# Data structures for association rules implementation in the R package **arules**

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#### 1 Introduction

Mining of frequent itemsets or association rules is a popular method for discovering interesting relations between variables in large databases. Apriori and Eclat (Agrawal, Imielinski, and Swami, 1993; Agrawal and Srikant, 1994) are two algorithms for fast evaluation of frequent itemsets and association rules given certain threshold values.

The R package **arules** provides an interface to the C programs by Christian Borgelt which implement the two algorithms and basic infrastructure for creating and suitably transforming the input and analyzing the results.

#### 2 Transaction data

The main application of association rules is for market basket analysis. In every transaction there are the items contained which are purchased at one visit. In one transaction there is usually only a small part of the items contained and the main interest is if the item is purchased or not. There are also other kinds of transaction data possible, as e.g., data containing the web pages visited during one session.

It can be assumed that transaction data consists of entries of the form

- transaction ID
- item ID
- user ID (optional)

Further information on transactions (as e.g., time, location), items (as e.g., category, price) or users (as e.g., socio-demographic variables: age, gender) might be available.

This kind of data can be transformed into a matrix with columns equal to the number of different transactions and rows equal to the number of different items. The matrix entries are the items present in the transaction. This matrix will in general be sparse.

Another application of mining association rules has been proposed for discovering interesting relationships between the values of different categorical variables. An example can be found in Hastie, Tibshirani, and Friedman (2001), where questionnaires data is used. Their data consists of ordinal, nominal and metric variables. A transaction for this kind of data consists of one customer and his characteristics and the items are the different values of all variables. This data can be transformed into a binary matrix, by first dichotomizing the metric variables and then coding each variable with k categories by k dummy variables.

The algorithms used for mining frequent itemsets or association rules need as input the transaction data transformed into a binary incidence matrix. A natural representation is a sparse matrix

format. For our implementation we choose the format cscMatrix which is defined in the R package Matrix. "csc" stands for compressed, sparse, column-oriented. It contains the indices of the rows unequal to zero, the pointers to the initial indices of elements in each column and the non-zero elements of the matrix. For a binary matrix the non-zero elements are all equal to 1.

Another possibility of storing binary matrices would be to interpret every column as a binary number with can be coerced into an integer in the decimal system. We could then represent each column by one number. The drawback is that the decimal number grows exponentially with  $2^n$  where n is the number of columns. In R this transformation from a binary vector to a decimal number is only possible for  $n \leq 30$ , because then the integer number limit is reached. As the number of items will in general be much larger we have decided not to use this sparse format to represent the data.

The natural format for the questionnaires data in R is a data.frame. In order to mine frequent itemsets/association rules with apriori or eclat this data needs to be transformed to an incidence matrix. In addition it has to be taken care that the information on the different variables and their levels is not lost. Transaction data is hence represented in arules as a S4 class named assMatrix containing the following S4 classes:

- cscMatrix: sparse matrix representation of the incidence matrix
- attributes: contains the factor names, their levels, and labels constructed by collapsing factor name and level name separated by a dot

The attributes are necessary in order to keep track of the factors and levels of a data frame which is transformed into the sparse matrix format. This allows the retransformation of a assMatrix into the original data frame.

Methods implemented are

- show
- coercion: from (binary) matrices and data frames (and the corresponding retransformation)
- extract ([)
- %in%
- as.character
- as.list

# 3 Mining frequent itemsets/association rules

Free reference implementations of the algorithms apriori and eclat in C are available by Christian Borgelt (Borgelt and Kruse, 2002; Borgelt, 2003). The code is called directly from R by the functions rapriori and reclat and no writing to external files is necessary for the data exchange. The input format of the data for the R functions is assMatrix (or a data format which can be

coerced to assMatrix). Furthermore, information on the parameters needs to be given. It is distinguished between parameters which change the characteristics of the itemsets/rules/hyperedges mined, as e.g. the minimum support or the target, and parameters which influence the performance of the algorithm, as e.g., an initial sorting of the items with respect to their frequency.

The parameters which influence the output are contained in the argument parameter, the other parameters are summarized in control. These arguments have to be either instances of the classes APparameter and APcontrol for rapriori and ECparameter and ECcontrol for reclat or data which can be coerced to these classes, as e.g., NULL which will give the default values or a named list (names equal to slot names to change the default values). In these classes each slot specifies a different parameter and the values are already validated (to a certain extend). Parameters which

have an integer specification are coerced before validity checking. The default values are equal to the defaults of the stand-alone C programs (cp. Borgelt, 2004) except that by default the original support definition is used for the specified minimum support required.

For rapriori there can be also the argument appearance specified, which determines which itemsets/rules/hyperedges are mined with respect to the items contained in the body or head. It will in general be specified by a named list containing

- default: character, one of "head", "body", "none", "both" (or one of the other names described in the apriori manual; Borgelt, 2004)
- head: character or integer, specifying the items which are allowed to appear in the head of the rules
- body: character or integer, specifying the items which are allowed to appear in the body of the itemsets/rules/hyperedges
- none: character or integer, specifying the items which are not allowed to be contained in the itemsets/rules/hyperedges
- both: character or integer, specifying the items which are allowed to be contained in the itemsets/rules/hyperedges

This list is coerced to an ASappearance object within rapriori where the attributes information in the assMatrix data is used for determining the column number if character values are given. The output of the functions rapriori and reclat is an object of class arules which contains the call, the (suitably modified) parameter specifications and the sets mined.

## 4 Sets: itemsets, rules, hyperedges

The output of rapriori or reclat are either itemsets (frequent itemsets, closed itemsets, maximal itemsets), rules or hyperedges. All three kinds contain the items involved and quality measures, as e.g., support, confidence, . . . .

In **arules** there is a virtual class sets defined which contains itemsets, rules and hyperedges as subclasses. It contains the following slots/classes:

- body: cscMatrix containing in each column the items which are in the body of the corresponding itemset/rule/hyperedge
- quality: data.frame with a row for each itemset/rule/hyperedge and a column for each quality measure (support, body.support, confidence, ...) returned
- rnb: integer number of rules
- attributes: containing the factor names, levels and labels from the input data

In addition rules contain

ullet head: cscMatrix containing in each column the items contained in the head of each rule

and itemsets contain

• trans: cscMatrix containing in each column the transactions which support the corresponding itemset (only returned by eclat if trans = TRUE)

Methods implemented are

• show

- summary
- extract ([)
- subset
- as.data.frame
- as.character

### 5 Example

**arules** provides the census data set from the UCI machine learning repository (Blake and Merz, 1998).

Transaction matrix in sparse format with dimension 132 48842

The census data set contains 48842 observations on 14 categorical variables. If the data frame is transformed into a binary incidence matrix using dummy coding the number of different items is 132.

We can use this data for calling rapriori and finding all rules with a minimum support of 0.05 and where all other parameters have the default values.

```
rules containing 19038 sets
derived using a minimum original support of 0.05
rapriori(data = census.assMatrix, parameter = list(support = 0.05))
Parameter specification:
 confidence minval smax arem aval originalSupport support minlen maxlen target
                       1 none FALSE
                0
                                                 TRUE
                                                          0.05
                                                                    1
                                                                            5 rules
  ext
 TRUE
Algorithmic control specification:
 filter tree heap memopt load sort verbose
    0.1 TRUE TRUE FALSE TRUE
The specified parameter values are validated and a support > 1 gives
> error <- try(rapriori(census.assMatrix, parameter = list(support = 1.3)))</pre>
> error
[1] "Error in validObject(.Object) : Invalid \"APparameter\" object: support = 1.3 > 1\n"
attr(,"class")
[1] "try-error"
The function subset can be used if we are only interested in the subset of rules which certain
variables or values of variables in the head/body or where a given quality measure fulfills a certain
criterion.
> sets <- rules@sets
> sets.sub <- subset(sets, subset = head %in% "sex" & lift > 1.4)
For an overview on the rules mined the function summary can be used which gives a description on
the number of rules, their lengths, the items contained in body and head and the quality measures.
> summary(sets.sub)
Object contains 612 rules.
body lengths:
      2
         3
  2 33 168 409
head:
Female.sex
             Male.sex
         1
                   611
body:
             Husband.relationship Married-civ-spouse.marital-status
                                470
                 none.capital-loss
                                         United-States.native-country
                                159
                                                                     158
```

none.capital-gain

155

White.race

157

#### (Other) 937

#### quality:

support	confidence	lift	body.support
Min. :0.05004	Min. :0.8005	Min. :1.401	Min. :0.05004
1st Qu.:0.06447	1st Qu.:0.9997	1st Qu.:1.495	1st Qu.:0.06585
Median :0.08276	Median :0.9999	Median :1.496	Median :0.08357
Mean :0.10863	Mean :0.9886	Mean :1.481	Mean :0.10956
3rd Qu.:0.12598	3rd Qu.:1.0000	3rd Qu.:1.496	3rd Qu.:0.12599
Max. :0.40365	Max. :1.0000	Max. :2.415	Max. :0.40367

The sets object can be coerced to a data frame which allows the printing of the rules similar to the output format of the C programs.

> as.data.frame(sets.sub)[1:3, ]

	head			body	support
54	Male.sex		Cr	aft-repair.occupation	0.11852504
145	Male.sex			Husband.relationship	0.40364850
436	Male.sex	Exec-manage	erial.occupation	Husband.relationship	0.06615208
	confidenc	e lift	body.support		
54	0.947153	31 1.416871	0.12513820		
145	0.999949	3 1.495851	0.40366897		
436	1.000000	00 1.495926	0.06615208		

### References

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