



## A Pattern Categorization Based Association Mining for Medical Databases

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### Abstract

Data mining is an effective method to find frequent patterns. In this paper we proposed, pattern categorization based Association mining for medical databases. Cancer data symptoms are consider for experimental purpose. We are using association rule for identifying valid and potentially useful patterns of symptoms for the medical database. Applying association rule we can generate individual and combine support of the symptom. Based on the support we find most frequent symptom in any patient after applying the association on 3 -30 days observation. Then we apply categorization so that we find the exact position of the symptom and calculate the diseases support. By considering the disease support we predict the disease. According to our experimental results we can achieve better prediction.

**Keywords:** Data Mining, Apriori, Association Rule, Medical Diagnosis

### 1. INTRODUCTION

The last few years Data Mining has become more and more popular. Together with the information age, the digital revolution made it necessary to use some heuristics to be able to analyze the large amount of data that has become available. Data Mining has especially become popular in the fields of forensic science, fraud analysis and healthcare, for it reduces costs in time and money. In this paper we use data mining to emphasize to discover knowledge that is not only accurate, but also comprehensible for the user [1], [2], [3]. According to Yang Jianxiong et al. [4] Comprehensibility is important whenever discovered knowledge will be used for supporting a human decision. After all, if discovered knowledge is not comprehensible for a user, it will not be possible to interpret and validate the knowledge. The Healthcare industry is among the most information intensive industries. Medical information, knowledge and data keep growing on a dailybasis.

It has been estimated that an acute care hospital may generate five terabytes of data a year [5]. The ability to use these data to extract useful information for quality healthcare is crucial.

Medical informatics plays a very important role in the use of clinical data. In such discoveries pattern recognition is important for the diagnosis of new diseases and the study of different patterns found when classification of data takes place. It is known that “Discovery of HIV infection and Hepatitis type C were inspired by analysis of clinical

courses unexpected by experts on immunology and hepatology, respectively” [6].

Data mining explores the hidden relationships and secret knowledge that cannot be observed and evaluated by the human beings easily and improves the quality of our lives by helping the experts showing the s cret relationships and correlations in the large databases [7][8].

Data mining has been played an important role in the intelligent medical systems[9][10]. The relationships of disorders and the real causes of the disorders and the effects of symptoms that are spontaneously seen in patients can be evaluated by the users via the constructed software easily. Large databases can be applied as the input data to the software by using the extendibility of the software. The effects of relationships that have not been evaluated adequately have been explored and the relationships of hidden knowledge laid among the large medical databases have been searched in this study by means of finding frequent items using candidate generation. The sets of sicknesses simultaneously seen in the medical databases can be reduced by using our non candidate approach.

The remaining of this paper is organized as follows. We discuss Association Rule Mining in Section 2. In Section 3 we discuss about preparation and stages . In section 4 we discuss about Recent Scenario. In section 5 we discuss about the proposed approach. Conclusions are given in Section 6. Finally references are given.

### 2. Recent Scenario

In 2011, M. Chaudhary et al. [11] proposed new and more optimized algorithm for online rule generation. The

advantage of this algorithm is that the graph generated in our algorithm has less edge as compared to the lattice used in the existing algorithm. The Proposed algorithm generates all the essential rules also and no rule is missing. The use of non redundant association rules help significantly in the reduction of irrelevant noise in the data mining process. This graph theoretic approach, called adjacency lattice is crucial for online mining of data. The adjacency lattice could be stored either in main memory or secondary memory.

The idea of adjacency lattice is to pre store a number of large item sets in special format which reduces disc I/O required in performing the query.

In 2011, Fu et al. [12] analyzes Real-time monitoring data mining has been a necessary means of improving operational efficiency, economic safety and fault detection of power plant.

Based on the data mining arithmetic of interactive association rules and taken full advantage of the association characteristics of real-time test-spot data during the power steam turbine run, the principle of mining quantificational association rule in parameters is put forward among the real-time monitor data of steam turbine.

Through analyzing the practical run results of a certain steam turbine with the data mining method based on the interactive rule, it shows that it can supervise steam turbine run and condition monitoring, and afford model reference and decision -making supporting for the fault diagnose and condition-based maintenance.

In 2011, Xin et al. [13] analyzes that use association rule learning to Process statistical data of private economy and analyze the results to improve the quality of statistical data of private economy. Finally the article provides some exploratory comments and suggestions about the application of association rule mining in private economy statistics.

In 2011, K. Zuhtuogullari et al. [14] proposed an extendable and improved itemset generation approach which has been constructed and developed for mining the relationships of the symptoms and disorders in the medical databases .

The algorithm of the developed software finds the frequent illnesses and generates association rules using Apriori algorithm. The developed software can be usable for large medical and health databases for constructing association rules for disorders frequently seen in the patient and determining the correlation of the health disorders and symptoms observed simultaneously.

In 2012, AshutoshDubey et al. [16] proposes an efficient method for knowledge discovery which is based on subset and superset approach. In this approach we also use dynamic minimum support so that we reduce the execution time. A frequent superset means it contains more transactions than the minimum support. It utilizes the concept that if the item set is not frequent but the superset may be frequent which is. It utilizes the behav-

ior that the less count may be frequent if we attached the less count with the higher order set.

Here we also provide the flexibility to find multiple minimum supports which is useful for comparison with associated items and dynamic support range. Our algorithm provides the flexibility for improved association and dynamic support. Comparative result shows the effectiveness of our algorithm.

In 2011, AshutoshDubey et al. [17] proposed a novel algorithm named Wireless Heterogeneous Data Mining (WHDM).

The entire system architecture consists of three phases:

- 1) Reading the Database.
- 2) Stores the value in Tbuf with different patterns.
- 3) Add the superset in the list and remove the related subset from the list.

Finally they find the frequent pattern patterns or knowledge from huge amount of data. They also analyze the better method or rule of data mining services which is more suitable for mobile devices.

In 2012, DevashriRaich et al. [18] introduce various intelligent computing techniques used for the medical diagnosis of diseases and a brief description about Nephritis and how its diagnosis could be done.

They suggested that computer science is getting more and more involved in medicines and health services. Various AI techniques and soft computing techniques are used for the diagnosis of particular diseases for the betterment of patient health. Various clinical decision support systems are also been devised by the help of AI.

In 2015, MohitSachan et al. [15] proposed a study on data mining. They suggest that the main aim of data mining is to extract useful patterns from huge amount of data. For this purpose some effective techniques like Apriori algorithm is presented and focus on the drawback.

To remove the above drawback, they present an improved non candidate single and multiple association approach for mining medical databases. The developed approach generates association rules for determining the relationships among the diseases observed synchronously.

The generated association rules are too significant for making early diagnosis for the correlated diseases. Some types of diseases can have triggering effects on different kinds of diseases. The symptoms and diseases which have stronger effect on each other can be determined and interpreted by the constructed system and the large and extended databases can be scanned effectively with the pruning property of the developed system.

### 3. Proposed Approach

In this study an improved association approach is used with single and multiple associations on symptoms observed on cancer diseases. The developed algorithm shows the relationships of the symptoms observed to-

gether by generating the item sets and constructing association rules using the frequent generation approach. The algorithm for the proposed concept in finding significant patterns for cancer prediction is presented in this section. In this framework administrator add the database, admin first add the disease and then the symptoms of the disease. According to the added value the parameter for that particular cancer is decided. This aim of our research is to apply the Association Rule Mining algorithm on patient symptoms for an efficient detection of Medical databases . In this we took cancer symptoms as medical database.

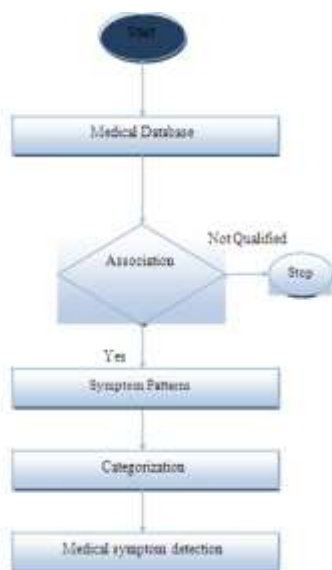


Figure 1: Flowchart of Association with Pattern Categorization

The flowchart of figure 1 shows the actual phenomena of our work. In this paper we took cancer symptoms as the cancer database. User selects the symptoms as he/she observed as shown in Figure 2. User enters the observation as per the days. of observation selected which is shown in figure 3. User enters 1 if he/she observe the symptom otherwise 0 as shown in figure 4. This can be converted in the tabular form as shown in figure 5.



Figure 2: Symptoms

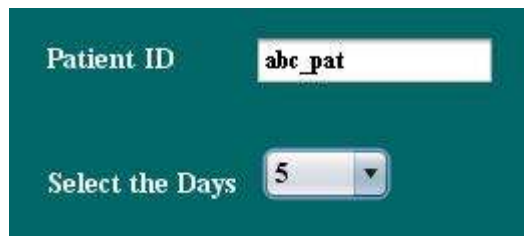


Figure 3: Observation Period

Figure 4: Observed Symptoms by the Patient

| TNo | FATIGUE | SWELLING | FEVER | NAUSEA | CHEST_PAIN |
|-----|---------|----------|-------|--------|------------|
| D1  | 1       | 1        | 1     | 0      | 1          |
| D2  | 1       | 1        | 1     | 0      | 1          |
| D3  | 1       | 1        | 1     | 1      | 1          |
| D4  | 1       | 0        | 1     | 1      | 1          |
| D5  | 1       | 1        | 0     | 0      | 1          |

Figure 5: Observed Symptoms

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Initialize: K = 1, C1 = all the 1- item sets;

Step 1: C1 to determine L1.
Step 2: L1 := {frequent 1- item sets}; k:=2; //k represents the pass number//
Step 3: while (Lk-1 ≠ ∅) do
begin
Step 4: Ck = gen_candidate_itemsets with the given Lk-1
Step 5: Prune (Ck)
Step 6: for all candidates in Ck do
count the number of transactions by using intersect method that are common in each item I ∈ Ck
Step 7: Lk := All candidates in Ck with minimum support ; k := k + 1;
end
Step 8: Frequent pattern = ∪k Lk ;

Step 9: General strategy: for each set find rule set that covers all instances in it (excluding instances not in the class). This approach is called a categorization approach because at each stage a rule is identified that covers some
  
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of the instances.

Step 10: General to specific rule induction : For each class C  
 Initialize E to the instance set  
 While E contains instances in class C  
 Create a rule R with an empty left-hand side that predicts class C  
 While R covers instances from classes other than C do:  
 For each attribute A not mentioned in R, and each value v,  
 Select A and v to maximize the accuracy Add (A = v) to R  
 Remove the instances covered by R from E

Step 11: Specific to general rule induction:  
 Pick up an instance and generalize it by repeatedly dropping conditions. Stop when all further generalizations lead to covering instances from other classes. Save the generalized instance as a rule.  
 Remove all instances covered by R and continue until all instances are covered.  
 When dropping conditions choose the ones that maximize rule coverage.  
 Problems: rule overlapping, rule subsumption.

Step 12: Then we find support based on the category , that is sup(x) of an itemset x is defined as the proportion of transactions in the data set which contain the item set.

Step 13: Discovery the percentage in the medical data set

Step 14: End

Figure 6: Algorithm for Association with Pattern Categorization

After the observed symptoms as obtained in figure 5. We apply our algorithm that is association with pattern categorization[Figure 6]. First we find the category of the individual as well as in the combination as shown in Figure 7, Figure 8, Figure 9 ,Figure 10 and figure 11. So that we achieve the frequent pattern list according to the support as shown in Figure 12.



Figure 8: Cancer Detection Pass2

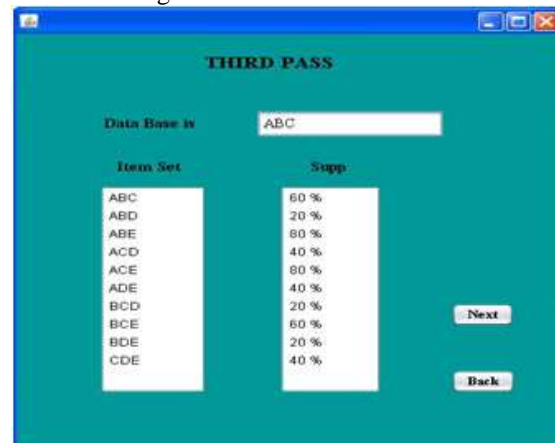


Figure 9: Cancer Detection Pass3

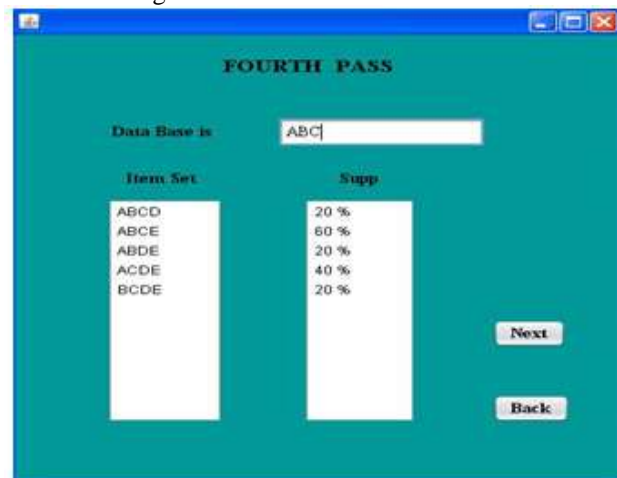


Figure 10: Cancer Detection Pass4

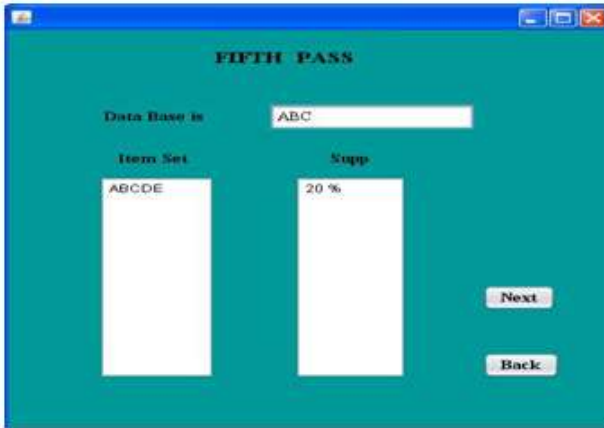


Figure 11: Cancer Detection Pass5

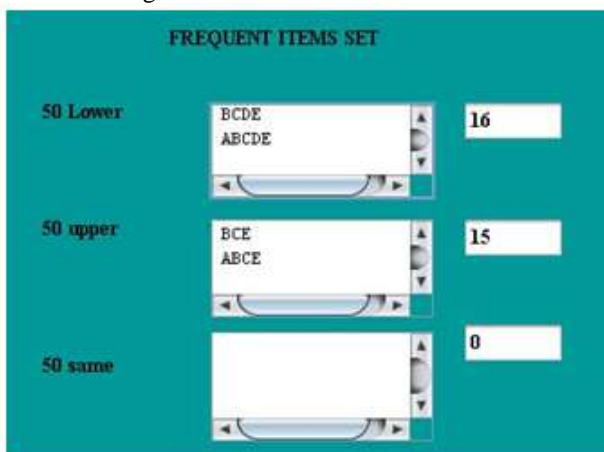


Figure 12: Frequent Item Set

#### 4. Result analysis

Then we apply categorization according to three diseases having highest support. In this we categorize by creating these disease as a separate class. Then we generate to specific rule induction for each class C. We initialize E to the instance set while E contains instances in class C. Then we create a rule R with an empty left-hand side that predicts class C. While R covers instances from classes other than C then iterate it. For each attribute A not mentioned in R, and each value v. Select A and v to maximize the accuracy Add (A = v) to R

Remove the instances covered by R from E. Then we pick up an instance and generalize it by repeatedly dropping conditions. Stop when all further generalizations lead to covering instances from other classes. Save the generalized instance as a rule. We also remove all instances covered by R and continue until all instances are covered. When dropping conditions choose the ones that maximize rule coverage.

Then we find support based on the category, that is  $sup(x)$  of an itemset x is defined as the proportion of transactions in the data set which contain the item set.



Figure 13: Disease Wise List

| Ser. No.           | Symptom     | Percentage |
|--------------------|-------------|------------|
| 1                  | Fatigue     | 100.0      |
| 2                  | pain_bones  | 0          |
| 3                  | Swelling    | 80.0       |
| 4                  | blood_clots | 0          |
| 5                  | chest_pain  | 100.0      |
| 6                  | Hoarseness  | 0          |
| 7                  | Fever       | 80.0       |
| 8                  | Weight_Loss | 0          |
| <b>LUNG_CANCER</b> |             | <b>45</b>  |

Figure 14: Chances of Cancer detection

| Ser. No.            | Symptom       | Percentage |
|---------------------|---------------|------------|
| 1                   | Nausea        | 40.0       |
| 2                   | Headache      | 0          |
| 3                   | Blurry_vision | 0          |
| 4                   | Fever         | 80.0       |
| 5                   | Seizures      | 0          |
| 6                   | Mental_Capa   | 0          |
| <b>BRAIN_CANCER</b> |               | <b>20</b>  |

Figure 15: Chances of Cancer detection

As shown in figure 14 and figure 15 we find the support weight as generated from the association. And according to the weight we then calculate support based on the symptom because it is dynamic in nature. And we achieve the chance of the symptom as shown in the figure 14 and figure 15. According to the final detection we



show the result as shown in figure 16. As per our observation our detection is better, because we comprise all individual items.

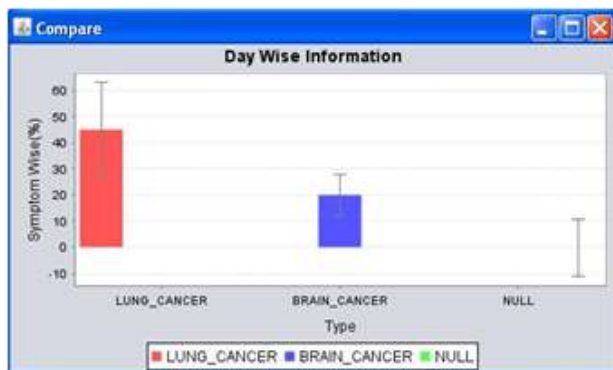


Figure 16: Cancer Chart

## 5. Conclusion

In addition to the classical approaches, the constructed approach can calculate the association rules from the desired item set number and this specification gives the system the opportunity to generate different association rules. In this paper we find the better association on medical databases. For experimental analysis we taken the cases of cancer symptoms. Based on our observations and result shown in the result analysis we provide better prediction of medical databases.

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