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An Inductive Database Prototype Based on Virtual Mining Views

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ABSTRACT
We present a prototype of an inductive database. Our system enables the user to query not only the data stored in the database but also generalizations (e.g. rules or trees) over these data through the use of virtual mining views. The mining views are relational tables that virtually contain the complete output of data mining algorithms executed over a given dataset. The prototype implemented into PostgreSQL currently integrates frequent itemset, association rule and decision tree mining. We illustrate the interactive and iterative capabilities of our system with a description of a complete data mining scenario.

Categories and Subject Descriptors: H.2.4 [Database Management]: System.


Keywords: Data Mining, Inductive Databases.

1. INTRODUCTION
Data mining is not a one-shot activity, but rather an iterative and interactive process. During the whole discovery process, typically, many different data mining tasks are performed, their results are combined, and possibly used as input for other data mining tasks. To support this knowledge discovery process, there is a need for integrating data mining with data storage and management. The concept of inductive databases (IDBs) has been proposed as a means of achieving such integration [7]. In [3, 4, 5, 6], we describe how such an inductive database can be designed in practice, by using virtual mining views.

In an IDB, one can not only query the data stored in the database, but also the patterns that are implicitly present in these data. One of the main advantages of our system is the flexibility of ad-hoc querying, that is, the user can specify new types of constraints and query the patterns and models in combination with the data itself and so forth. In this paper, we illustrate this feature with a data mining scenario in which we first learn a classifier over a given dataset and, afterwards, we look for correct association rules, which describes the misclassified examples w.r.t this classifier. Notice that the functionality of an inductive database goes far beyond that of data mining suites such as, Weka [10] and Yale [8]. These systems have the advantage of imposing one uniform data format for a group of algorithms. On the other hand, they do not allow ad-hoc queries.

The rest of the paper is organized as follows. Section 2 addresses the idea behind the development of our prototype. Section 3 presents the virtual mining views framework. In Section 4, we describe how the system is implemented. Finally, Section 5 describes a complete database scenario, which illustrates the interactive and iterative capabilities of our system.

2. DESCRIPTION OF THE SYSTEM
The system presented in this paper builds upon our preliminary work in [3, 5, 6]. In contrast to the numerous proposals for data mining query languages, we propose to integrate data mining into database systems without extending the query language. Instead, we extend the database schema with new tables containing, for instance, association rules, decision trees, or other descriptive or predictive models. As far as the user is concerned, these tables contain all possible patterns, trees, and models that can be learned over the data. Of course, such tables would in most cases be huge. Therefore, they are in fact implemented as views, called virtual mining views.

Whenever a query is formulated, selecting for instance association rules from these tables, a run of a data mining algorithm is triggered (e.g., Apriori [1]) to compute the result of the query, in exactly the same way that normal views in databases are only computed at query time, and only to the extent necessary for answering the query.

When the user formulates his or her mining query, the parser is invoked by the DBMS, creating an equivalent relational algebra expression. At this point, the expression is processed by the Mining Extension, which extracts from the query the constraints that can be pushed into the data mining algorithms. The output of these algorithms is then materialized in the virtual mining views. After the material-
The system is developed into PostgreSQL\textsuperscript{1} (written in C). We adapted the web-based administration tool PhpPgAdmin\textsuperscript{2}, in order to have a friendly interface to the system. When the user writes a query, PostgreSQL generates a data structure representing its corresponding relational algebra expression. After this data structure is generated, our Mining Extension is called. Here, we process the relational algebra structure, extract the constraints, trigger the data mining algorithms and materialize the results in the virtual mining views. Just after the materialization, the work-flow of the DBMS continues and the query is executed as if the patterns or models were there all the time.

The system is currently linked to algorithms for frequent itemset mining, association rule discovery and exhaustive decision tree learning [6]. The constraints represented as attributes of the mining views (size, accuracy, support, confidence) can be extracted and efficiently exploited by the integrated data mining algorithms.

Experiments published in [4] showed that the execution times of the queries are rather low which support our claim that the virtual mining views provide an elegant way to incorporate data mining capacities to database systems without changing the query language.

\textsuperscript{1}http://www.postgresql.org/
\textsuperscript{2}http://phppgadmin.sourceforge.net/
create table simple_mushroom
as select cap_color, odor, gill_size,
gill_color, spore_print_color, class
from mushroom;

alter table simple_mushroom add column id serial;

Figure 2: Pre-processing.

Figure 3: Mining views for table simple_mushroom.

5. AN ILLUSTRATIVE SCENARIO

In this section, we describe an extended scenario that explores the interactive and iterative capabilities of our system. The scenario consists of mining decision trees over the mushroom dataset [2] and also association rules over intermediate query results. We assume that table MUSHROOM is already stored in our system. It contains 8,124 tuples and 23 categorical attributes. The attribute class discriminates mushrooms from being poisonous or edible, while the other attributes describe features of the mushrooms, such as the color of the cap and odor.

Step 1: Pre-processing.

To illustrate how it is possible to preprocess data using our system, we create a new table called SIMPLE_MUSHROOM selecting only a subset of the attributes of the original table MUSHROOM.

Figure 2 shows the corresponding pre-processing query, followed by a query that adds an identifier to every example in the new table (it is worth noticing that the chosen attributes are known to be important for classification). The mining views automatically created for table SIMPLE_MUSHROOM can be visualized in the screenshot in Figure 3.

Step 2: Mining over decision trees.

In this step, we look for decision trees over table SIMPLE_MUSHROOM, targetting the attribute class and with maximum accuracy among those trees of size \( \leq 5 \).

The query is shown in Figure 4. The subquery selects the maximum accuracy achieved by the trees with size \( \leq 5 \) and accuracy \( \geq 90\% \) (the latter constraint is added in order to prune the search space of possible trees). The main query creates a table containing the trees with the characteristics mentioned above, having the pre-selected maximum accuracy. We use views Concepts and Treescharac in order to retrieve the concepts of the trees along with their characteristics. In the end, 3 trees are stored in table BEST_TREE. All of them have an accuracy of 95% and 5 nodes.

Step 3: Post-processing 1.

Having learned the trees with maximum accuracy in the previous step, we now want to explore the predictive capacity of these trees. Since all trees have the same accuracy, we choose the most balanced one, presented in Figure 5. This tree has treeid equal to 6.

Step 4: Post-processing 2.

Next, we want to store in the database the examples in SIMPLE_MUSHROOM that are misclassified w.r.t. the selected tree. To classify a new example using that tree, one simply looks up the concept that covers the new example. More generally, if we have a test set \( S \), all predictions of the examples in \( S \) are obtained by equi-joining \( S \) with the semantic representation of the tree given in the view Concepts. We join \( S \) to Concepts using a variant of the equi-join that requires that either the values are equal or there is a wildcard value ‘?’.

Figure 6 shows the corresponding query. To suit our purposes, the equi-join is made between tables SIMPLE_MUSHROOM and BEST_TREE, from which the tree with treeid = 6 is selected. The misclassified examples are those for which the prediction is different from the real class (stated in the last line of the query).

Step 5: Post-processing 3.

In this final step, we are interested in describing the mis-
create table misclassified_mushroom
as select test.*
    from simple_mushroom test, best_tree tree
where (test.cap_color = tree.cap_color
    or tree.cap_color = ' ?')
and (test.odor = tree.odor
    or tree.odor = ' ?')
and (test.gill_color = tree.gill_color
    or tree.gill_color = ' ?')
    and (test.size = tree.size
    or tree.size = ' ?')
    and (test.spore.print_color = tree.spore.print_color
    or tree.spore.print_color = ' ?')
and tree.treeid = 6
and test.class <> tree.class

create table mushroom_status
as select s.*, '0' as "status"
    from simple_mushroom s
    where exists
    (select * 
     from misclassified_mushroom m
     where s.id = m.id)
union
select s.*, '1' as "status"
    from simple_mushroom s
    where not exists
    (select * 
     from misclassified_mushroom m
     where s.id = m.id)

Figure 6: Query selecting the misclassified mushrooms.

classified examples obtained in the former step. To this end, first we create table MUSHROOM_STATUS in which every example in table SIMPLE_MUSHROOM is labeled as well classified (status = 1) or misclassified (status = 0). The query is shown in Figure 7.

select S.supp, S.sz, R.conf,
    C1.spore.print_color, C1.cap.color,
    C1.gill.size, C1.odor, C1.color = 
    "green", "purple"
from mushroom_status_sets S,
    mushroom_status_sets S1,
    mushroom_status_sets S2,
    mushroom_status_sets C1,
    mushroom_status_sets C2
where R.cida = S1.cid
and R.cidc = S2.cid
and C1.cid = R.cida
and C2.cid = R.cidc
and S.supp >= 15
and R.conf >= 100
and S.sz = 2 //total size of the rules
and S1.sz = 1 //size of the consequent
and C2.sz = 1
order by C2.size

Figure 8: Query over association rules.

Figure 9 shows a screenshot with the rules output by the query in Figure 8. As we are interested in describing the misclassified examples, we focus on the rules having consequent “status=0”. Clearly, these rules reveal odor=“musty”, gill.color=“green”, cap.color=“green”,“purple”), and spore.print.color=“green”,“purple”) as discriminative features to explain the misclassifications.

This ends the data mining scenario. In conclusion, we have developed a prototype of an inductive database based on virtual mining views. The advantages of our system are threefold: Firstly, the data are mined where they are located: in the database. Secondly, the user can specify in a declarative way (SQL queries) the patterns or models in which he or she is interested. Finally, thanks to the flexibility of ad-hoc querying of our system, the output of some queries can be used as input for subsequent queries.

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6. REFERENCES