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Frequent Document Mining Approach through Clustering

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Abstract-Now days, finding the association rule from large number of item-set become very popular issue in the field of data mining. To determine the association rule researchers implemented a lot of algorithms and techniques. FP-Growth is a very fast algorithm for finding frequent item-set. This paper, give us a new idea in this field. It replaces the role of frequent item-set to frequent sub graph discovery. It uses the processing of datasets and describes modified FP-algorithm for sub-graph discovery. The document clustering is required for this work. It can use self-similarity function between pair of document graph that similarity can use for clustering with the help of affinity propagation and efficiency of algorithm can be measure by F-measure function.

Keywords: clustering, document-graph, FP-growth, graph mining, frequent sub graphs clustering.

I. INTRODUCTION

Data mining techniques is used to extracting information from that data. Many researchers developed a lot of algorithm and techniques for finding useful information in the database. Frequent item-set mining is a core data mining operation and has been extensively studied over last decade. It plays an essential role in many important data mining tasks. Algorithms for frequent item-set mining form the basis for algorithms for a number of other mining problems, including association rule mining, correlations mining, and mining sequential and emerging patterns. Association rule mining technique is very effective data mining technique to finding the useful hidden information in the data, its aim to extract correlation, frequent pattern, association by transaction database or other data repository. This rules generated by datasets and it derives by measurement of support and confidence of every role, which define the frequency of that role. Association rule is a rule which is depends on the association relationship of objects and items. For example data item interrelation ship which is occurs simultaneous with other data items. This rule is calculated by data and association calculated with the help of probability. Share market and recommended system etc. are its practical applications.

To concept of document based clustering, association rule mining is very effective algorithm and useful approach. It utilizes that FP-Growth approach discovers frequent patterns and modified it to frequent sub graph discovery. Originally this algorithm is design to frequent item-set mine in market basket analysis. In this paper, analyse FP-Growth approach to document clustering. For graph mining, change in FP-Growth algorithm which discover frequent connected graph and perform better for thick connected graph. This Paper consists of five main sections. The second section describes the Basic Theory of this work. The frame work of this approach is describes in third section, Experimental Result analysis is forth section and finally section five is devoted to conclusion and future work.

II. BASIC THEORY

The association rule mining [2] approaches were established to determine frequent item-sets in market basket datasets [4].The two most common approaches to association rule mining are Frequent Pattern growth (FP-growth) [3]and apriori [5].The apriori algorithm uses prior information of frequent item-sets to produce the candidates for larger frequent item-sets. It relies on interactions between item-sets and subsets. If an item-set is frequent, then all of its subsets must also be frequent. But generating candidates and checking their support at each level of iteration can become expensive. FP-growth introduces a different approach here. Instead of generating the candidates, it compresses the database into a compact tree form, known as the FP-tree, and extracts the frequent patterns by traversing the tree.

In this work, exchanged the concept of the frequent item-set-in to frequent sub graph. However, there are some famous sub graph discovery systems like FSG (Frequent Sub graph Discovery) [6], g Span (graph-based Substructure pattern mining) [7], DSPM (Diagonally Sub graph Pattern Mining) [8], and SUBDUE [9]. It does allow us to believe that the concept of the creation of document-graphs and discovering frequent sub graphs to execute a sense-based clustering is currently feasible. All these systems deal with multiple aspects of efficient frequent sub graph mining. Most of them have been tested on real and artificial datasets of chemical compounds. Yet none of them has been applied to the mining of text data. It's developed a graph-based approach for



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document clustering using FP-growth. Since in this system the documents are represented by graphs, made the necessary changes to the FP-growth algorithm so that instead of generating frequent item-sets it can generate frequent sub graphs.

III. FRAME WORK OF APPROACH

Our overall method is briefly portrayed in this section. The system is divided into four major parts: Document Pre-processing, FP-Growth approach, Document clustering and Similarity Measures. Detailed mechanisms are described in the following subsections.

A. Document Pre-processing

The work use stemming to repossess the relationship between the keywords of a document. In stemming, words are related to each other based on different semantic relationships among them. This Paper use the relation of keywords, which is a super-ordinate or IS-A (“a kind of”) relationship. This work believes that, for sense-based document clustering, this relation is more suitable than the others because it provides the generalization of a concept or a word. It builds a document-graph by traversing the hierarchy of the keywords up to Entity. Once there have all of the document-graphs, create a Master Document Graph (MDG_{ph}) [11] based on the traversal of all of the keywords up to the topmost level.

Now use the MDG_{ph} for a number of reasons. Instead of taking the entirety of Words as the background knowledge. Use MDG_{ph} that contains the ontology related to the keywords only. This allows us to focus on the area of the Words which is applicable to our dataset. Another purpose why use the MDG_{ph} is that it can facilitate the use of the DFS codes. The code is produced by traversing the MDG_{ph} in Depth First Search (DFS) order. This gives us the DFS traversal order (DFS code) for the entire MDG_{ph} . The DFS codes of the edges of the MDG_{ph} are applied to the edges of the document-graphs. Thus, the DFS codes of the edges of the MDG_{ph} help us to mark each edge of the document-graphs in a DFS order specific manner. Therefore, the same sub graph acting in two document-graphs can be identified by the DFS codes of the edges of that subgraph. Additionally, the MDG_{ph} helps in testing the connectivity between nodes during the sub graph discovery process. The graph representation of the documents to a table consisting of the list of edges appearing in each document. This helps us to fit the frequent sub graph discovery task to the frequent item-set discovery problem of the market basket data analysis. Each edge of a document-graph is considered to be an item of a transaction, a sub graph is considered to be an item-set and each document is considered to be a transaction.

B. FP-growth Approach for Frequent Sub graph Discovery

The original FP-growth algorithm, when applied to our problem generates a set of frequent edges which do not necessarily constitute to a connected sub graph. Generation of all possible frequent patterns not only outputs all possible frequent sub graphs but also generates a lot of overhead in the form of all possible sets of frequent, but disconnected, edges. This causes unnecessary costs during the mining of the document-graphs as only looking for frequent sub graphs and these sets of frequent disconnected edges tolerate no useful information for the document clustering. The time and space required to generate and store these disconnected frequent edges have negative impact on the overall performance of the FP-growth approach. To overcome these problems it restricts the discovery of frequent item-sets in the FP-growth approach. Instead of discovering all possible frequent combination of edges we only discover the frequent edges that founds to a graph. We also control the maximum size of the frequent sub graph by the SP Threshold value.

The method Filter DGB reconstructs both the lists transaction DGB and header Table by eliminating the in frequent edges based on the minimum support provided by the user. It sorts the edges of transaction DGB and header Table in descending order by support. After this step, the top of the header covers the most frequent edges and the bottom contains the least frequent ones. The edges at the top level of the header table are the representative edges of the topmost levels of our MDG_{ph} . Since these edges denote very abstract relationships between synsets [10], they appear in too many document-graphs implying that they are not good candidates for clustering. In contrast, edges at the bottom of the header table are least frequent and denote relations between very specific concepts in the MDG_{ph} . Since they appear in very few document-graphs, they too provide less information for clustering. This motivated us to crop the edges a second time from the top and bottom of the header table before constructing the FP-tree. transactionDB is updated accordingly to reflect this change in the



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header table. After this refinement, create the FP-tree by calling the `FPTreeConstructor ()` method. Later, `FP-Mining ()` generates the frequent sub graphs by traversing the FP-tree.

In the original FP-growth approach a node from an FP-tree indicates an item. In that case, a node of the FP-tree contains the DFS-code of an edge. If the original FP-growth approach is directly used in graph-mining, it does not guarantee the connectivity property of a subgraph. It can generate a disconnected set of edges in a representation of a frequent subgraph. The difference between the original FP-growth approach and our modified approach is that the system crops the single paths of the FP-tree and maintains connectivity of edges for discovering frequent sub graphs.

If at any point, a single-path is encountered in the FP-tree, we crop nodes from the top of the single path based on the user-provided threshold $SPThreshold$. Removing a node from the single path refers to eliminating the corresponding edge denoted by that node. $SP Threshold$ provides control to the number of combinations of edges appearing in a single path. Depending on the connectivity of the edges, a combination of the edges may or may not generate a frequent sub graph. Let b be a sub graph for which a single path is generated by traversing the branches of the FP-tree ending with b . This says that the discovery of the newly joint frequent sub graphs is conditional on the frequent sub graph b . The supports of these new sub graphs are determined by the support of b before the merging.

The depth of the MDG_{ph} can reach up to 18 levels, which is the maximum height of the hierarchy of Words. Since our document-graphs contain hundreds of edges, the depth of the FP- tree can reach up to hundreds of levels depending on the number of edges in a document-graph. Conventional FP-growth generates all possible subsets of a single path for every edge-conditional FP-tree. Instead of accepting all possible combinations of a single path it only keeps the combinations of edges that generate connected frequent sub graphs. Whenever a single path is encountered in the FP-tree (or recursively generated conditional FP-trees), each of its combinations is generated and checked to make sure that it follows the connectivity constraint. This is done by first taking one of the edges from the combination of the single path and then adding it to a list called `connected List`. In the `Iterate ()` method, a neighbouring edge list, `neighborListi` is created for an edge i using the MDG_{ph} . The `neighborListi` is checked to see if it contains any edge from the combination of the conditional FP-tree's single path. If there are such edges it followed the same procedure for all of them until no new edges from the combination of the single path are added to the `connected List`. At the end, the size of the `connected List` is compared with the size of the combination. If both of their sizes are the same, then the whole combination must be connected generating a subgraph a . Then an attempt is made by step 6 through 9 to combine the subgraph b with subgraph a . The method is `Connected ()` returns true if a and b can be merged together to generate a connected subgraph. It should be noted that `connected` method checks for the connectivity of one combination at a time. If the edges under consideration of this combination do not compose a connected subgraph, but compose multiple disconnected sub graphs, then some other combinations of the single path will generate these smaller connected sub graphs. So, do not lose disjoint but smaller connected sub graphs of a larger disconnected combination of the single path.

Additionally, control the single path length by using $SP Threshold$ so that our FP-growth approach performs faster. The length of the single path can be as large as the number of all the edges appearing in a document-graph. Taking each combination of a large single path and checking the connectivity constraints is computationally intensive. The construction of the FP-tree forces the edges with higher frequencies to appear at the top of the FP-tree. So, it is more likely that nodes at the top levels of the FP-tree indicate edges at more abstract levels of the MDG_{ph} . After observing that a higher abstraction level of the MDG_{ph} does not provide enough information for reliable clustering, we restricted the generation of combination of a single path of the FP-tree to a certain length. ($SPThreshold$) When he mines the FP-tree, he starts from the edges appearing the least in the pruned header table. Thus, if he reaches a single path of length greater than $SPThreshold$, he prunes the upper part of the single path above the threshold and generates each combination of the lower part only. This mechanism prunes nodes of single paths of the FP-tree at the upper levels which are representative of top level edges of the MDG_{ph} .

C. Document Clustering

This used the discovered frequent sub graphs to cluster the document-graphs. These sub graphs can be viewed as concepts appearing frequently within the documents. If it evaluate the similarity of documents based on the co-occurrence of frequent sub graphs, and then use these similarity values to cluster the documents, it will get a

sense-based clustering of the text documents. He uses Hierarchical Agglomerative Clustering (HAC) [12] for the clustering phase of that work. Now it implements the Group Average method to cluster the documents where the distance between two clusters is defined by the average distance between points in both clusters.

A number of similarity measures exist that can be used to find the closest or most distant pair of documents to merge during HAC. Among them, the cosine measure [13] is the most frequently used one. It penalizes less in cases where the number of frequent sub graphs on each document differs significantly. Since the cosine measure focuses more on the components in the documents and is not influenced by the document length, it has been used widely in document clustering.

D. Similarity Measures

In this work chose this measure to compute the similarity between two document-graphs (DG1 and DG2) based on the frequent sub graphs (FS) appearing in them:

$$\text{Similarity}_{\text{cosine}}(\text{DG1}, \text{DG2}) = \frac{\text{count}(\text{FS}(\text{DG1}) \cap \text{FS}(\text{DG2}))}{\sqrt{\text{count}(\text{FS}(\text{DG1})) * \text{count}(\text{FS}(\text{DG2}))}}$$

To cluster the documents use a dissimilarity matrix which stores the dissimilarity between every pair of document-graphs using the formula, dissimilarity = 1 – similarity. The value of dissimilarity can range from 0 to 1. It can use self-similarity in work in place of similarity function this self-similarity can be define as K-means algorithm adopts the widely used similarity measurement Cosine coefficient, namely equation [1]. Because k-means needs to compute on the vector space model, for $X = (x_1, \dots, x_n)$ and $Y = (y_1, \dots, y_n)$, the component form of their similarity measurement can be expressed as:

$$S(X, Y) = \frac{\sum_{i=1}^n x_i y_i}{(\sum_{i=1}^n x_i^2)^{\frac{1}{2}} (\sum_{i=1}^n y_i^2)^{\frac{1}{2}}}$$

Where n is the number of the features in the whole vector space. Similar with k-means, AP (CC) and SAP (CC) also use Cosine coefficient (equation (2)) as the similarity between two documents. However, unlike k-means, a document in AP (CC) and SAP (CC) does not need to be represented into the whole vector space, but only into its own vector space. Therefore the similarity measurement computation complexity of the latter two algorithms is reduced greatly in respect to the one of k-means. The self-similarities of AP (CC) are defined as:

$$s(i, i) = \min_{1 \leq i, j \leq N, i \neq j} \{s(i, j)\} - \phi \left(\max_{1 \leq i, j \leq N, i \neq j} \{s(i, j)\} - \min_{1 \leq i, j \leq N, i \neq j} \{s(i, j)\} \right)$$

Where ϕ is an adjustable factor?

This similarity can be used for clustering with help of affinity propagation and efficiency of algorithm can be measured be F-Measure function using precision and recall value and Entropy function.

IV. EXPERIMENTAL RESULT ANALYSIS

A. Experimental Results of Modified FP Growth Approach

Table 1 contains some of our experimental results with the subset of 768 documents from 7 different groups of items. The avg. min_sup was kept 49.2 and we selected 568 items from the middle of the 768 documents to use them in our FP-growth approach.

Table 1- Parameters of Mining FP-Tree and Generate association rules.

No. of Groups	Min_sup	FP-Tree Storage (Bytes)	FP-Tree updates	FP-Tree Nodes	Generating Time (Seconds)	T-Tree Storage (Bytes)	No. of Frequent Sub graph Nodes
10	2.0	580	59	24	0.39	8992	639
25	5.0	602	186	25	0.44	8516	607
50	10.0	1284	364	56	0.34	8800	623
101	20.2	1702	759	75	0.27	8284	590



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200	40.0	2516	1534	112	0.33	8816	629
568	113.6	3770	4473	169	0.3	8360	596
768	153.6	4386	6090	197	0.3	8460	604

The Table 1 shows the outputs of FP-Growth algorithm in which taken a combination of groups like 10, 25, 50, 101, 200, 568, 768, and so on. This table shows the variation of minimum support, FP-Tree Storage in bytes, FP-tree updates after pruning nodes in the header table, No. of FP-Tree Nodes, time for generating association rules in seconds, T-Tree storage in Bytes and last No. of frequent Sub graph Nodes. Here take an experimental result of table 1 and plot a graph of minimum support value of all the no. of groups which show the efficiency of minimum support of FP-Growth algorithm.

B. Experimental Results of Document Clustering

We used the frequent sub graphs find by the FP-growth approach in the earlier section to cluster the corresponding documents took dissimilar numbers of keywords from each dataset for the clustering. To examine the accuracy of our clustering, we analyse the results with the commonly used frequency-based clustering technique. Table 2 contains our experimental results of document clustering with affinity propagation for different documents. Here table shows 8 different data files clustering outputs of documents.

Table 2 - Parameters of document files generated in clustering.

No. of Files	5	10	20	30	40	50	60	70
No. of Identified	1	2	6	8	12	15	18	19
Data point (net similarity)	5.7 e ⁻ ₃₀₆	4.4 6	12. 69	20. 32	25. 99	31. 88	39. 07	47. 55
Preference of selected exemplars	0	0	4.9 1	6.6 2	9.6 0	9.9 0	14. 26	15. 68
No. of iterations	126	131	114	108	121	114	122	120
Elapsed Time	2.5	2.7 2	2.2 2	2.1 4	2.3 9	2.2 9	2.6 4	2.4 1
No. of cluster	1	2	6	8	12	15	18	19
Fitness (net similarity)	5.7 e ⁻ ₃₀₆	4.4 6	17. 66	26. 95	35. 59	41. 79	53. 33	63. 23

Table 2 shows the overall parameters of document clustering which shows that when clustering method apply on the document file, the number of iteration and elapsed time in this method will be constant and overall process take average iteration for all clustering is 119.5 and average time is 2.41375 second for every clustering. Now consider result with respect to files and elapsed time, graph of elapsed time and documents is shown below.

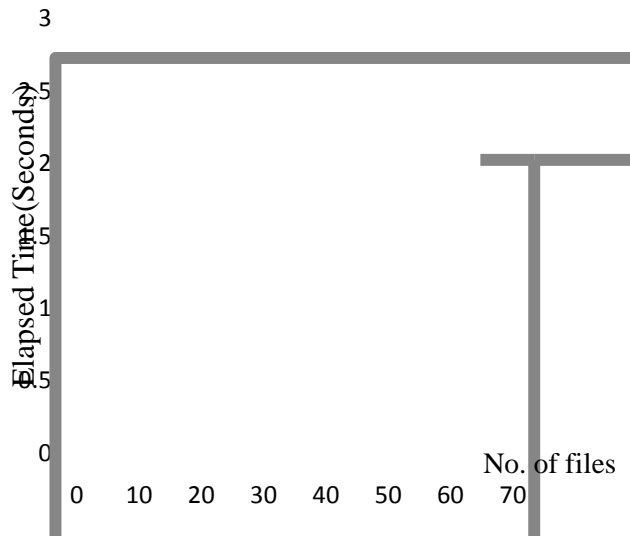


Fig 1 - Elapsed time in second for documents in clustering.

Now consider result with respect to files and no of iteration, graph of no. of iteration and document files is shown below.

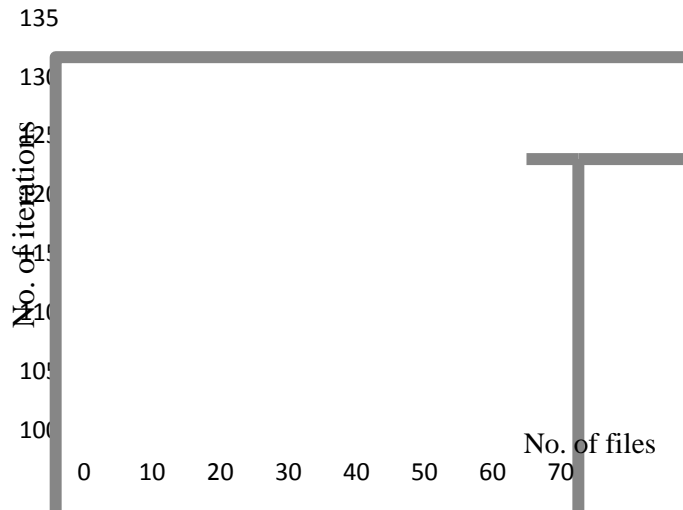


Fig 2 - No. of iterations for documents in clustering

For the number of documents, if we raise the number of keywords, the MDG_{ph} has the affinity to contain more edges. More keywords better illuminate the concept of the documents. In such case, the edges in the mid-level of the MDG_{ph} will have more relations between them. As a result, more sub graphs will appear in the middle of the MDG_{ph}. So, our methodology has the affinity to take the advantage of attachment of keywords by determining more sub graphs and using them in the clustering. It requires inclusion of thousands of keywords for better clustering.

V. CONCLUSION

In Data mining document clustering is very active research area. To find suitable information system much suitable idea has been implemented in document clustering. It is very challenging task to find human-like clustering. In this work, a graph based clustering with affinity propagation. That find a new way to clustering document based more on the keywords they contain document based clustering techniques mostly depend on the keywords. The work modifying the FP-mining algorithm to find the frequent sub graph with clustering affinity propagation in graph. It can evaluate some techniques for computing the combination of large scale paths which can improve the performance of mining algorithms. In future will plan to study more application related to that feature and try to implement better way and improve their performance.



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