the GDBN over a data set</i>
-----------------	---

Description

Given a bn.fit object, the size of the net and a data.set, performs exact forecasting over the initial evidence taken from the data set.

Usage

```
exact_inference(dt, fit, size, obj_vars, ini, len, prov_ev)
```

Arguments

dt	data.table object with the TS data
fit	bn.fit object
size	number of time slices of the net
obj_vars	variables to be predicted
ini	starting point in the data set to forecast.
len	length of the forecast
prov_ev	variables to be provided as evidence in each forecasting step

Value

the results of the forecast

exact_prediction_step *Performs exact inference in a time slice of the dbn*

Description

Given a bn.fit object and some variables, performs exact MVN inference over such variables in the net for a given time slice.

Usage

```
exact_prediction_step(fit, variables, evidence)
```

Arguments

fit	list with the mu and sigma of the MVN model
variables	variables to be predicted
evidence	a list with the provided evidence

Value

the inferred particles

expand_time_nodes *Extends the names of the nodes in t_0 to t_(max-1)*

Description

This method extends the names of the nodes to the given maximum and maintains the order of the nodes in each slice, so as to plotting the nodes in all slices relative to their homonyms in the first slice.

Usage

```
expand_time_nodes(name, acc, max, i)
```

Arguments

name	the names of the nodes in the t_0 slice
acc	accumulator of the resulting names in the recursion
max	number of time slices in the net
i	current slice being processed

Value

the extended names

fit_dbn_params	<i>Fits a markovian n DBN model</i>
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Description

Fits the parameters of the DBN via MLE or BGE. 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

the fitted net

Examples

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

fold_dt	<i>Widens the dataset to take into account the t previous time slices</i>
---------	---

Description

This 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
dt <- fold_dt(motor, size)
```

fold_dt_rec

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

Description

This 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. Recursive version not exported, the user calls from the handler 'fold_dt'

Usage

```
fold_dt_rec(dt, n_prev, size, slice = 1)
```

Arguments

dt	the data.table to be treated
n_prev	names of the previous time slice
size	number of time slices to unroll. Markovian 1 would be size 2
slice	the current time slice being treated. Should not be modified when first calling.

Value

the extended data.table

forecast_ts

Performs forecasting with the GDBN over a data set

Description

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

Usage

```
forecast_ts(
  dt,
  fit,
  size,
  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
obj_vars	variables to be predicted
ini	starting point in the data set 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

the results of the forecast

Examples

```
size = 3
data(motor)
dt_train <- motor[200:2500]
dt_val <- motor[2501:3000]
obj <- c("pm_t_0", "torque_t_0")
net <- learn_dbn_struct(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")
```

```
res <- suppressWarnings(forecast_ts(f_dt_val, fit, size,
  obj_vars = obj, print_res = FALSE, plot_res = FALSE))
```

learn_dbn_struc	<i>Learns the structure of a markovian n DBN model from data</i>
-----------------	--

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, blacklist = 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
blacklist	an optional matrix indicating forbidden arcs between nodes
...	additional parameters for rsmx2 function

Value

the structure of the net

Examples

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

merge_nets	<i>Merges and replicates the arcs in the static BN into all the time-slices in the DBN</i>
------------	--

Description

This will join the static net and the state transition net by replicating the arcs in the static net in all the time slices.

Usage

```
merge_nets(net0, netCP1, size, acc = NULL, slice = 1)
```

Arguments

<code>net0</code>	the structure of the static net
<code>netCP1</code>	the state transition net
<code>size</code>	the number of time slices of the net. Markovian 1 would be size 2
<code>acc</code>	accumulator of the results in the recursion
<code>slice</code>	current time slice that is being processed

Value

the merged nets

<code>motor</code>	<i>Multivariate time series dataset on the temperature of an electric motor</i>
--------------------	---

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 4th 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 12 columns.

Source

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

mvn_inference	<i>Performs inference over a multivariate normal distribution</i>
---------------	---

Description

Performs inference over a multivariate normal distribution given some evidence. After converting a Gaussian linear network to its MVN form, this kind of inference can be performed. It's recommended to use the [predict_bn](#) or [predict_dt](#) functions instead unless you need the posterior mean vector and covariance matrix.

Usage

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

Arguments

mu	the mean vector
sigma	the covariance matrix
evidence	a named vector with the values and names of the variables given as evidence

Value

the posterior mean and covariance matrix

Examples

```
as_named_vector <- function(dt){
  res <- as.numeric(dt)
  names(res) <- names(dt)

  return(res)
}
size = 3
data(motor)
dt_train <- motor[200:2500]
dt_val <- motor[2501:3000]
obj <- c("pm_t_0", "torque_t_0")

net <- learn_dbn_struct(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")

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

node_levels	<i>Defines a level for every node in the net</i>
-------------	--

Description

Calculates the levels in which the nodes will be distributed when plotting the structure. This level is defined by their parent nodes: a node with no parents will always be in the level 0. Subsequently, the level of a node will be one more of the maximum level of his parents.

Usage

```
node_levels(net, order, lvl = 1, acc = NULL)
```

Arguments

net	the structure of the network.
order	a topological order of the nodes, with the orphan nodes in the first place. See node.ordering
lvl	current level being processed
acc	accumulator of the nodes already processed

Value

a matrix with the names of the nodes in the first row and their level on the second

plot_dynamic_network	<i>Plots a dynamic Bayesian network in a hierarchical way</i>
----------------------	---

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.

Usage

```
plot_dynamic_network(structure, offset = 200)
```

Arguments

structure	the structure or fit of the network.
offset	the blank space between time slices

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_network	<i>Plots a Bayesian networks in a hierarchical way</i>
--------------	--

Description

Calculates the levels of each node and then plots them in a hierarchical layout in visNetwork.

Usage

```
plot_network(structure)
```

Arguments

structure the structure or fit of the network.

Examples

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

predict_bn	<i>Performs inference over a fitted GBN</i>
------------	---

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.

Usage

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

Arguments

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

Value

the mean of the particles for each row

Examples

```
size = 3
data(motor)
dt_train <- motor[200:2500]
dt_val <- motor[2501:3000]
net <- learn_dbn_struct(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")
res <- f_dt_val[, predict_bn(fit, .SD), by = 1:nrow(f_dt_val)]
```

predict_dt	<i>Performs inference over a test data set with a GBN</i>
------------	---

Description

Performs inference over a test data set, plots the results and gives metrics of the accuracy of the results.

Usage

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

Arguments

fit the fitted bn
dt the test data set
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

Value

the prediction results

Examples

```

size = 3
data(motor)
dt_train <- motor[200:2500]
dt_val <- motor[2501:3000]

# With a DBN
obj <- c("pm_t_0", "torque_t_0")
net <- learn_dbn_struct(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")
res <- suppressWarnings(predict_dt(fit, f_dt_val, obj_nodes = obj, verbose = FALSE))

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

Usage

```
time_rename(dt)
```

Arguments

dt the data.table to be treated

Value

the renamed data.table

Examples

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

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