Package: rCBA (via r-universe)

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Title CBA Classifier

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URL https://github.com/jaroslav-kuchar/rCBA

BugReports https://github.com/jaroslav-kuchar/rCBA/issues

Description Provides implementations of a classifier based on the ``Classification Based on Associations" (CBA). It can be used for building classification models from association rules. Rules are pruned in the order of precedence given by the sort criteria and a default rule is added. The final classifier labels provided instances. CBA was originally proposed by Liu, B. Hsu, W. and Ma, Y. Integrating Classification and Association Rule Mining. Proceedings KDD-98, New York, 27-31 August. AAAI. pp80-86 (1998, ISBN:1-57735-070-7).

Depends R (>= 3.1.3), rJava, arules

Imports R.utils, TunePareto, methods, stats, utils

License Apache License (== 2.0)

LazyData true

SystemRequirements Java (>= 8)

RoxygenNote 6.1.1

Encoding UTF-8

Collate 'init.R' 'build.R' 'buildFPGrowth.R' 'classification.R' 'fpgrowth.R' 'pruning.R' 'utils.R'

NeedsCompilation no

Config/pak/sysreqs default-jdk

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`build` *Build classifier function (Apriori-based)*

Description

Automatic build of the classification model using the Apriori algorithm from the `arules`

Usage

```
build(trainData, className = NA, pruning = TRUE, sa = list(),
  verbose = TRUE, parallel = TRUE)
```

Arguments

`trainData`	`data.frame` or `transactions` from `arules` with input data
`className`	column name with the target class - default is the last column
`pruning`	performing pruning while building the model
`sa`	simulated annealing setting. Default values: list(temp=100.0, alpha=0.05, tabuRuleLength=5, timeout=10)
`verbose`	verbose indicator
`parallel`	parallel indicator

Value

list with parameters and model as data.frame with rules

Examples

```
library("rCBA")
data("iris")

output <- rCBA::build(iris,sa = list(alpha=0.5), parallel=FALSE) # speeding up the cooling
model <- output$model

predictions <- rCBA::classification(iris, model)
table(predictions)
sum(as.character(iris$Species)==as.character(predictions), na.rm=TRUE) / length(predictions)
```

buildFPGrowth *Build classifier function (FP-Growth-based)*

Description

Automatic build of the classification model using the FP-Growth algorithm

Usage

```
buildFPGrowth(train, className = NULL, verbose = TRUE,
  parallel = TRUE)
```

Arguments

train	data.frame or transactions from arules with input data
className	column name with the target class - default is the last column
verbose	verbose indicator
parallel	parallel indicator

Value

list with parameters and model as data.frame with rules

Examples

```
library("rCBA")
data("iris")

output <- rCBA::buildFPGrowth(iris[sample(nrow(iris), 10),], "Species",
 parallel=FALSE, verbose=TRUE)
inspect(output$model)
```

classification *A classification function*

Description

A classification function

Usage

```
classification(test, rules, verbose = TRUE)
```

Arguments

test	data.frame or transactions from arules with input data
rules	data.frame with rules
verbose	verbose indicator

Value

vector with classifications

Examples

```
library("arules")
library("rCBA")
data("iris")

train <- sapply(iris, as.factor)
train <- data.frame(train, check.names=FALSE)
txns <- as(train,"transactions")

rules = apriori(txns, parameter=list(support=0.03, confidence=0.03, minlen=2),
appearance = list(rhs=c("Species=setosa", "Species=versicolor", "Species=virginica"),default="lhs"))

predictions <- rCBA::classification(train,rules)
table(predictions)
sum(as.character(train$Species)==as.character(predictions),na.rm=TRUE)/length(predictions)
```

---

fpgrowth *FP-Growth*

Description

FP-Growth algorithm - Jiawei Han, Jian Pei, and Yiwen Yin. Mining frequent patterns without candidate generation. SIGMOD Rec. 29, 2 (2000) <doi:10.1145/335191.335372>

Usage

```
fpgrowth(train, support = 0.01, confidence = 1, maxLength = 5,
  consequent = NULL, verbose = TRUE, parallel = TRUE)
```

Arguments

train	data.frame or transactions from arules with input data
support	minimum support
confidence	minimum confidence
maxLength	maximum length
consequent	filter consequent - column name with consequent/target class
verbose	verbose indicator
parallel	parallel indicator

Examples

```
library("rCBA")
data("iris")

train <- sapply(iris,as.factor)
train <- data.frame(train, check.names=FALSE)
txns <- as(train,"transactions")

rules = rCBA::fpgrowth(txns, support=0.03, confidence=0.03, maxLength=2, consequent="Species",
           parallel=FALSE)

predictions <- rCBA::classification(train,rules)
table(predictions)
sum(as.character(train$Species)==as.character(predictions),na.rm=TRUE)/length(predictions)

prunedRules <- rCBA::pruning(train, rules, method="m2cba", parallel=FALSE)
predictions <- rCBA::classification(train, prunedRules)
table(predictions)
sum(as.character(train$Species)==as.character(predictions),na.rm=TRUE)/length(predictions)
```

---

frameToRules *Conversion of* data.frame *to* rules *from* arules

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Description

Conversion of data.frame to rules from arules

Usage

```
frameToRules(model)
```

Arguments

model	data.frame with rules

Value

arules rules representation

Examples

```
library("rCBA")

model <- data.frame("rules" = c("{X=1} => {Y=1}","{X=0} => {Y=0}"),
                    "support" = c(0.5,0.5),
                    "confidence" = c(0.5,0.5),
                    "lift" = c(1.0,1.0))

rules <- rCBA::frameToRules(model)
```

```
inspect(rules)
```

---

pruning *A Pruning function*

Description

A Pruning function

Usage

```
pruning(train, rules, method = "m2cba", verbose = TRUE,
  parallel = TRUE)
```

Arguments

train	trainData data.frame or transactions from arules with input data
rules	data.frame with rules
method	pruning method m2cba(default)|m1cba|dcbrcba
verbose	verbose indicator
parallel	parallel indicator

Value

data.frame with pruned rules

Examples

```
library("arules")
library("rCBA")
data("iris")

train <- sapply(iris,as.factor)
train <- data.frame(train, check.names=FALSE)
txns <- as(train,"transactions")

rules = apriori(txns, parameter=list(support=0.03, confidence=0.03, minlen=2),
appearance = list(rhs=c("Species=setosa", "Species=versicolor", "Species=virginica"),default="lhs"))

print(length(rules))
prunedRules <- rCBA::pruning(train, rules, method="m2cba", parallel=FALSE)
print(length(prunedRules))
```

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