Package 'bnstruct'

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Type Package

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Author Francesco Sambo	
Maintainer Francesco Sambo <pre></pre>	
Description More about what it does (maybe more than one line)	
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add.observations<-

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Description

Add a list of observations to an InferenceEngine that already has observations, using a list composed by the two following vectors:

- observed.varsvector of observed variables;
- observed.valsvector of values observed for the variables in observed.vars in the corresponding position.

Usage

```
add.observations(x) \leftarrow value
```

Arguments

x an InferenceEngine.value the list of observations of the InferenceEngine.

Details

In case of multiple observations of the same variable, the last observation is the one used, as the most recent.

See Also

observations<-

asia_10000

asia

load Asia dataset.

Description

Wrapper for a loader for the Asia dataset, with only raw data.

Usage

```
asia()
```

Details

The dataset has 10000 items, no missing data, so no imputation needs to be performed.

Value

a BNDataset containing the Child dataset.

See Also

```
asia_10000
```

Examples

```
dataset <- asia()
print(dataset)</pre>
```

asia_10000

Asia dataset.

Description

The Asia dataset contains 10000 complete (no missing data, no latent variables) randomly generated items of the Asia Bayesian Network. No imputation needs to be performed, so only raw data is present.

Usage

```
asia_10000
```

Format

a BNDataset with raw data slow filled.

belief.propagation 5

Details

The data the BNDataset object is built from is located in files pkg_folder/extdata/asia_10000.header and pkg_folder/extdata/asia_10000.data.

References

S. Lauritzen, D. Spiegelhalter. Local Computation with Probabilities on Graphical Structures and their Application to Expert Systems (with discussion). Journal of the Royal Statistical Society: Series B (Statistical Methodology), 50(2):157-224, 1988.

See Also

asia

belief.propagation

perform belief propagation.

Description

Perform belief propagation for the network of an InferenceEngine, given a set of observations when present. In the current version of bnstruct, belief propagation can be computed only over a junction tree.

Usage

```
belief.propagation(ie, ...)
## S4 method for signature 'InferenceEngine'
belief.propagation(ie, net = NULL,
   observed.vars = NULL, observed.vals = NULL, return.potentials = FALSE)
```

Arguments

```
ie an InferenceEngine object.

net a BN object.

observed.vars list of observed variables.

observed.vals values taken by variables listed in observed.vars.

return.potentials

if TRUE only the potentials are returned, instead of the default BN.

potential further arguments of methods.
```

Value

updated InferenceEngine object.

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Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
bn <- BN(dataset)
ie <- InferenceEngine(bn)
ie <- belief.propagation(ie)

observations(ie) <- list("observed.vars"=("A","G","X"), "observed.vals"=c(1,2,1))
belief.propagation(ie)

## End(Not run)</pre>
```

bn

get the BN object contained in an InferenceEngine.

Description

Return a network contained in an InferenceEngine.

Usage

```
bn(x)
## S4 method for signature 'InferenceEngine'
bn(x)
```

Arguments

Х

an InferenceEngine.

Value

the BN object contained in an InferenceEngine.

BN-class

BN class.

Description

BN class.

Instantiate a BN object.

BN-class 7

Usage

```
## S4 method for signature 'BN'
initialize(.Object, dataset = NULL, ...)
BN(dataset = NULL, ...)
```

Arguments

name name of the Bayesian Network.

dataset a BNDataset object containing the dataset the network is built upon, if any. The remaining parameters are considered only if a starting dataset is provided.

... potential further arguments of methods.

.0bject An object: see the Details section.

Details

The constructor may be invoked without parameters – in this case an empty network will be created, and its slots will be filled manually by the user. This is usually viable only if the user already has knowledge about the network structure.

Value

BN object.

Slots

```
name: name of the network

num.nodes: number of nodes in the network

variables: names of the variables in the network

discreteness: TRUE if variable is discrete, FALSE if variable is continue

node.sizes: if variable i is discrete, node.sizes[i] contains the cardinality of i, if i is instead discrete the value is the number of states variable i takes when discretized

cpts: list of conditional probability tables of the network

dag: adjacency matrix of the network

wpdag: weighted partially dag
```

Examples

```
## Not run:
net.1 <- BN()

dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
net.2 <- BN(dataset)

## End(Not run)</pre>
```

8 BNDataset-class

bn<-

set the original BN object contained in an InferenceEngine.

Description

Add an original network to an InferenceEngine.

Usage

```
bn(x) \leftarrow value
```

Arguments

x an InferenceEngine.

value the BN object contained in an InferenceEngine.

BNDataset-class

BNDataset class.

Description

Contains the all of the data that can be extracted from a given dataset: raw data, imputed data, raw and imputed data with bootstrap.

```
initialize a BNDataset object.
constructor for BNDataset object
```

Usage

```
## $4 method for signature 'BNDataset'
initialize(.Object, ...)
BNDataset(name = "", ...)
```

Arguments

.Object an empty BNDataset. name name of the dataset.

... potential further arguments of methods.

Details

Dataset should be provided in the following format... (describe)

BNDataset-class 9

Value

```
a BNDataset object.
BNDataset object.
```

Slots

```
name: name of the dataset
header.file: name and location of the header file
data.file: name and location of the data file
variables: names of the variables in the network
node.sizes: cardinality of each variable of the network
num. variables: number of variables (columns) in the dataset
discreteness: TRUE if variable is discrete, FALSE if variable is continue
num.items: number of observations (rows) in the dataset
has.rawdata: TRUE if the dataset contains data read from a file
has.impdata: TRUE if the dataset contains imputed data (computed from raw data)
raw.data: matrix containing raw data
imputation: TRUE if it dataset contains imputed data
imputed.data: matrix containing imputed data
has.boots: dataset has bootstrap samples
boots: list of bootstrap samples
has.imp.boots: dataset has imputed bootstrap samples
imp.boots: list of imputed bootstrap samples
num.boots: number of bootstrap samples
```

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
## End(Not run)</pre>
```

10 boots<-

boots

get list of bootstrap samples of a BNDataset.

Description

Return the list of samples computed from raw data of a dataset.

Usage

```
boots(x)
## S4 method for signature 'BNDataset'
boots(x)
```

Arguments

Х

a BNDataset object.

Value

the list of bootstrap samples.

See Also

```
has.boots, has.imp.boots, imp.boots
```

boots<-

set list of bootstrap samples of a BNDataset.

Description

Add to a dataset a list of samples from raw data computed using bootstrap.

Usage

```
boots(x) <- value</pre>
```

Arguments

x a BNDataset object.

value the list of bootstrap samples.

bootstrap 11

bootstrap Perform bootstrap.

Description

Create a list of num. boots samples of the original dataset.

Usage

```
bootstrap(object, ...)
## S4 method for signature 'BNDataset'
bootstrap(object, num.boots = 100, seed = 0,
  imputation = FALSE, k.impute = 10, na.string.symbol = "?", ...)
```

Arguments

object the BNDataset object.

num.boots number of sampled datasets for bootstrap.

seed random seed.

imputation TRUE if imputation has to be performed.

na.string.symbol character that denotes NA in the dataset (useful only if imputation == TRUE).

k.impute number of neighbours to be used; for discrete variables we use mode, for continuous variables the median value is instead taken (useful only if imputation == TRUE).

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
dataset <- bootstrap(dataset, num.boots = 1000)
## End(Not run)</pre>
```

potential further arguments of methods.

12 child

```
build.junction.tree build a JunctionTree.
```

Description

Starting from the adjacency matrix of the directed acyclic graph of the network contained in an InferenceEngine, build a JunctionTree for the network and store it into an InferenceEngine.

Usage

```
build.junction.tree(object, ...)
## S4 method for signature 'InferenceEngine'
build.junction.tree(object, ...)
```

Arguments

```
object an InferenceEngine object.
... potential further arguments for methods.
```

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
net <- BN(dataset)
eng <- InferenceEngine()
eng <- build.junction.tree(eng)
## End(Not run)</pre>
```

child

load Child dataset.

Description

Wrapper for a loader for the Child raw dataset; also perform imputation.

Usage

```
child()
```

Details

The dataset has 5000 items, with random missing values (no latent variables). BNDataset object contains the raw dataset and imputed dataset, with k=10 (see impute for related explanation).

child_NA_5000

Value

a BNDataset containing the Child dataset.

See Also

```
child_NA_5000
```

Examples

```
dataset <- child()
print(dataset)</pre>
```

child_NA_5000

Child dataset.

Description

The Child dataset contains 5000 randomly generated items with missing data (no latent variables) of the Child Bayesian Network. Imputation is performed, so both raw and imputed data is present.

Usage

```
child_NA_5000
```

Format

a BNDataset with a raw and imputed data slow filled with 5000 items.

Details

The data the BNDataset object is built from is located in files $pkg_folder/extdata/extdata/Child_data_na_5000$. heade and $pkg_folder/extdata/extdata/Child_data_na_5000$. data.

References

D. J. Spiegelhalter, R. G. Cowell (1992). Learning in probabilistic expert systems. In Bayesian Statistics 4 (J. M. Bernardo, J. 0. Berger, A. P. Dawid and A. F. M. Smith, eds.) 447-466. Clarendon Press, Oxford.

See Also

child

14 cpts<-

cpts

get the list of conditional probability tables of a BN.

Description

Return the list of conditional probability tables of the variables of a BN object. Each probability table is associated to the corresponding variable, and its dimensions are named according to the variable they represent.

Usage

```
cpts(x)
## S4 method for signature 'BN'
cpts(x)
```

Arguments

Χ

an object.

Details

Each conditional probability table is represented as a multidimensional array. The ordering of the dimensions of each variable is not guaranteed to follow the actual conditional distribution. E.g. dimensions for conditional probability P(C|A,B) can be either (C,A,B) or (A,B,C), depending on if some operations have been performed, or how the probability table has been computed. Users should not rely on dimension numbers, but should instead select the dimensions using their names.

Value

list of the conditional probability tables of the desired object.

cpts<-

set the list of conditional probability tables of a network.

Description

Set the list of conditional probability tables of a BN object.

Usage

```
cpts(x) <- value
```

Arguments

x an object.

value list of the conditional probability tables of the object.

dag 15

Details

Each conditional probability table is represented as a multidimensional array. To retrieve single dimensions (e.g. to compute marginals), users should provide dimensions names.

dag

get adjacency matrix of a network.

Description

Return the adjacency matrix of the directed acyclic graph representing the structure of a network.

Usage

```
dag(x)
## S4 method for signature 'BN'
dag(x)
```

Arguments

Х

an object.

Value

matrix containing the adjacency matrix of the directed acyclic graph representing the structure of the object.

dag<-

set adjacency matrix of an object.

Description

Set the adjacency matrix of the directed acyclic graph representing the structure of a network.

Usage

```
dag(x) \leftarrow value
```

Arguments

Х

an object.

value

matrix containing the adjacency matrix of the directed acyclic graph representing the structure of the object.

16 data.file<-

data.file

get data file of a BNDataset.

Description

Return the data filename of a dataset (with the path to its position, as given by the user). The data filename may contain a header in the first row, containing the list of names of the variables, in the same order as in the header file. After the header, if present, the file contains a data.frame with the observations, one item per row.

Usage

```
data.file(x)
## S4 method for signature 'BNDataset'
data.file(x)
```

Arguments

Х

a BNDataset.

Value

data filename of the dataset.

See Also

```
data.file
```

data.file<-

set data file of a BNDataset.

Description

Set the data filename of a dataset (with the path to its position, as given by the user). The data filename may contain a header in the first row, containing the list of names of the variables, in the same order as in the header file. After the header, if present, the file contains a data.frame with the observations, one item per row.

Usage

```
data.file(x) \leftarrow value
```

Arguments

x a BNDataset. value data filename. discreteness 17

See Also

```
header.file<-
```

discreteness

get status (discrete or continuous) of the variables of an object.

Description

Get a vector representing the status of the variables (with their names) of a BN or BNDataset. Elements of the vector are c if the variable is continue, and d if the variable is discrete.

Usage

```
discreteness(x)
## S4 method for signature 'BNDataset'
discreteness(x)
## S4 method for signature 'BN'
discreteness(x)
```

Arguments

x an object.

Value

vector containing, for each variable of the desired object, c if the variable is continue, and d if the variable is discrete.

discreteness<-

set status (discrete or continuous) of the variables of an object.

Description

Set the list of variable status for the variables in a network or a dataset.

Usage

```
discreteness(x) \leftarrow value
```

Arguments

x an object.

value a vector of elements in {c,d} for continuous and discrete variables (respec-

tively).

18 em

 em

expectation-maximization algorithm.

Description

Learn parameters of a network using the Expectation-Maximization algorithm.

Usage

```
em(x, dataset, ...)
## S4 method for signature 'InferenceEngine,BNDataset'
em(x, dataset, threshold = 0.001,
   k.impute = 10, ...)
```

Arguments

Х	an InferenceEngine.
dataset	observed dataset with missing values for the Bayesian Network of x.
threshold	threshold for convergence, used as stopping criterion.
k.impute	number of neighbours to be used; for discrete variables we use mode, for continuous variables the median value is instead taken.
	further potential arguments for method.

Value

a list containing: an InferenceEngine with a new updated network ("InferenceEngine"), and the imputed dataset ("BNDataset").

Examples

```
## Not run:
em(x, dataset)
## End(Not run)
```

get.boot 19

get.boot

get selected element of bootstrap list.

Description

Given a BNDataset, return the sample corresponding to given index.

Usage

```
get.boot(dataset, index, imputed, ...)
## S4 method for signature 'BNDataset,numeric'
get.boot(dataset, index, imputed = TRUE, ...)
```

Arguments

dataset a BNDataset object.

index the index of the requested sample.

imputed TRUE if samples from imputed dataset are to be used.
... potential further arguments of methods (ignored).

See Also

bootstrap

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
dataset <- bootstrap(dataset, num.boots = 1000)

for (i in 1:num.boots(dataset))
    print(get.boot(dataset, i))

## End(Not run)</pre>
```

get.data

get data of a BNDataset.

Description

Return data contained in a BNDataset object, if any. Preference is given to imputed data, if available, because the imputed dataset is (supposed to be), in general, more useful. To obtain specifically raw or imputed data, one must revert to get.raw.data() and get.imputed.data(), respectively.

20 get.imputed.data

Usage

```
get.data(x)
## S4 method for signature 'BNDataset'
get.data(x)
```

Arguments

x a BNDataset.

See Also

has.data, has.raw.data, has.imputed.data, get.raw.data, get.imputed.data

Examples

```
## Not run:
x <- BNDataset()
x <- read.dataset(x, "file.header", "file.data")
get.data(x) # returns raw dataset, the only one present in dataset
x <- impute(x)
get.data(x) # returns imputed dataset, since it is present now
## End(Not run)</pre>
```

get.imputed.data

get imputed data of a BNDataset.

Description

Return imputed data contained in a BNDataset object, if any.

Usage

```
get.imputed.data(x)
## S4 method for signature 'BNDataset'
get.imputed.data(x)
```

Arguments

x a BNDataset.

See Also

has.data, has.raw.data, has.imputed.data, get.data, get.raw.data

get.most.probable.values

```
get.most.probable.values
```

compute the most probable values to be observed.

Description

Return an array containing the values that each variable of the network is more likely to take, according to the CPTS. In case of ties take the first value.

Usage

```
get.most.probable.values(x, ...)
## S4 method for signature 'BN'
get.most.probable.values(x, ...)
## S4 method for signature 'InferenceEngine'
get.most.probable.values(x, ...)
```

Arguments

```
x a BN or InferenceEngine object.
```

... potential further arguments of methods.

Value

array containing, in each position, the most probable value for the corresponding variable.

Examples

```
## Not run:
# try with a BN object x
get.most.probable.values(x)
# now build an InferenceEngine object
eng <- InferenceEngine(x)
get.most.probable.values(eng)
## End(Not run)</pre>
```

22 has.boots

get.raw.data

get raw data of a BNDataset.

Description

Return raw data contained in a BNDataset object, if any.

Usage

```
get.raw.data(x)
## S4 method for signature 'BNDataset'
get.raw.data(x)
```

Arguments

Х

a BNDataset.

See Also

has.data, has.raw.data, has.imputed.data, get.data, get.imputed.data

has.boots

check whether a BNDataset has bootstrap samples or not.

Description

Return TRUE if the given dataset contains samples for bootstrap, FALSE otherwise.

Usage

```
has.boots(x)
## S4 method for signature 'BNDataset'
has.boots(x)
```

Arguments

Х

a BNDataset object.

Value

TRUE if dataset has bootstrap samples.

See Also

```
has.imp.boots, boots, imp.boots
```

has.data 23

has.data

check if a BNDataset contains any data.

Description

Check whether a BNDataset object actually contains raw or imputed data.

Usage

```
has.data(x)
## S4 method for signature 'BNDataset'
has.data(x)
```

Arguments

Х

a BNDataset.

See Also

```
has.raw.data, has.imputed.data, get.data, get.raw.data, get.imputed.data
```

Examples

```
## Not run:
x <- BNDataset()
has.data(x) # FALSE

x <- read.dataset(x, "file.header", "file.data")
has.data(x) # TRUE

## End(Not run)</pre>
```

has.imp.boots

check whether a BNDataset has bootstrap samples from imputed data or not.

Description

Return TRUE if the given dataset contains samples for bootstrap from inputed dataset, FALSE otherwise.

Usage

```
has.imp.boots(x)
## S4 method for signature 'BNDataset'
has.imp.boots(x)
```

24 has.imputed.data

Arguments

```
x a BNDataset object.
```

Value

TRUE if dataset has bootstrap samples from imputed data.

See Also

```
has.boots, boots, imp.boots
```

has.imputed.data

check if a BNDataset contains impited data.

Description

Check whether a BNDataset object actually contains imputed data.

Usage

```
has.imputed.data(x)
## S4 method for signature 'BNDataset'
has.imputed.data(x)
```

Arguments

x a BNDataset.

See Also

```
has.data, has.raw.data, get.data, get.raw.data, get.imputed.data
```

Examples

```
## Not run:
x <- BNDataset()
has.imputed.data(x) # FALSE

x <- read.dataset(x, "file.header", "file.data")
has.imputed.data(x) # FALSE, since read.dataset() actually reads raw data.

x <- impute(x)
has.imputed.data(x) # TRUE

## End(Not run)</pre>
```

has.raw.data 25

has.raw.data

check if a BNDataset contains raw data.

Description

Check whether a BNDataset object actually contains raw data.

Usage

```
has.raw.data(x)
## S4 method for signature 'BNDataset'
has.raw.data(x)
```

Arguments

Χ

a BNDataset.

See Also

has.data, has.imputed.data, get.data, get.raw.data, get.imputed.data

Examples

```
## Not run:
x <- BNDataset()
has.raw.data(x) # FALSE

x <- read.dataset(x, "file.header", "file.data")
has.raw.data(x) # TRUE, since read.dataset() actually reads raw data.
## End(Not run)</pre>
```

header.file

get header file of a BNDataset.

Description

Return the header filename of a dataset (with the path to its position, as given by the user), present if the dataset has been read from a file and not manually inserted. The header file contains three rows:

- 1. list of names of the variables, in the same order as in the data file;
- 2. list of cardinalities of the variables, if discrete, or levels for quantization if continuous;
- 3. list of status of the variables: c for continuous variables, d for discrete ones.

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Usage

```
header.file(x)
## S4 method for signature 'BNDataset'
header.file(x)
```

Arguments

x a BNDataset.

Value

header filename of the dataset.

See Also

data.file

header.file<-

set header file of a BNDataset.

Description

Set the header filename of a dataset (with the path to its position, as given by the user). The header file has to contain three rows:

- 1. list of names of the variables, in the same order as in the data file;
- 2. list of cardinalities of the variables, if discrete, or levels for quantization if continuous;
- 3. list of status of the variables: c for continuous variables, d for discrete ones.

Further rows are ignored.

Usage

```
header.file(x) <- value
```

Arguments

x a BNDataset. value header filename.

See Also

```
data.file<-
```

imp.boots 27

imp.boots

get list of bootstrap samples from imputed data of a BNDataset.

Description

Return the list of samples computed from raw data of a dataset.

Usage

```
imp.boots(x)
## S4 method for signature 'BNDataset'
imp.boots(x)
```

Arguments

Х

a BNDataset object.

Value

the list of bootstrap samples from imputed data.

See Also

```
has.boots, has.imp.boots, boots
```

imp.boots<-</pre>

set list of bootstrap samples from imputed data of a BNDataset.

Description

Add to a dataset a list of samples from imputed data computed using bootstrap.

Usage

```
imp.boots(x) \leftarrow value
```

Arguments

x a BNDataset object.

value the list of bootstrap samples from imputed data.

28 imputed.data<-

impute

Impute a BNDataset raw data with missing values.

Description

Impute a BNDataset raw data with missing values.

Usage

```
impute(object, ...)
## S4 method for signature 'BNDataset'
impute(object, k.impute = 10)
```

Arguments

object the BNDataset object.

k. impute number of neighbours to be used; for discrete variables we use mode, for con-

tinuous variables the median value is instead taken.

... potential further arguments of methods.

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
dataset <- impute(dataset)
## End(Not run)</pre>
```

imputed.data<-</pre>

add imputed data.

Description

Insert imputed data in a BNDataset object.

Usage

```
imputed.data(x) <- value
```

Arguments

```
x a BNDataset.
```

value a matrix of integers containing a dataset.

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Details

Users are encouraged to not use this method whenever possible, in favour of read.dataset with flag imputation = TRUE.

See Also

```
has.data, has.imputed.data, get.data, read.dataset
```

InferenceEngine-class InferenceEngine class.

Description

InferenceEngine class.

Constructor method of InferenceEngine class.

constructor for InferenceEngine object

Usage

```
## S4 method for signature 'InferenceEngine'
initialize(.Object, ...)

InferenceEngine(bn = NULL, observations = NULL, ...)
```

Arguments

. Object an empty InferenceEngine object.

bn a BN object.

observations a list of observations composed by the two following vectors:

- observed.vars:vector of observed variables;
- observed.vals:vector of values observed for the variables in observed.vars in the corresponding position.

... potential further arguments of methods.

Value

an InferenceEngine object.

InferenceEngine object.

jpts

Slots

```
junction.tree: junction tree adjacency matrix.
num.nodes: number of nodes in the junction tree.
cliques: list of cliques composing the nodes of the junction tree.
triangulated.graph: adjacency matrix of the original triangulated graph.
jpts: inferred joint probability tables.
bn: original Bayesian Network (as object of class BN) as provided by the user, or learnt from a dataset. NULL if missing.
updated.bn: Bayesian Network (as object of class BN) as modified by a belief propagation computation. In particular, it will have different conditional probability tables with respect to its original version. NULL if missing.
observed.vars: list of observed variables, by name or number.
observed.vals: list of observed values for the corresponding variables in observed.vars.
```

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
bn <- BN(dataset)
eng <- InferenceEngine(bn)

obs <- list(c("A","G,"X),c(1,2,1))
eng.2 <- InferenceEngine(bn, obs)

## End(Not run)</pre>
```

jpts get the list of joint probability tables compiled by an InferenceEngine.

Description

Return the list of joint probability tables for the cliques of the junction tree obtained after belief propagation has been performed.

Usage

```
jpts(x)
## S4 method for signature 'InferenceEngine'
jpts(x)
```

Arguments

x an InferenceEngine.

jpts<-

Details

Each joint probability table is represented as a multidimensional array. To retrieve single dimensions (e.g. to compute marginals), users should not rely on dimension numbers, but should instead select the dimensions using their names.

Value

the list of joint probability tables compiled by the InferenceEngine.

```
jpts<- set the list of joint probability tables compiled by an
InferenceEngine.
```

Description

Add a list of joint probability tables for the cliques of the junction tree.

Usage

```
jpts(x) <- value</pre>
```

Arguments

```
x an InferenceEngine.value the list of joint probability tables compiled by the InferenceEngine.
```

Details

Each joint probability table is represented as a multidimensional array. To retrieve single dimensions (e.g. to compute marginals), users should provide dimension names.

```
jt.cliques get the list of cliques of the junction tree of an InferenceEngine.
```

Description

Return the list of cliques containing the variables associated to each node of a junction tree.

Usage

```
jt.cliques(x)
## S4 method for signature 'InferenceEngine'
jt.cliques(x)
```

32 junction.tree

Arguments

x an InferenceEngine.

Value

the list of cliques of the junction tree contained in the InferenceEngine.

jt.cliques<-

set the list of cliques of the junction tree of an InferenceEngine.

Description

Add to the InferenceEngine a list containing the cliques of variables composing the nodes of the junction tree.

Usage

```
jt.cliques(x) \leftarrow value
```

Arguments

x an InferenceEngine.

value the list of cliques of the junction tree contained in the InferenceEngine.

junction.tree

get the junction tree of an InferenceEngine.

Description

Return the adjacency matrix representing the junction tree computed for a network.

Usage

```
junction.tree(x)
## S4 method for signature 'InferenceEngine'
junction.tree(x)
```

Arguments

x an InferenceEngine.

Details

Rows and columns are named after the (variables in the) cliques that each node of the junction tree represent.

junction.tree<-

Value

the junction tree contained in the InferenceEngine.

See Also

```
build.junction.tree
```

junction.tree<-

set the junction tree of an InferenceEngine.

Description

Set the adjacency matrix of the junction tree computed for a network.

Usage

```
junction.tree(x) <- value
```

Arguments

x an InferenceEngine.

value the junction tree to be inserted in the InferenceEngine.

layering

return the layering of the nodes.

Description

Compute the topological ordering of the nodes of a network, in order to divide the network in layers.

Usage

```
layering(x, ...)
## S4 method for signature 'BN'
layering(x, ...)
## S4 method for signature 'InferenceEngine'
layering(x, updated.bn = TRUE, ...)
```

Arguments

a BN or InferenceEngine object.

updated.bn TRUE if x is an InferenceEngine and the updated network is chosen (kept only

for compatibility with other methods).

. . . potential further arguments for methods.

34 learn.params

Value

a vector containing layers the nodes can be divided into.

Examples

```
## Not run:
dataset <- BNDataset(name="MyDataset")
dataset <- read.dataset(dataset, "file.header", "file.data")
x <- BN(dataset)
layering(x)
eng <- InferenceEngine(x)
layering(x, updated.bn=TRUE)
## End(Not run)</pre>
```

learn.params

learn the parameters of a BN.

Description

Learn the parameters of a BN object according to a BNDataset using MAP (Maximum A Posteriori) estimation.

Usage

```
learn.params(bn, dataset, ...)
## S4 method for signature 'BN,BNDataset'
learn.params(bn, dataset, ess = 1)
```

Arguments

```
bn a BN object.
dataset a BNDataset object.
ess Equivalent Sample Size value.
```

... potential further arguments of methods.

Value

new BN object with conditional probabilities.

learn.structure 35

Examples

```
## Not run:
## first create a BN and learn its structure from a dataset
dataset <- BNDataset(name = "MyDataset")
dataset <- read.dataset(dataset, "file.header", "file.data")
bn <- BN()
bn <- learn.structure(bn, dataset)
bn <- learn.params(bn, dataset, ess=1)
## End(Not run)</pre>
```

learn.structure

learn the structure of a network.

Description

Learn the structure (the directed acyclic graph) of a BN object according to a BNDataset. Currently, two algorithms are supported (can be specified using the algo option): 'sm', the Silander-Myllymaki exact algorithm, and 'mmhc', the Max-Min Hill-Climbing heuristic algorithm (default). Three scoring functions are also provided: 'BDeu', the Bayesian-Dirichlet equivalent uniform score, 'AIC', the Akaike Information criterion, and 'BIC', the Bayesian Information criterion.

Usage

```
learn.structure(bn, dataset, ...)
## S4 method for signature 'BN,BNDataset'
learn.structure(bn, dataset, algo = "mmhc",
    scoring.func = "BDeu", alpha = 0.05, ess = 1, bootstrap = FALSE,
    layering = c(), max.fanin.layers = NULL,
    max.fanin = num.variables(dataset), cont.nodes = c(), raw.data = FALSE,
    num.boots = 100, imputation = TRUE, k.impute = 10,
    na.string.symbol = "?", seed = 0)
```

Arguments

bn a BN object. dataset a BNDataset.

algo the algorithm to use. Currently, one among sm (Silander-Myllymaki) and mmhc

(Max-Min Hill Climbing, default).

scoring. func the scoring function to use. Currently, one among BDeu, AIC, BIC.

alpha confidence threshold (only for mmhc).

ess Equivalent Sample Size value.
bootstrap TRUE to use bootstrap samples.

num.boots number of bootstrap samples to generate, if needed.

36 learn.structure

layering vector containing the layers each node belongs to (only for sm). max.fanin.layers matrix of available parents in each layer (only for sm). max.fanin maximum number of parents for each node (only for sm). cont.nodes use an empty vector. raw.data TRUE to learn the structure from the raw dataset. Default is to use imputed dataset (if available, otherwise the raw dataset will be used anyway). imputation TRUE if imputation is needed; if bootstrap=TRUE, imputed samples will be also used. na.string.symbol symbol for NA values (missing data). k.impute number of neighbours to be used; for discrete variables we use mode, for continuous variables the median value is instead taken. seed random seed. potential further arguments of methods.

Details

The Silander-Myllymaki algorithm can take a very long time, and it is not feasible for networks of more than 20-30 nodes. It is strongly recommended that valid layering, max.fanin.layers and max.fanin parameters are passed to the method if algo = 'sm' is given as parameter to the method.

Value

new BN object with DAG.

Examples

marginals 37

marginals

compute the list of inferred marginals of a BN.

Description

Given an InferenceEngine, it returns a list containing the marginals for the variables in the network, according to the propagated beliefs.

Usage

```
marginals(x, ...)
## S4 method for signature 'InferenceEngine'
marginals(x, ...)
```

Arguments

x an InferenceEngine
... potential further arguments of methods.

Value

a list containing the marginals of each variable, as probability tables.

Examples

```
## Not run:
eng <- InferenceEngine(net)
marginals(eng)
## End(Not run)</pre>
```

name

get name of an object.

Description

Return the name of an object, of class BN or BNDataset.

Usage

```
name(x)
## S4 method for signature 'BNDataset'
name(x)
## S4 method for signature 'BN'
name(x)
```

38 node.sizes

Arguments

x an object.

Value

name of the object.

name<-

set name of an object.

Description

Set the name slot of an object of type BN or BNDataset.

Usage

```
name(x) \leftarrow value
```

Arguments

x an object.

value the new name of the object.

node.sizes

get size of the variables of an object.

Description

Return a list containing the size of the variables of an object. It is the actual cardinality of discrete variables, and the cardinality of the discretized variable for continuous variables.

Usage

```
node.sizes(x)
## S4 method for signature 'BNDataset'
node.sizes(x)
## S4 method for signature 'BN'
node.sizes(x)
```

Arguments

Χ

an object.

Value

vector contaning the size of each variable of the desired object.

node.sizes<-

node.sizes<-

set the size of variables of an object.

Description

Set the size of the variables of a BN or BNDataset object. It represents the actual cardinality of discrete variables, and the cardinality of the discretized variable for continuous variables.

Usage

```
node.sizes(x) \leftarrow value
```

Arguments

x an object.

value vector containing the size of each variable of the object.

num.boots

get number of bootstrap samples of a BNDataset.

Description

Return the number of bootstrap samples computed from a dataset.

Usage

```
num.boots(x)
## S4 method for signature 'BNDataset'
num.boots(x)
```

Arguments

x a BNDataset object.

Value

the number of bootstrap samples.

40 num.items

num.boots<-

set number of bootstrap samples of a BNDataset.

Description

Set the length of the list of samples of a dataset computed using bootstrap.

Usage

```
num.boots(x) \leftarrow value
```

Arguments

x a BNDataset object.value the number of bootstrap samples.

num.items

get number of items of a BNDataset.

Description

Return the number of items in a dataset, that is, the number of rows in its data slot.

Usage

```
num.items(x)
## S4 method for signature 'BNDataset'
num.items(x)
```

Arguments

x a BNDataset object.

Value

number of items of the desired dataset.

num.items<-

num.items<-

set number of items of a BNDataset.

Description

Set the number of observed items (rows) in a dataset.

Usage

```
num.items(x) \leftarrow value
```

Arguments

x a BNDataset object.

value number of items of the desired dataset.

num.nodes

get number of nodes of an object.

Description

Return the name of an object, of class BN or InferenceEngine.

Usage

```
num.nodes(x)
## S4 method for signature 'BN'
num.nodes(x)
## S4 method for signature 'InferenceEngine'
num.nodes(x)
```

Arguments

Χ

an object.

Value

number of nodes of the desired object.

42 num.variables

num.nodes<-

set number of nodes of an object.

Description

Set the number of nodes of an object of type BN (number of nodes of the network) or InferenceEngine (where parameter contains the number of nodes of the junction tree).

Usage

```
num.nodes(x) <- value
```

Arguments

x an object.

value the number of nodes in the object.

num.variables

get number of variables of a BNDataset.

Description

Return the number of the variables contained in a dataset. This value corresponds to the value of num. nodes of a network built upon the same dataset.

Usage

```
num.variables(x)
## S4 method for signature 'BNDataset'
num.variables(x)
## S4 method for signature 'BNDataset'
num.variables(x)
```

Arguments

x a BNDataset object.

Value

number of variables of the desired dataset.

See Also

num.nodes

num.variables<- 43

num.variables<-

set number of variables of a BNDataset.

Description

Set the number of variables observed in a dataset.

Usage

```
num.variables(x) <- value</pre>
```

Arguments

x a BNDataset object.

value number of variables of the dataset.

observations

get the list of observations of an InferenceEngine.

Description

Return the list of observations added to an InferenceEngine.

Usage

```
observations(x)
## S4 method for signature 'InferenceEngine'
observations(x)
```

Arguments

Χ

an InferenceEngine.

Details

Output is a list in the following format:

- observed.varsvector of observed variables;
- observed.valsvector of values observed for the variables in observed.vars in the corresponding position.

Value

the list of observations of the InferenceEngine.

44 plot

observations<-

set the list of observations of an InferenceEngine.

Description

Add a list of observations to an InferenceEngine, using a list of observations composed by the two following vectors:

- observed.varsvector of observed variables;
- observed.valsvector of values observed for the variables in observed.vars in the corresponding position.

Usage

```
observations(x) <- value
```

Arguments

```
x an InferenceEngine.value the list of observations of the InferenceEngine.
```

Details

Replace previous list of observations, if present. In order to add evidence, and not just replace it, one must use the add.observations<- method.

In case of multiple observations of the same variable, the last observation is the one used, as the most recent.

See Also

```
add.observations<-
```

plot

plot a BN as a picture.

Description

Plot the network as a picture to default output.

Usage

```
## S4 method for signature 'BN'
plot(x, use.node.names = TRUE, frac = 0.2,
   max.weight = max(dag(x)), node.col = rep("white", ncol(dag(x))),
   plot.wpdag = FALSE)
```

print 45

Arguments

```
x a BN object.

use.node.names TRUE if node names have to be printed. If FALSE, number are used instead.

frac fraction

max.weight max.weight

node.col list of (R) colors for the nodes.

plot.wpdag if TRUE plot the network according to the WPDAG computed using bootstrap instead of the DAG.

... potential further arguments of methods.
```

Examples

print

print an object to stdout.

Description

print an object to stdout.

Usage

```
print(x, ...)
## S4 method for signature 'BNDataset'
print(x, show.raw.data = FALSE,
    show.imputed.data = FALSE, ...)
## S4 method for signature 'BN'
print(x, ...)
## S4 method for signature 'InferenceEngine'
print(x, engine = "jt", ...)
```

Arguments

```
x an object.

show.raw.data when x is a BNDataset, print also raw dataset, if available.

show.imputed.data when x is a BNDataset, print also imputed dataset, if available.
```

46 raw.data<-

engine when x is an InferenceEngine, specify the inference engine to be shown. Currently only engine = 'jt' is supported.

... potential other arguments.

query

query BN given observations

Description

```
query BN given observations
```

Usage

```
query(x, ...)
## S4 method for signature 'BN'
query(x, observed.vars, observed.vals)
```

Arguments

```
    x a BN.
    observed.vars vector of observed variables.
    observed.vals vector of observed values for corresponding variables in observed.vars.
    potential further arguments for method.
```

Value

most probable values given observations

raw.data<-

add raw data.

Description

Insert raw data in a BNDataset object.

Usage

```
raw.data(x) <- value</pre>
```

Arguments

x a BNDataset.

value a matrix of integers containing a dataset.

read.dataset 47

Details

Users are encouraged to not use this method whenever possible, in favour of read.dataset.

See Also

```
has.data, has.raw.data, get.data, read.dataset
```

read.dataset

Read a dataset from file.

Description

File has to be in format (describe...)

Usage

```
read.dataset(object, header.file, data.file, ...)
## S4 method for signature 'BNDataset,character,character'
read.dataset(object, header.file,
   data.file, imputation = FALSE, header.flag = FALSE,
   na.string.symbol = "?", sep.symbol = "", k.impute = 10,
   bootstrap = FALSE, num.boots = 100, seed = 0, ...)
```

Arguments

object the BNDataset object. header.file the header file. data.file the data file. na.string.symbol character that denotes NA in the dataset. sep.symbol separator among values in the dataset. header.flag TRUE if the first row of dataset file is an header (e.g. it contains the variable names). imputation TRUE if imputation has to be performed. k.impute number of neighbours to be used; for discrete variables we use mode, for continuous variables the median value is instead taken (useful only if imputation == TRUE). bootstrap TRUE if bootstrap has to be performed; prepares a list of datasets sampled from the original one. number of sampled datasets for bootstrap (useful only if bootstrap == TRUE). num.boots seed random seed (useful only if bootstrap == TRUE). potential further arguments of methods.

48 save.to.eps

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, header="file.header", dataset="file.data")
## End(Not run)</pre>
```

save.to.eps

save a BN picture as .eps file.

Description

Save an image of a Bayesian Network as an .eps file.

Usage

```
save.to.eps(x, filename)
## S4 method for signature 'BN,character'
save.to.eps(x, filename)
```

Arguments

```
x a BN object
filename name (with path, if needed) of the file to be created
```

See Also

plot

Examples

```
## Not run:
save.to.eps(x, "out.eps")
## End(Not run)
```

test.updated.bn 49

test.updated.bn

check if an updated BN is present in an InferenceEngine.

Description

Check if an InferenceEngine actually contains an updated network, in order to provide the chance of a fallback and use the original network if no belief propagation has been performed.

Usage

```
test.updated.bn(x)
## S4 method for signature 'InferenceEngine'
test.updated.bn(x)
```

Arguments

.,

an InferenceEngine.

Value

TRUE if an updated network is contained in the InferenceEngine, FALSE otherwise.

Examples

```
## Not run:
dataset <- BNDataset()
dataset <- read.dataset(dataset, "file.header", "file.data")
bn <- BN(dataset)
ie <- InferenceEngine(bn)
test.updated.bn(ie) # FALSE

observations(ie) <- list("observed.vars"=("A","G","X"), "observed.vals"=c(1,2,1))
ie <- belief.propagation(ie)
test.updated.bn(ie) # TRUE

## End(Not run)</pre>
```

updated.bn

get the updated BN object contained in an InferenceEngine.

Description

Return an updated network contained in an InferenceEngine.

50 variables

Usage

```
updated.bn(x)
## S4 method for signature 'InferenceEngine'
updated.bn(x)
```

Arguments

x an InferenceEngine.

Value

the updated BN object contained in an InferenceEngine.

updated.bn<-

set the updated BN object contained in an InferenceEngine.

Description

Add an updated network to an InferenceEngine.

Usage

```
updated.bn(x) \leftarrow value
```

Arguments

x an InferenceEngine.value the updated BN object contained in an InferenceEngine.

variables

get variables of an object.

Description

Get the list of variables (with their names) of a BN or BNDataset.

Usage

```
variables(x)
## S4 method for signature 'BNDataset'
variables(x)
## S4 method for signature 'BN'
variables(x)
```

variables<- 51

Arguments

x an object.

Value

vector of the variables names of the desired object.

variables<-

set variables of an object.

Description

Set the list of variable names in a BN or BNDataset object.

Usage

```
variables(x) \leftarrow value
```

Arguments

x an object.

value vector containing the variable names of the object. Overwrites num.nodes slot

if non-matching.

wpdag

get the WPDAG of an object.

Description

Return the weighted partially directed acyclic graph of a network, when available (e.g. when bootstrap on dataset is performed).

Usage

```
wpdag(x)
## S4 method for signature 'BN'
wpdag(x)
```

Arguments

an object.

Value

matrix contaning the WPDAG of the object.

52 wpdag<-

wpdag<-

set WPDAG of the object.

Description

Set the weighted partially directed acyclic graph of a network (e.g. in case bootstrap on dataset is performed).

Usage

```
wpdag(x) \leftarrow value
```

Arguments

x an object.

value matrix containing the WPDAG of the object.

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