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***Abstract*—Discovering pattern sets or global patterns is an attractive issue from the pattern mining community in order to provide useful information. By combining local patterns satisfying a joint meaning, this approach produces patterns of higher level and thus more useful for the end-user than the usual local patterns. In parallel, recent works investigating relationships between data mining and constraint programming (CP) show that the CP paradigm is a powerful framework to model and mine patterns in a declarative and generic way. We present a constraint-based language which enables us to define queries in a declarative way addressing patterns sets and global patterns. By specifying what the task is, rather than providing how the solution should be computed, it is easy to process by stepwise refinements to successfully discover global patterns. The usefulness of the approach is highlighted by several examples coming from the clustering based on associations. All primitive constraints of the language are modeled and solved using the SAT framework. We illustrate the efficiency of our approach through several experiments.**

I. Introduction

The process of extracting useful patterns from data, called *pattern mining*, is an important tool for data analysis and has been used in a wide range of applications and domains. A large amount of work has been developed and many pattern extraction problems are now identified and understood from both theoretical and computational perspectives. Local pattern discovery has become a growing field [19] and several paradigms are available for producing extensive collections of patterns such as the constraint-based pattern mining [20] or condensed representations of patterns [3]. Because of the exhaustive nature of the techniques, the pattern collections provide a fairly complete picture of the information content of the data. However, the approach suffers from limitations. First, the collections of patterns still remain too large for an individual and global analysis performed by the data analyst. Secondly, the so-called local patterns represent fragmented information whereas patterns expected by the data analyst require to consider simultaneously several local patterns. That is why combining local patterns to get global patterns is highly attractive.

The data mining literature includes several methods to take into account the relationships between patterns and produce global patterns or pattern sets [4], [9]. Recent approaches - constraint-based pattern set mining [4], pattern teams [15] and selecting patterns according to the added value of a new pattern given the currently selected patterns [2] - aim at reducing the redundancy by selecting patterns from the initial large set of local patterns on the basis of their usefulness in the context of the other selected patterns. Nevertheless, these methods are mainly based on the reduction of the redundancy or specific aims such as classification processes. The difficulty of the task may explain the use of heuristic functions and the lack of complete and correct methods to mine global patterns. Indeed, mining local patterns requires the exploration of a large search space but mining global patterns is even harder because solutions satisfying each pattern must be compared. Clearly, the lack of generic approaches restrains the discovery of useful global patterns because the user has to develop a new method each time he wants to extract a new kind of global patterns. It explains why this issue deserves our attention.

In this paper, we propose a constraint-based language to discover patterns combining several local patterns. The key idea is to propose a declarative and generic approach to ask queries: the user models a problem by specifying a set of constraints and expresses his queries thanks to constraints over terms built from constants, variables, operators, and function symbols. Queries and built-in constraints of the language are encoded and solved using the SAT framework. Our approach takes benefit of the recent progress on cross-fertilization between data mining and Constraint Programming [11], [12], [13], [22]. The definition of a constraint-based language offers the great advantage to provide a declarative method to address different pattern mining problems: it is enough to change the specification in term of constraints. We illustrate the approach by several examples coming from the clustering based on associations. With simple query refinements, the data analyst is able to easily produce clusterings satisfying different properties. By specifying what the task is, rather than providing how the solution should be computed, the process greatly facilitates the search of global patterns and the discovery of knowledge.

This paper is organized as follows. Section II describes the constraint-based language and shows how queries and constraints can be defined using terms and built-in constraints. Starting from the clustering example, Section III depicts the process of successive refinements which enables us to easily address several kinds of clustering and the discovery of global models. Section IV describes how queries and built-in

constraints of the language are modeled and solved using the SAT framework. Section V demonstrates the efficiency of our approach through several experiments. We review related work in Section VI.

II. A Constraint-based Language

This section describes the constraint-based language we propose. Terms are built using constants, variables, operators, and function symbols. Constraints are relations over terms that can be satisfied or not. The data analyst can define new function symbols. We show how queries and constraints can be written using built-in constraints.

A. Definitions and example

Let $\mathcal{I}$ be a set of n distinct literals called items, an itemset (or *pattern*) is a non-null subset of $\mathcal{I}$. The language of itemsets corresponds to $\mathcal{L}_\mathcal{I} = 2^\mathcal{I} \backslash \emptyset$. A *transactional dataset* is a multi-set of m itemsets of $\mathcal{L}_\mathcal{I}$. Each itemset, usually called a *transaction* or object, is a database entry. For instance, Table I gives a transactional dataset $\mathcal{T}$ with m=11 transactions $t_1, \ldots, t_{11}$ described by n=8 items A, B, C, D, E, F, G, H. Interestingness measures such as the frequency and the area [8] are commonly used to evaluate the relevance of patterns.

Definition 1. (frequency) The frequency of a pattern X_i is the number of transactions that X_i covers in $\mathcal{T}$: $\texttt{freq}(X_i) = |\{t \in \mathcal{T} \mid X_i \subseteq t\}|$.

Definition 2. (area) Let X_i be a pattern, $\texttt{area}(X_i) = \texttt{freq}(X_i) \times \texttt{size}(X_i)$ where $\texttt{size}(X_i)$ denotes the cardinality of X_i.

From our running example (cf. Table I), $\texttt{freq}(\{A, E\}) = 3$ and $\texttt{freq}(\{C, F, G, H\}) = 1$. The *frequency constraint* $\texttt{freq}(X_i) \geq$ *minfr* focuses on patterns occurring in the dataset a number of times exceeding a given minimal threshold *minfr*. On the other hand, 9 patterns satisfy the constraint $\texttt{area}(X_i) \geq 6$: $\{A, E, G\}$, $\{B, E, G\}$, $\{C, E, G\}$, $\{C, E, H\}$, $\{E, G\}$, $\{C, E\}$, $\{C, H\}$, $\{E\}$, $\{G\}$. The goal of constraint-based pattern mining is to discover all the patterns of $\mathcal{L}_\mathcal{I}$ satisfying a given constraint. The rest of this section depicts our proposition to express constraints. Contrary to the previous examples, several patterns can be involved in the following constraints.

B. Terms

Terms are built from constants, variables, operators, and function symbols.

- **constants** are either numerical values (as threshold *minfr*), or items (as A) or patterns (as $\{A, B\}$) or transactions (as t_7).
- **variables**, noted X_i, for $1 \leq i \leq k$, represent the unknown patterns.
- **operators**:
  - set operators as $\cap, \cup, \setminus, \ldots$
  - numerical operators as $+, -, \times, /, \ldots$

Trans.	Items						
t_1	A			D		F	
t_2	A				E	F	
t_3	A				E		G
t_4	A				E		G
t_5		B			E		G
t_6		B			E		G
t_7			C		E		G
t_8			C		E		G
t_9			C		E			H
t_{10}			C		E			H
t_{11}			C			F	G	H

Table I
Transactional dataset $\mathcal{T}$.

- **function symbols** involving one or several patterns: `freq/1`, `size/1`, `cover/1`, `overlapItems/2`, `overlapTransactions/2`, ...

Examples of terms:

- $\texttt{freq}(X_1) \times \texttt{size}(X_1)$ (i.e. the `area` of X_1)
- $\texttt{freq}(X_1 \cup X_2) \times \texttt{size}(X_1 \cap X_2)$ (i.e., the overlapping between $\texttt{area}(X_1)$ and $\texttt{area}(X_2)$)
- $\texttt{freq}(X_1) - \texttt{freq}(X_2)$

1) Built-in function symbols: The constraint based language owns predefined (built-in) function symbols[1] like:

- $\texttt{cover}(X_i) = \{t \mid t \in \mathcal{T}, X_i \subseteq t\}$ is the set of transactions covered by X_i.
- $\texttt{freq}(X_i) = \mid \{t \mid t \in \mathcal{T}, X_i \subseteq t\} \mid$
- $\texttt{size}(X_i) = \mid \{j \mid j \in \mathcal{I}, j \in X_i\} \mid$
- $\texttt{overlapItems}(X_i, X_j) = \mid X_i \cap X_j \mid$ is the number of items shared by both X_i and X_j.
- $\texttt{overlapTransactions}(X_i, X_j) = \mid \texttt{cover}(X_i) \cap \texttt{cover}(X_j) \mid$ is the number of transactions covered by both X_i and X_j.

2) User-defined function symbols: The data analyst can define new function symbols using constants, variables, operators and existing function symbols (built-in or previously defined ones). Examples:

- $\texttt{area}(X_i) = \texttt{freq}(X_i) \times \texttt{size}(X_i)$
- $\texttt{coverage}(X_i, X_j) = \texttt{freq}(X_i \cup X_j) \times \texttt{size}(X_i \cap X_j)$
- interestingness measures can be straightforwardly defined. Here is the example of the `growth-rate` well-used in contrast mining [21]. Let $D_1, D_2 \subset \mathcal{T}$ be 2 sets of transactions (i.e., classes) and $\texttt{freq'}(X_i, D_j)$ the frequency of X_i into D_j, the growth-rate of X_i in D_1 is $gr_1(X_i) = \frac{|D_2| \times \texttt{freq'}(X_i, D_1)}{|D_1| \times \texttt{freq'}(X_i, D_2)}$

C. Constraints and Queries

Constraints are relations over terms that can be satisfied or not. There are three kinds of built-in constraints:

[1]Only function symbols used in Section III are introduced in this paper.

- numerical constraints like: $<, \leq, =, \neq, \geq, >, \ldots$
Examples:

- $\texttt{freq}(X_1) \leq 10$
- $\texttt{size}(X_2) = 2 \times \texttt{size}(X_3)$
- $\texttt{area}(X_1) < \texttt{size}(X_2) \times \texttt{freq}(X_3)$

- set constraints like: $=, \neq, \in, \notin, \subset, \subseteq, \ldots$
Examples:

- $A \in X_1$
- $X_1 \cup X_2 \subset X_3$
- $X_1 = X_2 \cap X_4$

- dedicated constraints like:

- $\texttt{closed}(X_i)$ is satisfied iff X_i is a closed[2] pattern.
- $\texttt{coverTransactions}([X_1, ..., X_k])$ is satisfied iff each transaction is covered by at least one pattern (i.e. $\bigcup_{1 \leq i \leq k} \texttt{cover}(X_i) = \mathcal{T}$),
- $\texttt{coverItems}([X_1, ..., X_k])$ is satisfied iff every item belongs to at least one pattern (i.e. $\bigcup_{1 \leq i \leq k} X_i = \mathcal{I}$).
- $\texttt{canonical}([X_1, ..., X_k])$ is satisfied iff for all i s.t. $1 \leq i < k$, pattern X_i is less than pattern X_{i+1} with respect to the lexicographic order.

Queries are formulae built using constraints and logical connectors: $\wedge$ (conjunction) and $\vee$ (disjunction).

III. Mining by refining contraint-based queries

A major strength of our approach is to provide a simple and efficient way to declare and refine queries. In practice, the data analyst starts by writing a first query Q_1. Then, he successively refines the query (deriving Q_{i+1} from Q_i) until he considers that relevant information has been extracted. We illustrate this approach with the clustering problem. Clustering aims at partitioning data into groups (clusters) so that transactions are similar inside each cluster but different between clusters [7]. We selected clustering because it is an important and popular data mining task and, by nature, clustering proceeds by iteratively refining queries until a satisfactory solution is found. Our approach is also well-suited to integrate constraints handled in constraint-based clustering [1].

A. Modeling a clustering query

The closed patterns are well-designed for clustering based on associations because a closed pattern gathers the maximum amount of similarity between a set of transactions. Thus, a closed pattern is a candidate cluster. The standard clustering problem can then be formalized as: "to find a set of k closed patterns $X_1, X_2, ..., X_k$ (i.e., clusters) covering all transactions without any overlap on these transactions".

Our constraint-based language offers the constraints to express this query: the $\texttt{closed}(X_i)$ constraints enforce each unknown pattern X_i to be closed, the $\texttt{coverTransactions}([X_1, ..., X_k])$ constraint ensures to cover all the transactions. To avoid an overlap between transactions, we add for each couple of patterns (X_i, X_j) s.t. $i<j$ the $\texttt{overlapTransactions}(X_i, X_j)=0$ constraint. This constraint states that there exists no transaction covered by both X_i and X_j.

Moreover, a clustering problem intrinsically owns a lot of symmetrical solutions. Let $s = (p_1, p_2, ..., p_k)$ be a solution containing k patterns p_i, any permutation σ of these k patterns $\sigma(s) = (p_{\sigma(1)}, p_{\sigma(2)}, ..., p_{\sigma(k)})$ is also a solution. The $\texttt{canonical}([X_1, ..., X_k])$ constraint is used to avoid computing symmetrical solutions. This constraint ensures that, for all i s.t. $1 \leq i < k$, pattern X_i is before pattern X_{i+1} with respect to the lexicographic order. The $\texttt{canonical}([X_1, ..., X_k])$ constraint plays an important role. As for a clustering involving k clusters, the number of symmetrical solutions is $k!$, it is crucial to break the symmetries to avoid obtaining a huge number of redundant solutions. Moreover, this constraint performs an efficient filtering by drastically reducing the size of the search space.

Finally, we get the following query (Q_1) modeling the initial clustering problem:

$$\begin{cases} \wedge_{1 \leq i \leq k} \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1, ..., X_k]) \wedge \\ \wedge_{1 \leq i < j \leq k} \texttt{overlapTransactions}(X_i, X_j) = 0 \wedge \\ \texttt{canonical}([X_1, ..., X_k]) \end{cases}$$

On our running example, with k=3 patterns, query Q_1 provides 5 solutions (see Table II).

Sol.	X_1	X_2	X_3
s_1	{C, F, G, H}	{E}	{A, D, F}
s_2	{A, F}	{C, H}	{E, G}
s_3	{C, E, H}	{E, G}	{F}
s_4	{A, F}	{C, E, H}	{G}
s_5	{A}	{B, E, G}	{C}

Table II
Set of different clusterings.

Sol.	X_1	X_2	X_3
s'_1	{C, F, G, H}	{E}	{F}
s'_2	{A, D, F}	{C, F, G, H}	{E}
s'_3	{A, F}	{C, F, G, H}	{E}
s'_4	{A, E, F}	{E}	{F}
s'_5	{A, D, F}	{E}	{F}
s'_6	{A}	{B, E, G}	{C}
s'_7	{A, F}	{C, E, H}	{G}
s'_8	{A, F}	{C, H}	{G}
s'_9	{A, F}	{C, H}	{E, G}
s'_{10}	{C, E, H}	{E, G}	{F}
s'_{11}	{C, H}	{E, G}	{F}
s'_{12}	{C, H}	{F}	{G}
s'_{13}	{C, E, H}	{F}	{G}

Table III
Set of different clusterings for query Q_4.

[2]Let Tr_i be the set of transactions covered by pattern X_i. X_i is closed iff X_i is the largest ($\subset$) pattern covering Tr_i.

B. Refining queries

By only refining queries on a clustering, the data analyst can easily produce other clusterings satisfying different properties. This section illustrates this feature of our approach that facilitates the building of global patterns and the discovery of knowledge. From the initial query Q_1, we derive queries Q_2 and Q_3 avoiding clusterings with non-frequent patterns and clusterings with small size patterns.

1) Removing solutions with non-frequent patterns: When a cluster has a low frequency, it lacks of representativity and the clustering is not considered as reliable. From Q_1, it is easy to add frequency constraints ensuring that each pattern is frequent. With a frequency threshold δ_1=2, we get Q_2:

$$\begin{cases} \wedge_{1\leq i\leq k}\ \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapTransactions}(X_i,X_j)=0 \wedge \\ \texttt{canonical}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i\leq k}\ \texttt{freq}(X_i)\geq \delta_1 \end{cases}$$

Pattern $\{C,F,G,H\}$ of solution s_1 has a frequency of 1 and thus is removed. With Q_2, there remain 4 solutions (s_2, s_3, s_4, and s_5, see Table II).

2) Removing solutions with small size patterns: A clustering with at least one cluster X_i of small size[3] is not considered as useful because X_i does not ensure enough similarity between transactions associated to X_i. It is simple to add constraints requiring that the size of each pattern is higher than a minimal size. From Q_2, with a minimal size threshold δ_2=2, we obtain the query Q_3:

$$\begin{cases} \wedge_{1\leq i\leq k}\ \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapTransactions}(X_i,X_j)=0 \wedge \\ \texttt{canonical}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i\leq k}\ \texttt{freq}(X_i)\geq \delta_1 \wedge \\ \wedge_{1\leq i\leq k}\ \texttt{size}(X_i)\geq \delta_2 \end{cases}$$

With the query Q_3, there is only one solution: $s2$ with X_1=$\{A,F\}$, X_2=$\{C,H\}$ and X_3=$\{E,G\}$ (cf. Table II).

C. Solving other Clustering Problems

In the same way, it is easy to express other clustering problems [1] such as soft clustering, co-clustering, and soft co-clustering.

*1) **Soft clustering**:* This problem is a relaxed version of the clustering where small overlaps on transactions (less than a threshold δ_T) are allowed. The query Q_4 (soft version of Q_1) models this problem:

$$\begin{cases} \wedge_{1\leq i\leq k}\ \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapTransactions}(X_i,X_j)\leq \delta_T \wedge \\ \texttt{canonical}([X_1,...,X_k]) \end{cases}$$

[3]Moreover, clustering with clusters of size 1 often reflects values coming from the binarization of an attribute (such as A, B and C in Table I) and are useless.

With k=3 and a maximal overlap between transactions δ_T=1, Q_4 produces 13 solutions (see Table III).

With s'_1, the overlaps are the transaction t_{11} (covered by X_1 and X_3) and the transaction t_2 (covered by X_2 and X_3) (see Tables III and I). Removing solutions with non frequent patterns (with δ_1=2) leads to 8 solutions (from s'_6 to s'_{13}). By adding a minimal size constraint δ_2=2, only the solution s'_9 remains (which is also the solution s_2 of the initial clustering problem, see Section III-B2).

*2) **Co-clustering**:* The co-clustering task consists in finding k clusters covering both the set of transactions and the set of items, without any overlap on transactions or on items. Query Q_5 expresses this problem:

$$\begin{cases} \wedge_{1\leq i\leq k}\ \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapTransactions}(X_i,X_j)=0 \wedge \\ \texttt{coverItems}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapItems}(X_i,X_j)=0 \wedge \\ \texttt{canonical}([X_1,...,X_k]) \end{cases}$$

*3) **Soft co-clustering**:* This problem is a relaxed version of the co-clustering, allowing small overlaps on transactions (less than δ_T) and on items (less than δ_I). The query Q_6 (soft version of Q_4 and Q_5) models this task:

$$\begin{cases} \wedge_{1\leq i\leq k}\ \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapTransactions}(X_i,X_j)\leq \delta_T \wedge \\ \texttt{coverItems}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapItems}(X_i,X_j)\leq \delta_I \wedge \\ \texttt{canonical}([X_1,...,X_k]) \end{cases}$$

*4) **Balanced clustering**:* In clustering, we generally prefer solutions in which the frequencies of the clusters do not differ too much from each other. Query Q_7 describes clusterings with balanced frequencies. For any couple of clusters (X_i,X_j), their difference of frequencies must be lower than a threshold $\Delta\times m$ where Δ is a percentage. Looking for clusterings with balanced size clusters could be achieved in a same way.

$$\begin{cases} \wedge_{1\leq i\leq k}\ \texttt{closed}(X_i) \wedge \\ \texttt{coverTransactions}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \texttt{overlapTransactions}(X_i,X_j)=0 \wedge \\ \texttt{canonical}([X_1,...,X_k]) \wedge \\ \wedge_{1\leq i<j\leq k}\ \ |\ \texttt{freq}(X_i)-\texttt{freq}(X_j)\ |\leq \Delta\times m \end{cases}$$

IV. Modeling and Solving as a SAT Problem

Satisfiability (SAT) is the problem of determining if the variables of a given boolean formula can be assigned in such a way as to make the formula be evaluated to `True`. A formula is in conjunctive normal form (CNF) if it is a conjunction of clauses, where a clause is a disjunction of literals and a literal is either a variable x_i or its negation $\neg x_i$.

Even if the SAT problem is NP-complete, efficient and scalable algorithms for SAT, that were developed over the

last decade, have contributed to dramatic advances in the ability to automatically solve problem instances involving tens of thousands of variables and millions of constraints. That is why, we have chosen to transform a query into a CNF and then use a SAT solver to find its solutions.

In the remainder of this section, let $(d_{t,i})$ be the (m,n) boolean matrix where ($d_{t,i}$=`True`) iff ($i \in t$). Queries are modeled in two steps. First, unknown patterns are modeled using boolean variables and matrix $(d_{t,i})$. Then, each built-in constraint its expressed using a CNF.

A. Modeling unknown patterns

Let $X_1, X_2, ..., X_k$ be the k patterns we are looking for. In a same way as [13], [22], the link between the data set $\mathcal{T}$ and an unknown pattern X_j is performed by introducing two kinds of boolean variables:

- $X_{1,j}, X_{2,j}, ..., X_{n,j}$ s.t. ($X_{i,j}$=`True`) iff $(i \in X_j)$
- $T_{1,j}, T_{2,j}, ..., T_{m,j}$ s.t. ($T_{t,j}$=`True`) iff $(X_j \subset t)$

The relation $(X_j \subset t)$ can be transformed into the following CNF:

$$\bigwedge_{\{i \in \mathcal{I} | \neg d_{t,i}\}} \neg X_{i,j} \tag{1}$$

The relationship between X_j and $\mathcal{T}$ is modeled by stating that, for each transaction t, ($T_{t,j}$=`True`) iff X_j covers t:

$$\forall t \in \mathcal{T}, T_{t,j} \Leftrightarrow (X_j \subset t) \tag{2}$$

Using Eq. 1, the left to right implication of Eq. 2 can be transformed into the following CNF:

$$\bigwedge_{t \in \mathcal{T}} (\bigwedge_{\{i \in \mathcal{I} | \neg d_{t,i}\}} (\neg T_{t,j} \vee \neg X_{i,j})) \tag{3}$$

Using Eq. 1, the right to left implication of Eq. 2 can be transformed into the following CNF:

$$\bigwedge_{t \in \mathcal{T}} (\bigvee_{\{i \in \mathcal{I} | d_{t,i}\}} X_{i,j} \vee T_{t,j}) \tag{4}$$

Finally, Eq. 3 and Eq. 4 must hold for every X_j, $1 \leq j \leq k$. So, the SAT encoding of a query with k unknown patterns requires $k \times m \times n$ binary clauses.

B. Constraints as boolean formulae

This section provides the boolean formulae associated to built-in constraints. The following ones have a straightforward encoding:

- $X_p = X_q \rightarrow \wedge_{i \in \mathcal{I}} (X_{i,p} \Leftrightarrow X_{i,q})$
- $i_o \in X_p \rightarrow X_{i_o,p}$
- $X_p \cap X_q = X_r \rightarrow \wedge_{i \in \mathcal{I}} (X_{i,r} \Leftrightarrow X_{i,p} \wedge X_{i,q})$
- $X_p \cup X_q = X_r \rightarrow \wedge_{i \in \mathcal{I}} (X_{i,r} \Leftrightarrow X_{i,p} \vee X_{i,q})$
- $X_p \backslash X_q = X_r \rightarrow \wedge_{i \in \mathcal{I}} (X_{i,r} \Leftrightarrow X_{i,p} \wedge \neg X_{i,q})$
- `coverItems`$([X_1, ..., X_k]) \rightarrow \wedge_{i \in \mathcal{I}} (\vee_{j \in [1..k]} X_{i,j})$
- `coverTransactions`$([X_1, ..., X_k]) \rightarrow \wedge_t (\vee_j T_{t,j})$

All threshold constraints are modelled using the *sorting network* approach (for technical details, see [6]) in order to prevent prohibitive grounding. Using such an encoding, the size of the CNF modelling a threshold constraint is independent from the value of the threshold but depends on the maximal value for the considered measure. For example, the size of the CNF for constraint (`freq`$(X_i) \geq \delta_1$) is $O(m \times log(m))$, and does not depend on δ_1. The size of the CNF for constraint (`size`$(X_j) \leq \delta_2$) is $O(n \times log(n))$, and does not depend on δ_2. Threshold constraints for measures like `overlapItems` and `overlapTransactions` are encoded in the same way.

dataset	#transactions	#items	density
`Australian`	690	125	0.40
`Mushroom`	8124	119	0.19
`Soybean`	630	50	0.32
`Primary-Tumor`	336	36	0.48
`Zoo`	101	36	0.44
`Meningitis`	329	82	0.26

Table IV
DESCRIPTION OF THE DATASETS.

C. Ensuring completeness while using a SAT solver

Given a CNF, SAT solvers either find one instanciation (and only one) for the variables evaluating the formula to `True`, or prove there is no such an instanciation. In order to ensure the completeness of our approach, restarts are performed.

Let $\mathcal{F}$ be the CNF modeling a query Q. $\mathcal{F}$ is the conjunction of the CNF associated to the modeling of unknown patterns (see Section IV-A) and the CNFs associated to the modeling of the constraints involved in Q (see Section IV-B). Resolution begins with $\mathcal{F}$. Then, after having obtained the i-th solution s_i, its negation $\neg s_i$ is added to the (current) CNF and resolution is restarted in order to look for another solution. The process ends when a failure occurs, i.e. when all solutions have been found.

Using restarts may seem too naive, but in practice is powerful (see experiments in Section V-A). This is because the CNF $\mathcal{F}$ contains much binary clauses (see Section IV-A), and so, filtering by unit propagation will be very effective.

The SAT solver `MiniSat`[4] [5] has been used for experiments, because its implementation proved easy to modify and `MiniSat` is one of the most efficient SAT solvers.

V. EXPERIMENTS

Recalling that the key contribution of this paper is to propose a constraint-based language to model and mine global patterns in a declarative way. Therefore, the goal of the experiments is to provide better insights on the use of our approach in order to discover global patterns with the example of clustering based on associations.

[4] `http://minisat.se/`

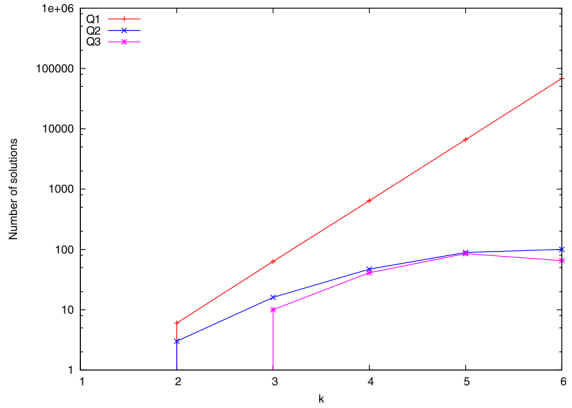

Figure 1. Number of solutions for refinements Q_1 to Q_3 (Soybean).

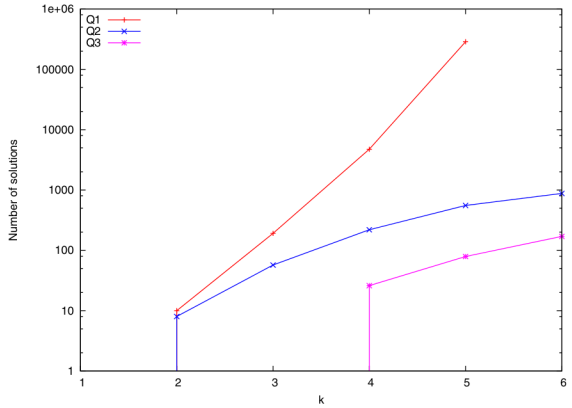

Figure 2. Number of solutions for refinements Q_1 to Q_3 (Meningitis).

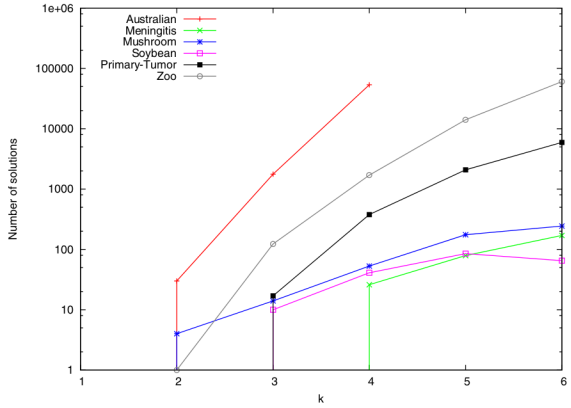

Figure 3. Number of solutions to query Q_3 on several datasets.

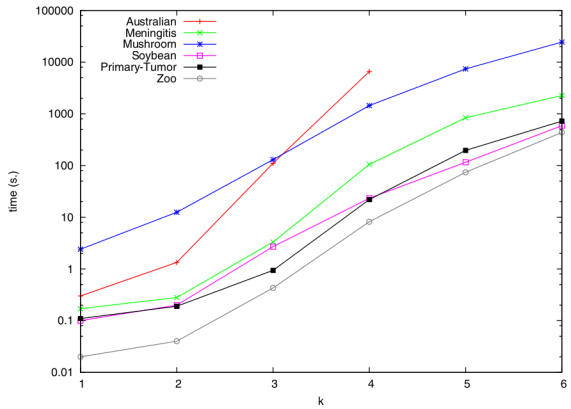

Figure 4. Computing times for query Q_3 on several datasets.

Experiments were performed on several benchmarks from the UCI repository[5] and also a real-world dataset Meningitis gathering 329 children hospitalized for acute meningitis. Characteristics of datasets are presented in Table IV. Experiments were conducted on a PC having a 2.83 GHz Intel Core 2 Duo Processor and 4GB of RAM, running Ubuntu Linux.

A. Clustering Queries

This section illustrates the successive query refinement depicted in Section III and leading to the query Q_3.

Fig. 1 (resp. Fig. 2) gives the total number of solutions for queries Q_1, Q_2, and Q_3 for the dataset Soybean (resp. Meningitis) according to k (i.e., the number of patterns). Applying the stepwise refinements from query Q_1 to query Q_3 drastically reduces the number of clusterings and highlights on the most promising ones (note that the y-axis is a logarithmic scale).

Fig. 3 depicts the number of solutions to query Q_3 on several datasets. In some cases, the query can be satisfied by a large number of solutions and we do not report results when there are more than one million of solutions. It especially happens with Australian because this dataset provides a huge number of closed patterns. In such situations, the query should be refined (by increasing the minimal frequency and/or size thresholds or adding new constraints).

The completeness of our approach is ensured by restarting the SAT solver. Each time a new solution is found, its negation is stored and added to the current CNF. Memory consumption becomes too high above millions of solutions. From a practical point of view, as the data analyst is interested in extracting a small number of solutions, it does not make sense to perform mining providing millions of patterns. Once again, the stepwise refinement query should be used to avoid these situations and to focus on the most relevant patterns.

Fig. 4 gives the computing times on several datasets, showing the efficiency of our approach. Even for rather large datasets like Mushroom, computing times are still affordable.

B. Soft Clustering

Fig. 5 and Fig. 6 provide soft clustering results with query Q_4 on datasets Soybean and Primary-Tumor.

[5] http://www.ics.uci.edu/~mlearn/MLRepository.html

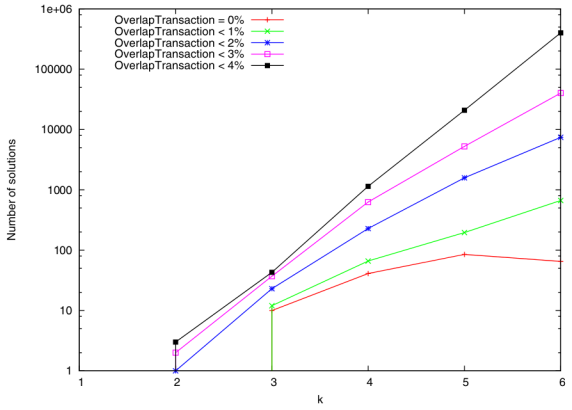

Figure 5. No. of sol. according to k for several δ_T values (Soybean).

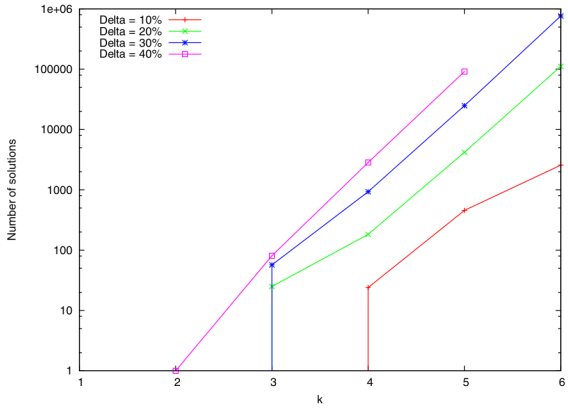

Figure 7. No. of sol. according to k for several Δ values (Zoo).

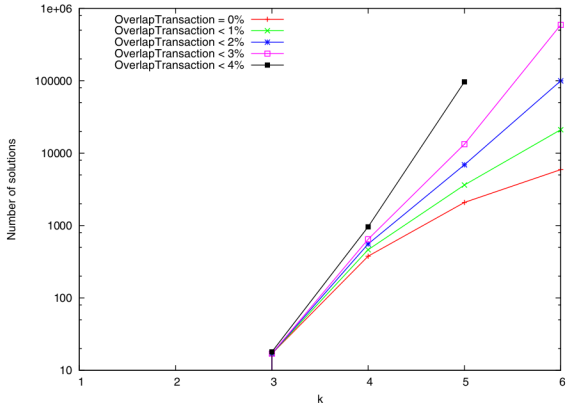

Figure 6. No. of sol. according to k for several δ_T values (Primary-Tumor).

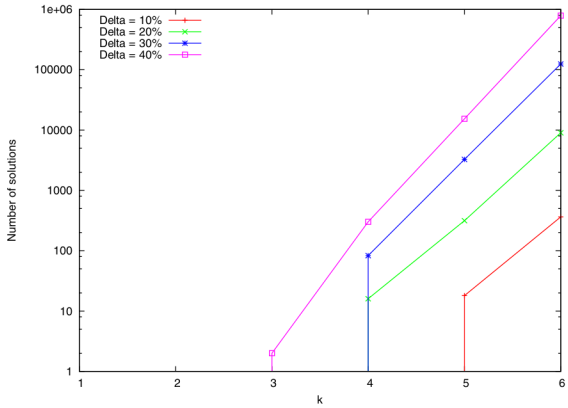

Figure 8. No. of sol. according to k for several Δ values (Primary-Tumor).

Curves indicate the number of solutions according to k and the maximal overlap threshold δ_T on transactions. As expected, the number of solutions increases with the size of the overlap. The number of solutions can be easily controlled by adjusting (decreasing/increasing) δ_T.

C. Balanced Clustering

We performed experiments with query Q_7. Fig. 7 and Fig. 8 display curves indicating the number of solutions on datasets Zoo and Primary-Tumor according to k and Δ. Adding the balanced frequencies constraint strongly reduces the number of solutions. For dataset Zoo, only a gap Δ=40% allows to find a solution for k=2. Moreover, with k=3 and k=4, there is no solution with a gap Δ=10%.

VI. Related work

The data mining literature includes many approaches to reduce the number of produced patterns such as the condensed representations of patterns [3], the compression of the dataset by exploiting Minimum Description Length Principle [24], the discovery of k representative patterns with probabilistic models for summarizing frequent patterns [18]. These approaches mainly aim at reducing the redundancy between patterns and often focus on frequent patterns.

Taking into account the relationships between patterns is explicitly required to produce global patterns or pattern sets [2], [4], [9]. A large number of these methods are based on two-step techniques. The first step generates an exhaustive collection of local patterns and then the patterns are heuristically post-processed to select a smaller subset of complementary relevant patterns such as in associative classification [17]. There are few methods without heuristic to mine complete and correct pattern sets or global patterns and in practice running techniques are devoted to specific kinds of global patterns [16], [23]. General data mining frameworks based on the notion of local patterns to design global models are presented in [9], [14]. These frameworks help to analyze and improve current methods in the area.

Recent works [11], [12], [13], [22] have shown the cross-fertilization between data mining and Constraint Programming (CP). Indeed, CP provides a general declarative methodology for modeling and solving constraint problems. CP facilitates the design of generic methods handling several patterns, that is a key point in pattern set mining. Techniques

for mining itemsets and n-ary patterns (i.e., the combination of n patterns) have been proposed. Looking for declarative techniques in pattern mining was also recently investigated in [10]: by using relational algebra, the authors propose an algebraic framework for pattern discovery for expressing a wide range of queries.

VII. Conclusion and Future Work

We have proposed a constraint-based language allowing to easily express different mining tasks in a declarative way. Thanks to the declarative process, extending or changing the specification to refine the results and discover more relevant patterns or address new global patterns is very simple. Moreover, all constraints can be combined together and new constraints can be added. The efficiency and the flexibility of our approach is shown on several examples coming from clustering based on associations. Thanks to query refinements, the data analyst is able to produce clusterings satisfying different constraints and generating more meaningful clusters.

As future work, we want to enrich our constraint-based language with further constraints to capture and model a wider range of data mining tasks. The scalability of the approach to larger values of k and larger datasets will also be investigated. Another promising direction is to integrate optimisation criteria in our framework.

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