

# Package ‘dbnR’

March 25, 2020

**Type** Package

**Title** Dynamic Bayesian Network Learning and Inference

**Version** 0.3.4

**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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## R topics documented:

acc_successions . . . . .	2
add_attr_to_fit . . . . .	3
approximate_inference . . . . .	3
approx_prediction_step . . . . .	4
calc_mu . . . . .	5
calc_mu_cpp . . . . .	5
calc_sigma . . . . .	6
calc_sigma_cpp . . . . .	6
check_time0_formatted . . . . .	7
create_blacklist . . . . .	7
dynamic_ordering . . . . .	8
exact_inference . . . . .	8
exact_prediction_step . . . . .	9
expand_time_nodes . . . . .	9
fit_dbn_params . . . . .	10
fold_dt . . . . .	10
fold_dt_rec . . . . .	11
forecast_ts . . . . .	11
learn_dbn_struct . . . . .	13
merge_nets . . . . .	13
motor . . . . .	14
mvn_inference . . . . .	14
node_levels . . . . .	15
plot_dynamic_network . . . . .	16
plot_network . . . . .	16
predict_bn . . . . .	17
predict_dt . . . . .	18
time_rename . . . . .	19
<b>Index</b>	<b>20</b>

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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.
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**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)
```

**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

**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>
----------------	-------------------------------------

---

**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 <a href="#">bn.fit</a> 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"
)
```

**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

**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_struct	<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_struct(dt, size = 2, ...)
```

**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
...	additional parameters for <code>rsmx2</code> function

**Value**

the structure of the net

**Examples**

```
data("motor")
net <- learn_dbn_struct(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**

net0	the structure of the static net
netCP1	the state transition net
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 merged nets

---

motor	<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

*Defines a level for every node in the net*


---

**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 <a href="#">node.ordering</a>
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 *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.

### 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 *Plots a Bayesian networks in a hierarchical way*

---

### 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

*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.

**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`*Performs inference over a test data set with a GBN*

---

### 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

<code>fit</code>	the fitted bn
<code>dt</code>	the test data set
<code>obj_nodes</code>	the nodes that are going to be predicted. They are all predicted at the same time
<code>verbose</code>	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	<i>Renames the columns in a data.table so that they end in '_t_0'</i>
-------------	---

---

**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)
```

# Index

## \*Topic **datasets**

motor, [14](#)

acc\_successions, [2](#)

add\_attr\_to\_fit, [3](#)

approx\_prediction\_step, [4](#)

approximate\_inference, [3](#)

bn.fit, [10](#)

calc\_mu, [5](#)

calc\_mu\_cpp, [5](#)

calc\_sigma, [6](#)

calc\_sigma\_cpp, [6](#)

check\_time0\_formatted, [7](#)

create\_blacklist, [7](#)

dynamic\_ordering, [8](#)

exact\_inference, [8](#)

exact\_prediction\_step, [9](#)

expand\_time\_nodes, [9](#)

fit\_dbn\_params, [10](#)

fold\_dt, [10](#)

fold\_dt\_rec, [11](#)

forecast\_ts, [11](#)

learn\_dbn\_struct, [13](#)

merge\_nets, [13](#)

motor, [14](#)

mvn\_inference, [14](#)

node\_ordering, [15](#)

node\_levels, [15](#)

plot\_dynamic\_network, [16](#)

plot\_network, [16](#)

predict\_bn, [14](#), [17](#)

predict\_dt, [14](#), [17](#), [18](#)

rsmx2, [13](#)

time\_rename, [19](#)