

Package ‘AgroBayes’

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Type Package

Title Generates static and dynamic Bayesian networks for forecasting crop results

Version 0.1.0

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Description This package permits, from variables related to agricultural production, to generate static and dynamic Bayesian networks that allow forecasting crop results. The package allows comparing the effectiveness of models generated from some network structure learning methods. The networks are generated based on functions available in the bnlearn (static networks) and dbnR (dynamic networks) packages. The forecasting of the crop production can be done from sets of data geographically separated (specific areas of the plantation) and also from chronological cuts (phenological phases).

All functions identified with the term TEST FUNCTION in the documentation and with a name starting with 'test', were added to the package to run the demonstration of the package's functionalities. The demo.Rmd provides an interactive example of how the package works.

The final user must organize the dataset in dataframes representing the interval referring to a phenological phase and group them in lists according to the area of the plantation site.

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 7.2.3

Suggests knitr,
rmarkdown,
testthat (>= 3.0.0)

VignetteBuilder knitr

Config/testthat/edition 3

Imports bnlearn, dbnR, Hmisc, dplyr, caTools, ggplot2, data.table, caret, stats, graphics, grDevices

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createDbn	<i>Creates dynamic Bayesian networks for performance evaluation</i>
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Description

Using harvest data from all phenological phases of the cultivar, from a specific area of the plantation, dynamic Bayesian networks are generated (using the `dbnR::learn_dbn_struct` functions with `natPsoho` and `dmmhc` methods), trained (using the `dbnR::fit_dbn_params` function) and evaluated for performance (using `dbnR::forecast_ts` and `caret::defaultSummary`). Two networks are created, one using `natPsoho` learning method and other using `dmmhc` method. The DBN are tested using `dbnR::forecast_ts` with exact and approx methods then, from the return of these functions, the RMSE, R-squared and MAE metrics are calculated and returned

Usage

```
createDbn(area)
```

Arguments

area A dataframe of continuous data, from a specific area of the plantation to be used

Value

A dataframe with the dynamics networks metrics (RMSE, R-squared, MAE)

Examples

```
metricsDbn = createDbn(area_1)
```

createNetworks	<i>Creates Bayesian networks for performance evaluation</i>
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Description

Using harvest data from a phenological phase of the cultivar, from a specific area of the plantation, Bayesian network are generated (using the `bnlearn::hc` and `bnlearn::mmhc` functions), trained (using the `bnlearn::bn.fit` function) and evaluated for performance (using [validateNetwork](#)). Four networks are created, two from the pre-established topology and two learned only from the presented data.

Usage

```
createNetworks(areaphase, blacklist, whitelist)
```

Arguments

areaphase	the dataframe of discretized data to be used
blacklist	a dataframe of character string with two columns, it is passed as a parameter to bnlearn learn functions in order to avoid these arcs composing the final network
whitelist	a dataframe of character string with two columns, it is passed as a parameter to bnlearn learn functions in order to guarantee these arcs composing the final network

Value

Network evaluation metrics, as calculated in the validateNetwork function

Examples

```
metricsArea1Phase1 = list()
blacklist = data.frame(
  from = c("X_1", "X_1",
           "X_2", "X_2",
           "X_3", "X_3",
           "harvest", "harvest", "harvest" ),
  to = c("X_2", "X_3", #from v1
         "X_1", "X_3", #from v2
         "X_1", "X_2", #from v3
         "X_1", "X_2", "X_3")) #from col
whitelist = data.frame(
  from = c("X_1", "X_2", "X_3"),
  to = c("harvest", "harvest", "harvest"))
areaphase = data.frame(area1_phase_1)
metricsArea1Phase1 = createNetworks (areaphase, blacklist, whitelist)
```

runNetworks

Executes [createNetworks](#) function to an area

Description

Executes the [createNetworks](#) function for all phenological phases of an area. It also organizes the generated metrics in a dataframe.

Usage

```
runNetworks(arealist, blacklist, whitelist)
```

Arguments

arealist	list of dataframes of all phenological phase on a specific area of the plantation
blacklist	a dataframe of character string with two columns, it is passed as a parameter to bnlearn learn functions in order to avoid these arcs composing the final network
whitelist	a dataframe of character string with two columns, it is passed as a parameter to bnlearn learn functions in order to guarantee these arcs composing the final network

Value

A dataframe organizing the network metrics generated in the `createNetworks` function. The lines represent the network performance learned in each phenological phase of the area (arealist)

Examples

```
blacklist = data.frame(
  from = c("X_1", "X_1",
           "X_2", "X_2",
           "X_3", "X_3",
           "harvest", "harvest", "harvest" ),
  to = c("X_2", "X_3", #from v1
         "X_1", "X_3", #from v2
         "X_1", "X_2", #from v3
         "X_1", "X_2", "X_3")) #from harvest
whitelist = data.frame(
  from = c("X_1", "X_2", "X_3"),
  to = c("harvest", "harvest", "harvest"))
arealist <- list(areal)
metricsArea1 <- createNetworks (arealist, blacklist, whitelist)
```

 validateNetwork

Generates Bayesian networks performance evaluation

Description

Using the functions available in the repository <https://github.com/KaikeWesleyReis/bnlearn-multivar-prediction>, calculates the metrics of the four Bayesian networks generated in the `createNetworks` function when executed in context of `runNetworks` function.

Usage

```
validateNetwork(
  test,
  train,
  dag_fitted1,
  dag_fitted2,
```

```
    dag_fitted3,  
    dag_fitted4  
  )
```

Arguments

test	dataframe to be used to test the Bayesian networks. It is composed of a 25 the createNetworks function.
train	dataframe to be used to train the Bayesian networks. It is composed of a 75 the createNetworks function.
dag_fitted1	Fitted Bayesian network to be tested
dag_fitted2	Fitted Bayesian network to be tested
dag_fitted3	Fitted Bayesian network to be tested
dag_fitted4	Fitted Bayesian network to be tested

Value

List of values returned from bnMetricsMultiVarPrediction. See more in <https://github.com/KaikeWesleyReis/bnlearn-multivar-prediction-metrics#bnmetricsmultivarprediction>

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