

Package: r.blip (via r-universe)

September 11, 2024

Title R binding of blip (Bayesian network Learning Improved Project)

Version 1.1

Description Offers many approaches to score-based structure learning of Bayesian networks.

Depends R (>= 3.0.0)

Imports foreign, bnlearn (>= 4.0)

SystemRequirements Java (>= 1.5)

License GPLv3

Encoding UTF-8

LazyData true

RoxxygenNote 6.1.1

Repository <https://mauro-idsia.r-universe.dev>

RemoteUrl <https://github.com/mauro-idsia/r.blip>

RemoteRef HEAD

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blip*Bayesian Learning Package - Main function.***Description**

Used by most of the functions in the r.blip binding, provides access to the included jar file.

Usage

```
blip(args)
```

Arguments

args	Vector of arguments to be passed to the jar
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Details

The arguments vector is formatted in a system call to the included jar file. Should not be called directly by the user, unless you know exactly what you are doing. In that case, call directly the blip jar.

blip.learn*Learns a BN***Description**

Fully learns a Bayesian networks.

Usage

```
blip.learn(dat, scorer.method = "is", solver.method = "winasobs",
           indeg = 6, time = 3600, allocated = 80, scorefunction = "bic",
           alpha = 1, cores = 1, verbose = 0)
```

Arguments

dat	dataframe from which to learn the parent sets.(required)
scorer.method	Method to be used for scoring the parent sets. Possible values: "is" (independence selection), "sq" (sequential selection). (default: is)
solver.method	Method to be used for structure exploration. Possible values: "winasobs", "winobs", "asobs", "obs". (default: winasobs)
indeg	Maximum number of parents (default: 6)
time	Execution time (default: 3600)
allocated	Percentage of the total execution time dedicated to parent set exploration (default: 80)

<code>scorefunction</code>	Chosen score function. Possible choices: BIC, BDeu (default: bic)
<code>alpha</code>	(if BDeu is chosen) equivalent sample size parameter (default: 1.0)
<code>cores</code>	Number of machine cores to use. If 0, all are used. (default: 1)
<code>verbose</code>	Verbose level (default: 0)

Details

The input data is required to be complete and discrete. Accordingly missing values in the input `data.frame` will be ignored, and all numeric values will be converted to integers.

Value

The learned Bayesian network in the bnlearn format.

Examples

```
bn <- blip.learn(read.table('data/child-5000.dat', sep = ' '), time = 10)
```

<code>blip.learn.tw</code>	<i>Learns a BN with a treewidth bound</i>
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Description

Fully learns a Bayesian networks with a treewidth bound.

Usage

```
blip.learn.tw(dat, scorer.method = "is", solver.method = "kmax",
  treewidth = 4, time = 3600, allocated = 80,
  scorefunction = "bic", alpha = 1, cores = 1, verbose = 0)
```

Arguments

<code>dat</code>	dataframe from which to learn the parent sets.(required)
<code>scorer.method</code>	Method to be used for scoring the parent sets. Possible values: "is" (independence selection), "sq" (sequential selection). (default: is)
<code>solver.method</code>	Method to be used for bounded-treewidth structure exploration. Possible values: "kmax", "kg", "ka". (default: kmax)
<code>treewidth</code>	Maximum treewidth (default: 4)
<code>time</code>	Execution time (default: 3600)
<code>allocated</code>	Percentage of the total execution time dedicated to parent set exploration (default: 80)
<code>scorefunction</code>	Chosen score function. Possible choices: BIC, BDeu (default: bic)
<code>alpha</code>	(if BDeu is chosen) equivalent sample size parameter (default: 1.0)
<code>cores</code>	Number of machine cores to use. If 0, all are used. (default: 1)
<code>verbose</code>	Verbose level (default: 0)

Details

The input data is required to be complete and discrete. Accordingly missing values in the input data.frame will be ignored, and all numeric values will be converted to integers.

Value

The learned Bayesian network in the bnlearn format.

<code>blip.scorer</code>	<i>Parent set exploration</i>
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Description

Generates the cache of parent sets from a given data source

Usage

```
blip.scorer(dat, method = "is", indeg = 6, time = 3600,
            scorefunction = "bic", alpha = 1, cores = 1, verbose = 1)
```

Arguments

<code>dat</code>	dataframe from which to learn the parent sets.(required)
<code>method</code>	Method to be used. Possible values: "is" (independence selection), "sq" (sequential selection). (default: is)
<code>indeg</code>	Maximum number of parents (default: 6)
<code>time</code>	Maximum Execution time (default: 3600)
<code>scorefunction</code>	Chosen score function. Possible choices: BIC, BDeu (default: bic)
<code>alpha</code>	(if BDeu is chosen) equivalent sample size parameter (default: 1.0)
<code>cores</code>	Number of machine cores to use. If 0, all are used. (default: 1)
<code>verbose</code>	Verbose level (default: 0)

Details

Usually the first step in the learning of a Bayesian network.

The input data is required to be complete and discrete. Accordingly missing values in the input data.frame will be ignored, and all numeric values will be converted to integers.

Value

Cache of parent sets

blip.solver *Structure Optimization*

Description

Find an optimal structure from the cache of parent sets

Usage

```
blip.solver(jkl, method = "winasobs", time = 3600, cores = 1,  
verbose = 1)
```

Arguments

jk1	cache of pre-computed parent sets.(required)
method	Method to be used. Possible values: "winasobs", "winobs", "asobs", "obs". (default: winasobs)
time	Maximum Execution time (default: 3600)
cores	Number of machine cores to use. If 0, all are used. (default: 1)
verbose	Verbose level (default: 0)

Details

The input data is required to be complete and discrete. Accordingly missing values in the input data.frame will be ignored, and all numeric values will be converted to integers.

Value

Structure

blip.solver.tw *Structure Optimization - treewidth bound*

Description

Find an optimal structure from the cache of parent sets

Usage

```
blip.solver.tw(jkl, method = "kmax", treewidth = 4, time = 3600,  
cores = 1, verbose = 1)
```

Arguments

<code>jk1</code>	cache of pre-computed parent sets.(required)
<code>method</code>	Method to be used. Possible values: "kmax", "kg", "ka". (default: kmax)
<code>treewidth</code>	Maximum treewidth (default: 4)
<code>time</code>	Maximum Execution time (default: 3600)
<code>cores</code>	Number of machine cores to use. If 0, all are used. (default: 1)
<code>verbose</code>	Verbose level (default: 0)

Details

The input data is required to be complete and discrete. Accordingly missing values in the input `data.frame` will be ignored, and all numeric values will be converted to integers.

Value

`Structure`

<code>read.jkl</code>	<i>Read a Jkl file (parent sets cache)</i>
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Description

Read a Jkl file (parent sets cache)

Usage

```
read.jkl(path, data)
```

<code>read.str</code>	<i>Read a res file for bnlearn loading</i>
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Description

Read a res file for bnlearn loading

Usage

```
read.str(path, data)
```

`write.jkl`

Write a Jkl file (parent sets cache)

Description

Write a Jkl file (parent sets cache)

Usage

`write.jkl(path, jkl, data)`

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