

Practical Problems and Solutions in Hospital Information System Data Mining

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Abstract. Information mining from textual data becomes a very challenging task when the structure of the text record is very loose without any rules. Doctors often use natural language in medical records. Therefore it contains many ambiguities due to non-standard abbreviations and synonyms. The medical environment itself is also very specific: the natural language used in textual description varies with the personality creating the record (there are many personalized approaches), however it is restricted by terminology (i.e. medical terms, medical standards, etc.). Moreover, the typical patient record is filled with typographical errors, duplicates, ambiguities, syntax errors and many nonstandard abbreviations.

This paper describes the process of mining information from loosely structured medical textual records with no apriori knowledge. The paper concerns mining a large dataset of ~50,000–140,000 records × 20 attributes in relational database tables, originating from the hospital information system (thanks go to the University Hospital in Brno, Czech Republic) recording over 11 years. This paper concerns only textual attributes with free text input, that means 650,000 text fields in 16 attributes. Each attribute item contains approximately 800–1,500 characters (diagnoses, medications, anamneses, etc.). The output of this task is a set of ordered/nominal attributes suitable for automated processing that can help in asphyxia prediction during delivery.

The proposed technique has an important impact on reduction of the processing time of loosely structured textual records for experts.

Note that this project is an ongoing process (and research) and new data are still received from the medical facility, justifying the need for robust and fool-proof algorithms.

In the preliminary analysis of the data, classical approaches such as basic statistic measures, word (and word sequence) frequency analysis, etc., have been used to simplify the textual data and provide a preliminary overview of the data. Finally, an ant-inspired self-organizing approach has been used to automatically provide a simplified dominant structure, presenting structure of the records in the human readable form that can be further utilized in the mining process as it describes the vast majority of the records.

Keywords: Swarm Intelligence, Ant Colony, Text Data Mining, Information Retrieval, Hospital Information System.

1 Introduction

1.1 Motivation

In many industrial, business, healthcare and scientific areas we witness the boom of computers, computational appliances, personalized electronics, high-speed networks, increasing storage capacity and data warehouses. Therefore a huge amount of various data is transferred and stored, often mixed from different sources, containing different data types, unusual coding schemes, and seldom come without any errors (or noise) and omissions. Massively parallel distributed storage systems are used nowadays to provide computational nodes with data in reasonable time.

There are also problems with on-time data availability for a computational node. Especially in text processing, the impact of automated methods is crucial. In contrary to classical methods, nature-inspired methods offer many techniques, that can increase speed and robustness of classical methods.

1.2 Nature Inspired Methods

Nature inspired metaheuristics play an important role in the domain of artificial intelligence, offering fast and robust solutions in many fields (graph algorithms, feature selection, optimization, clustering, feature selection, etc). Stochastic nature inspired metaheuristics have interesting properties that make them suitable to be used in data mining, data clustering and other application areas.

In the last two decades, many advances in the computer sciences have been based on the observation and emulation of processes of the natural world. The origins of *bioinspired informatics* can be traced to the development of perceptrons and artificial life, which tried to reproduce the mental processes of the brain and biogenesis respectively, in a computer environment [1]. Bioinspired informatics also focuses on observing how the nature solves situations that are similar to engineering problems we face.

With the boom of high-speed networks and increasing storage capacity of database clusters and data warehouses, a huge amount of various data can be stored. *Knowledge discovery* and *Data mining* is not only an important scientific branch, but also an important tool in industry, business and healthcare. These techniques target the problematic of processing huge datasets in reasonable time – a task that is too complex for a human. Therefore computer-aided methods are investigated, optimized and applied, leading to the simplification of the processing of the data. The main goal of computer usage is data reduction preserving the statistical structure (clustering, feature selection), data analysis, classification, data evaluation and transformation.

Ant Algorithms. Ant colonies inspired many researchers to develop a new branch of stochastic algorithms: *ant colony inspired algorithms*. Based on the ant metaphor, algorithms for both static and dynamic combinatorial optimization, continuous optimization and clustering have been proposed. They show many properties similar to the natural ant colonies, however, their advantage lies in incorporating the mechanisms, that allowed the whole colonies to effectively survive during the evolutionary process.

1.3 Knowledge Extraction

Several techniques to extract knowledge from raw data have been developed in the past. These techniques have various and multiple origins: some result from the statistical analysis of the data, the regressions, decision trees, etc.; some resulting from the artificial intelligence such as the expert systems, intelligent agents, fuzzy logic, etc.

Plenty of nature inspired methods are studied and developed in present. One category is represented by methods, that are inspired by the behavior of ant colonies. These methods have been applied to many problems (often NP-hard). Review can be seen in [6] and [2]. We concentrate on the state-of-the-art nature methods inspired by the social behavior of insect communities, by the swarm intelligence, brain processes and other real nature processes.

Text Extraction. The accuracy for relation extraction in journal text is typically about 60 % [7]. A perfect accuracy in text mining is nearly impossible due to errors and duplications in the source text. Even when linguists are hired to label text for an automated extractor, the inter-linguist disparity is about 30 %. The best results are obtained via an automated processing supervised by a human [9].

Ontologies have become an important means for structuring knowledge and building knowledge-intensive systems. For this purpose, efforts have been made to facilitate the ontology engineering process, in particular the acquisition of ontologies from texts.

Additionally, medical records are specific. Although the medical doctors are members of an association, no real measures relating the unification of terminology and disambiguation is really made. The level of semantic interoperability is low, making the automated information retrieval a really nontrivial task.

Doctors often use natural language in medical records. Therefore it contains many ambiguities due to non-standard abbreviations and synonyms. For example, the information of *diabetes mellitus* is often expressed as *DM*, *DM2*, *DM 2*, *dia II*, *DMII*, etc. Even the national nomenclature (a translation of the ICD10 system) contains nonstandard and ambiguous abbreviations. The medical environment itself is also very specific: the natural language used in textual description varies with the personality creating the record (there are many personalized approaches), however it is (not strictly) restricted by terminology (i.e. medical terms, medical standards, etc.).

2 Input Dataset Overview

The dataset consists of a set of approx. 50 to 120 thousand records (structured in different relational DB tables; some of them are not input, therefore the range is mentioned) \times approx. 20 attributes. Each record in an attribute contains about 800 to 1500 characters of text (diagnoses, patient state, anamneses, medications, notes, references to medical stuff, etc.). For textual mining, 16 attributes are suitable (contain sufficiently large corpus).

The database export is a 10+ year export from the hospital information system. Anonymization has been performed in order not to reveal the sensitive patient data. As the patient ID is considered sensitive data, the records and other signals available have been referenced using a MD5 hash.

The overview of one small (in field length) attribute is visualized in Fig. [1]. Only a subsample (about 5 %) of the dataset could be displayed in this paper, as the whole set would render into a incomprehensible black stain. The vertices (literals) are represented as colored circle, the size reflects the literal (i.e. word) frequency. Edges represent transition states between literals (i.e. the sequence of 2 subsequent words in a sentence/record); edge stroke shows the transition rate (probability) of the edge. The same holds for all figures showing the transition graph, only a different visualization approach has been used.

It is clear, that human interpretation and analysis of the textual data is very fatiguing, therefore any computer aid is highly welcome.

3 Motivation

The task of this work is to provide the researchers with a quick automated or semi-automated view on the textual records. Textual data are not easy to visualize. The word frequency method is simple, but did not provide easily interpretable data. A frequency of multiple words is also a valuable input, however contains many duplications and does not really contribute in the process of definition of regular expressions for further mining. Therefore we decided to extract information in the form of a transition graph.

Such graphs allow as to induce a set of rules for information retrieval. These rules serve for extraction of (boolean/nominal) attributes from the textual rules. These attributes are used in automated rule discovery and can be further used for recommendation. The overall goal of the project is asphyxia prediction during delivery. High asphyxia might lead to several brain damage of the neonate and when predicted, caesarean section might be indicated on time.

4 Graph Explanation

In this paper we describe *transition graphs*. These are created for each attribute. An attribute consists of many records in form of a sentence. By *sentence* we hereby mean a sequence of literals, not a sentence in a linguistic form. The records are compressed – unnecessary words (such as verbs *is*, *are*) are omitted.

In this paper, only the attribute describing the anesthetics during deliveries visualized, as it is the simplest one.

Vertices of the transition graph represent the words (separated by spaces) in the records. For each word (single or multiple occurrence) a vertex is created and its potency (number of occurrences is noted). For example, the words *mesocaine*, *anesthetics*, *not*, *mL* form a vertex. Note that also words as *mesocain*, *mezokain* and other versions of the word *mesocaine* are present. For a number (i.e. sequence of digits) a special literal `_NUMBER_` is used.

Edges are created from single records (sentences entered). For example the sentence *mesocaine 10 mL* would add edges from vertex *mesocaine* to vertex `_NUMBER_` and from vertex `_NUMBER_` to the vertex *mL* (or the edge count is increased in case it exists). For all records, the count of the edges is also useful. It provides an overview on the inherent structure of the data – the most often word transitions.

Note that only a small subsection of the records of only one attribute is visualized in the paper. When displaying all records, the graphs are unprintable in the common paper formats and usually render as a black stain, therefore the ink-to-information ratio is very high. But it is totally unreadable. Images are supplied in a vector format, so they should be zoomed in correctly.

5 Nature Inspired Techniques

Social insects, i. e. ant colonies, show many interesting behavioral aspects, such as self-organization, chain formation, brood sorting, dynamic and combinatorial optimization, etc. The coordination of an ant colony is of local nature, composed mainly of indirect communication through pheromone (also known as *stigmergy*, the term has been introduced by Grassé et al. [8]), although direct interaction communication from ant to ant (in the form of antennation) and direct communication have also been observed [11].

The high number of individuals and the decentralized approach to task coordination in the studied species means that ant colonies show a high degree of parallelism, self-organization and fault tolerance. In studying these paradigms, we have high chance to discover inspiration concepts for many successful meta-heuristics.

5.1 Ant Colony Optimization

Ant Colony Optimization (ACO) [6] is an optimization technique that is inspired by the foraging behavior of real ant colonies. Originally, the method was introduced for the application to discrete and combinatorial problems.

Ant Colony Methods for Clustering. Several species of ant workers have been reported to form piles of corpses (cemeteries) to clean up their nests. This aggregation phenomenon is caused by attraction between dead items mediated by the ant workers.

This approach has been modeled in the work of Deneubourg et al. [5] and in the work of Lumer and Faieta [10] to perform a clustering of data.

ACO_DTree Method. The ACO_DTree method is a hybrid evolutionary approach for binary decision tree construction [4]. The tree is induced using the known data and can be further used for unsupervised clustering later: each leaf of the classification tree can be interpreted as a cluster. The algorithm uses a population of classification trees that is evolved using an evolutionary approach. Creation of the trees is driven by a pheromone matrix, which uses the ACO paradigm.

This approach has been utilized (with improvements and adaptation to the specific problem area) to simplify the structure of the vast dataset by finding the most important state transitions between literals, producing a probabilistic transitional model. The output structure is presented to the analyst for further processing/iteration.

For clustering, the ACO_DTree method [4,3] and ACO inspired clustering [10] variations have been successfully used. A self-organizing map has also been tested, but performed poorly.

New solutions are constructed continuously. It is a stochastic decision process based on the pheromone amount sensed by the ants. As in nature, the pheromone slowly evaporates over time (over iterations) in order to avoid getting stuck in local minimum and to adapt to dynamically changing environment. Daemon actions represent *background* actions which consist mainly of pheromone deposition. The amount is proportional to the quality of solution (and appropriate adaptive steps).

Main parameters of the algorithm are (with major importance to the method proposed): pheromone lay rate, pheromone evaporate rate, number of solutions created (number of ants), number of iterations, etc.

6 Automated Processing

Automated layout of transition graph is very comfortable for an expert, however the contents of the attribute is so complicated, that a human intervention is inevitable. Examples of automated layout can be seen in Fig. [1].

The figure Fig. [1] shows a transitional graph where only positioning based on the word distance from the sentence start is used. Although it might look correct, note that the same words are mispositioned in the horizontal axis.

7 Expert Intervention

A human intervention and supervision over the whole project is indiscutable. Therefore also human (expert) visualization of the transition graph has been studied.

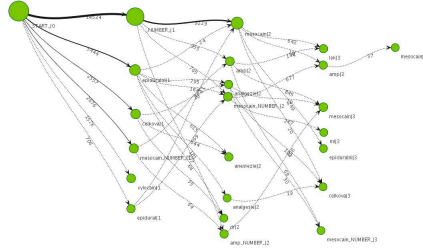


Fig. 1. A fully automated transition graph showing the most important relations in one textual attribute. No clustering has been used. The layout is based on the word distance from the start of the sentence. Note the mis-alignment of the similar/same words. Refer to section [2].

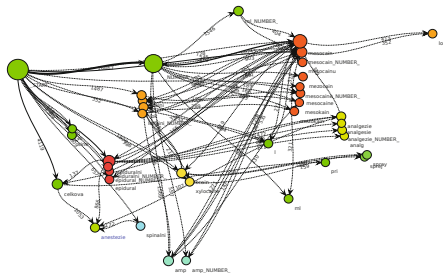


Fig. 2. An expert (human) organized transition graph (sub-graph) showing the most important relations in one textual attribute. Refer to section [2].

The vertices in a human-only organization are (usually) organized depending on the position in the text (distance from the starting point) as they have the highest potency. Number literal (a wildcard) had the highest potency, as many quantitative measures are contained in the data (age, medication amount, etc.). Therefore it has been fixed to the following literal, spreading into the graph via multiple nodes (i.e. a sequence *mesocain 10 mL* become two vertices – *mesocain_NUMBER_* and *mL*). This allowed to organize the chart visualization in more logical manner. Time needed to organize such graph was about 5–10 minutes. The problem is that the transition graph contains loops, therefore the manual organization is not straightforward.

An aid of a human expert has been used in semi-automated approach (see Fig. [3]) where the automated layout has been corrected by the expert. The correction time has been about 20–30 seconds only.

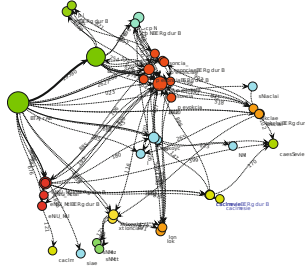


Fig. 3. A semi-automated (corrected by a human expert) organized transition graph showing the most important relations in one textual attribute. Refer to section [2].

8 Results and Conclusion

The main advantage of the nature inspired concepts lies in automatic finding relevant literals and group of literals that can be adopted by the human analysts and furthermore improved and stated more precisely. The use of induced probabilistic models in such methods increased the speed of loosely structured textual attributes analysis and allowed the human analysts to develop lexical analysis grammar more efficiently in comparison to classical methods. The speedup (from about 5–10 minutes to approx 20–30 seconds) allowed to perform more iterations, increasing the yield of information from data that would be further processed in rule discovery process. However, the expert intervention in minor correction is still inevitable. The results of the work are adopted for rule discovery and are designed to be used in expert recommendation system. A secondary output of this project is the gained knowledge for design of interoperable medical systems.

9 Discussion and Future Work

The future work is to evaluate the DB analyst’s utilization and aid of such graphs in more accurate way. The graphs serve as a bases for extraction rule proposal. However the only relevant measure is the time to reorganize the transitional graphs. The subjective opinion is very expressive and is not coherent. Next, the semantic meaning of the attributes will be extracted and verified followed by rule discovery mining.

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References

1. Adami, C.: Introduction to Artificial Life. Springer (1998)
2. Blum, C.: Ant colony optimization: Introduction and recent trends. *Physics of Life Reviews* 2(4), 353–373 (2005)
3. Bursa, M., Huptych, M., Lhotska, L.: Ant colony inspired metaheuristics in biological signal processing: Hybrid ant colony and evolutionary approach. In: *Biosignals 2008-II*, vol. 2, pp. 90–95. INSTICC Press, Setubal (2008)
4. Bursa, M., Lhotska, L., Macas, M.: Hybridized swarm metaheuristics for evolutionary random forest generation. In: *Proceedings of the 7th International Conference on Hybrid Intelligent Systems 2007 (IEEE CSP)*, pp. 150–155 (2007)
5. Deneubourg, J.L., Goss, S., Franks, N., Sendova-Franks, A., Detrain, C., Chretien, L.: The dynamics of collective sorting robot-like ants and ant-like robots. In: *Proceedings of the First International Conference on Simulation of Adaptive Behavior on From Animals to Animats*, pp. 356–363. MIT Press, Cambridge (1990)
6. Dorigo, M., Stutzle, T.: *Ant Colony Optimization*. MIT Press, Cambridge (2004)
7. Freitag, D., McCallum, A.K.: Information extraction with hmms and shrinkage. In: *Proceedings of the AAAI Workshop on Machine Learning for Information Extraction* (1999)
8. Grasse, P.P.: La reconstruction du nid et les coordinations inter-individuelles chez *bellicositermes natalensis* et *cubitermes* sp. la thorie de la stigmergie: Essai d'interpretation des termites constructeurs. *Insectes Sociaux* 6, 41–81 (1959)
9. Lafferty, J., McCallum, A., Pereira, F.: Conditional random fields: Probabilistic models for segmenting and labeling sequence data. In: *Proceedings of the ICML*, pp. 282–289 (2001); text processing: interobserver agreement among linguists at 70
10. Lumer, E.D., Faieta, B.: Diversity and adaptation in populations of clustering ants. In: *From Animals to Animats: Proceedings of the 3th International Conference on the Simulation of Adaptive Behaviour*, vol. 3, pp. 501–508 (1994)
11. Trianni, V., Labella, T.H., Dorigo, M.: Evolution of Direct Communication for a *Swarm-bot* Performing Hole Avoidance. In: Dorigo, M., Birattari, M., Blum, C., Gambardella, L.M., Mondada, F., Stützle, T. (eds.) *ANTS 2004*. LNCS, vol. 3172, pp. 130–141. Springer, Heidelberg (2004)