International Journal of Engineering Applied Sciences and Technology, 2017
Vol. 2, Issue 3, ISSN No. 2455-2143, Pages 1-4
Published Online January-February 2017 in IJEAST (http://www.ijeast.com)

DESIGN OF HIERARCHICAL GENEALOGY WITH SUB GRAPH MINING

Miss. Monika Deshpandey
Department of Computer Science & Engineering,
G.H. Raisoni Academy of Engineering &Technology
Nagpur, Maharashtra, India

Mrs. Sonali Bodkhe (Assit.proff.)
Department of Computer Science & Engineering,
G.H. Raisoni Academy of Engineering &Technology
Nagpur, Maharashtra, India

Abstract-- **Whenever a new topic is searched by researchers, they go to literature review area for a new topic. Researchers are unfamiliar to all those topics are. To make literature survey they have to collect all information regarding domain which is relevant to topic but there are huge amount of data and number of papers present in the different citation. For researchers is difficult to study each and every papers with their references of same and different citation. To solve these problems, this paper is designed which is working on, how to find out relative papers with respected query. This paper will be focused on creation of their genealogy.First working process is to find out the relevant matching paper according to keywords which is given by researchers. This is the pre-processing part. After that user goes for, discrimination of survey paper and implementation of paper for this all papers have to extract according to seminal papers it will create genealogy of those paper, by making association and interlinking among all matching documents on the basis of references of each paper. This Genealogy will help user to get a quick look at which papers are relevant to given query of research, and association among them, so that user will focus on those documents only. In this proposed work user neither stray on unwanted documents nor waste the time for searching the particular topic, which may increases scalability and efficiency of searching keywords.**

Keywords— **Sub graph matching, Genealogy of papers, Analysis of data, Construction of sub graph, Yearly analysis of reference papers Construction of Efficient Genealogy.**

I. INTRODUCTION

Researching for a new topic, problem that researchers face is a literature survey problem for the topic which may useful or not. Literature Review put an important role on a desired topic for research, which shows, what are the latest issues are left to work on and also give direction to work in future scop. A research trend gives deep sense of the topic, i.e., what type of research has been done before and what the new issues are at this time. Once the preparation of literature survey it can predict what will be the current research trend, and that will give a hint to which direction to go. It will be successful perform research. It will make research more efficient and effective.

Through search button, it will display number of papers which is to be read or observe will be difficult for human being. Therefore, it is important to identify seminal papers on their search topic queried by user and find the relationships among them. Therefore, this work is necessary to find relevant papers on the research topic, which make relationships among those entire topics. And there should be a provision which isolates the survey paper from implementation paper.

There is N number of papers present which represented in citation. Every paper contains some keywords. These keywords are assigned by some weights for measuring information and match the keyword with relevant paper. For example if user wants to search a survey paper which is in social network citation, and other relevant paper is also in that citation so this method creates a graph “Qg.” Stands for query graph of a large graph G.

There are large amount of data present in web which shows different area of domain. This web not only shows textual information but also show the link documents. This link document shows lots of data which is useful for searching

Relevant data the RDF frame is using in this proposed method and RWR framework is also used to contain data in graphic form. This work mainly describe to create research paper genealogy which remove the difficulty of the literature survey at large contain and will help the researcher to easily take the movement of the research.

There three problems are Specify: for finding the relative papers, distinguish of survey paper and Implementation of paper, and genealogy creation of same topic from relevant papers. This proposed work will mainly help in analysing and visualising in this domain i.e. creating research paper genealogy which remove the difficulty of finding at large contain and will help the researcher to easily take the movement of the research topic. This proposed system will give quick result for searching keyword with better visual presentation.

II. RELATED WORK

The inherent computational complexity arises. Finding Random graphs have been generated using the multiplicative linear random number generator algorithm finding a homomorphic image of a pattern graph in a target graph for this some algorithms are applied which is useful to remove unsuccessful mapping which retrieves sub graphs that are structurally isomorphic to the query graph, and meanwhile satisfy the condition of vertex pair matching with weighted (dynamic) set similarity. [1]. Graph X-Ray (G-Ray), a fast method that finds sub- graphs that either match the desirable query pattern exactly, or as well as possible.

Traditional SQL-based methods, as well as more re-cent graph indexing methods, will return no answer when an exact instance of a pattern does not exist. This phenomenon is interception for finding intermediate nodes. The non-intermediate nodes will be referred to as matching nodes. Observing the effectiveness and efficiency is low in this method for finding graph index by index; using some new techniques to find efficient data in less time [2].This paper presents a novel technique for approximate matching of large graph queries.

There is proposing a novel indexing method that incorporates graph structural information in a hybrid index structure. TALE is a general tool for approximate sub graph matching queries, and can be easily customized to meet the requirement of different applications. Our empirical evaluations demonstrate the improved effectiveness and efficiency of TALE over existing methods for searching the data into large nodes in a specific domain [3].

This is an approximate sub graph matching technique that finds sub graphs in the database that are similar to the query, allowing for node mismatches, node gaps, and graph structural differences. Torque [4] is a topology working with genealogical graphs is no exception in this senses the approaches of the first category follow the main idea behind the Apriori algorithm for mining frequent item sets. More specifically, they rely on the apriori property, according to which all the sub patterns of a frequent sub graph pattern are also frequent.

Thus, to enumerate candidate patterns, they apply breadth-first search to generate sub graphs of size (k+1), by joining two sub graphs of the previous level. Graph visualization can help to form an overview of relational patterns and detect data structure much faster than data in a tabular form. The form in which the graph is presented has a significant impact on how the graph is understood and the time that is necessary to achieve this.[5] In this approach all nodes placed close to one another might be interpreted by the user as a true relationship whether or not this relationship exists. Some graphic algorithms are used in this proposed work which 2D rotating and scale matrices.

That takes time to generate graph in parent child form [6]. A query language for graph databases which supports graphs as the basic unit of information. Sun et al. utilized graph exploration and parallel computing to process sub graph matching query on a billion node graphs. This is one of good approach for making graphs with number of nodes which would be prepare as an online in could [7]. The quality of fan charts is immediately appealing, but unfortunately it does not grow fast enough. In the layout of this family tree that emphasizes temporal data.

The ancestors and descendants are laid out radially around a centred person. The layout also supports dynamic interaction with the family tree. Rendering of large data is needed in this method it is Observed. And make good densely populated family trees [8]. A SPARQL query is converted into a corresponding sub graph matching query. For speeding up query processing, development of a novel index, together with some effective pruning rules and efficient search algorithms.

The RDF (Resource Description Framework) data model was proposed for modeling Web objects as part of developing the se- mantic web. RDF format used to support structural queries. This approach work on speed of the query searching of training set. Recently, an efficient and robust sub graph isomorphism algorithm. Finding book keeping cost is reducing. DBpedia is one of an approach for extracting structure information. Those are present in the web, all information contain by the Wikipedia.

This is work in sophisticated queries which link with data set which available in the web in could server, interlinking methods are used in this project [9].It is used in static data. Weighted data string similarity queries are useful. In this applications like data cleaning and integration. For finding approximate matches in the presence of typographical mistakes, multiple data formatting conventions, of data transformation errors. Index construction and updating is hard. Because heaviest first algorithm for weighted intersection based on prefix and suffix lists [10].

An algorithm for pattern-matching on arbitrary graphs that is based on reducing the problem of finding a ho-momorphic image of a pattern graph in a target graph, to that of finding ho-momorphic images of every connected component of the pattern in the target. Finding depiction of relationships in a large family is challenging, as is generally the case with large graphs. For analysing the nature of genealogical graphs, characterized difficulties to draw and presented novel graphical representations for them [11].

An algorithm used to find a set of seminal papers on a given topic that construct a genealogy of the seminal paper by using the influence measure and citation information.TurboISO [12] was proposed graph alignment algorithms for biological networks, which can be used to solve isomorphism problems. Approximate sub graph matching query usually concerns the structure information and allows some of the vertices or edges not being matched exactly. Closure-tree [13] is the first graph index that supports both sub graph queries and graph similarity queries.

However, a query graph is much smaller than the data graph in subgraph isomorphism problems, while the two graphs usually have similar size in graph alignment problems [14].To solve subgraph isomorphism problems, graph alignment algorithms introduce additional cost as they should first find candidate subgraph of similar size from the large data graph [15]. In addition, existing exact subgraph matching and graph alignment algorithms do not consider weighted set similarity on vertices, which will cause high post processing cost of set similarity computation [16].

While doing research for Similarity measure which is an important issue in proposed design reviewed, Cosine Similarity for text based measures, which is used for text clustering along with K-Clustering Method for similarity computation on the basis of terms in abstract, keyword and body of paper [17]. Again for Link based measure, which takes help of citations in paper for similarity computation between papers, for that purpose reviewed?

The performance and scalability issues are sophisticated in algorithm. The parent child relationship [18] had generated. When question arises to search data which is in linked together for this keyword query routing approach is used for searching linked data. Novel method is used in this project to improve quality of keyword searching. Multilevel ranking method is used in this project for finding relevant information. For making efficient graph some graph joined methods like merge join are used in this approach, this approach creates the complexity of nodes when combination elements are increased. Ranking keyword routing plan is useful to show efficient data in graph form [19].

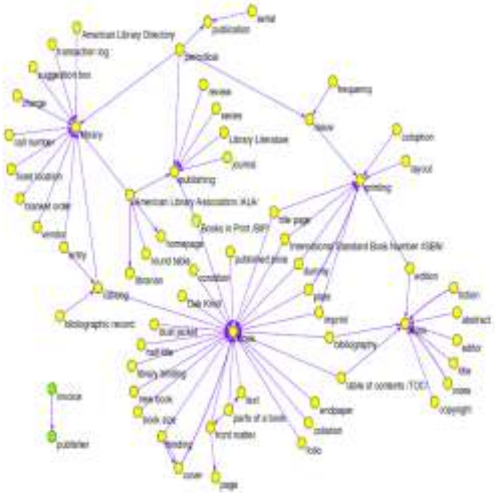

Fig1- ODLIS dictionary Graph

ODLIS is stands for Online Dictionary of Library and Information Science. This is useful to find out data in Information technology and computer categories. This is an online library where any query is typed by user and relevant information is extracted by this dictionary. This all process is done in the form of graph. This is one of the sections which can manage all data and information for searching queries. This is an idea to create graph which shows relevant data according to user query.

III. PROPOSED WORK

In proposed system, It will find out all research papers in the same domain, then it will find out all seminal paper in same domain which influence current research topic mostly then it will distinguishes between good research paper and original research papers of high quality. For that purpose it will identify survey papers. After that it will find out Genealogy of papers which depicts relationship among papers to show influence relation among them.

This genealogy may be including with Survey paper and all other research paper or may be excluding survey paper and including all other papers according to user wish. When a researcher insert a interested research topic keyword. In its database all sample paper will be available; it will match that keyword with all papers in database. Then it will use clustering algorithm to extract papers belonging the same topic. After that it will parse all document which it get after applying clustering algorithm and it will take out its citation information from those paper.

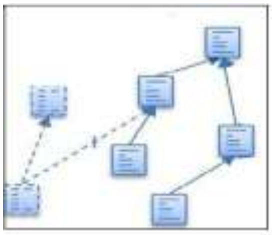

Fig2-genealogy construction of relevant Paper.

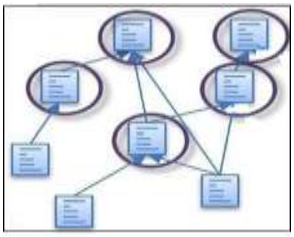

Fig3-Finding relevant Paper

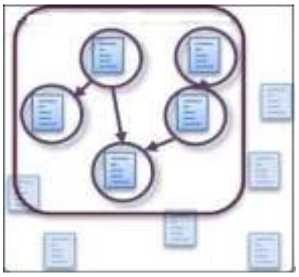

Fig4- construction of relevant Paper

Then it will apply ranking algorithm on citation information, from that it will assign ranking score to each paper, and on that basis which papers influence topic most and which papers are not. Analyses of different phases are used to generate graphs. Some mock-ups are using to analysis all phases. By using of architecture design and implementation part will be created. Database design is the most important part of this project. Lots of data required to show the genealogy of seminal papers. Those all data would be contains of many papers which is collected is a data base, user can find out there important topic according to their choices.

IV. PROPOSED METHODOLOGY

After getting seminal paper including or excluding survey papers, it will try to find out relationship among papers on the basis of similarity measures, that means which paper citing which paper will be considered into it. But in earlier methods it was direction dependent that means it may be in-link or out-link with references to two papers which to be examined for relationship between them but in proposed system it will use direction independent method for similarity measure. It will find out genealogy among papers which will help for quick literature survey to find out current research trend. There are following steps to show the proposed work of this project which are-

1. Collection of data sets
2. Pre-processing the data sets
3. Classification of data sets
4. Reduce time to make graph.
5. Result and Analysis of efficiency.

In propose method have to create sample database, which consist number of documents, these documents are in form of pdf file.

Text Pre-processing is an important part for pre-processing of input keywords for asking query, so that it will extract only important and relevant word from input keywords. Some approaches take higher time complexity but gives a higher accuracy than others. This application has been used to determine the categories of documents on the basis of query. After getting all relevant matching documents, it moves for construction of genealogy among all matching documents.

There is a provision for user to choose paper type. Here we have two paper types one is Survey and another is Implementation, It's on user which type of paper they want to include in their research paper survey and according to that Genealogy will be created. Now what does mean of Genealogy, in Genealogy it will show interlinking between all documents, which we get after extracting all matching documents in previous step.

V. CONCLUSION

The proposed work is about to collection of relevant matching research .According to this proposed work, all papers with pre-processed query will be related to any type of content, which will help the user to make quick look at which papers are relevant to any topic of research, user will focus on those documents and literature review which is only useful for them, they wouldn't be work in unwanted document.

This would be helpful for researchers to get similar paper in short time, by this the process of searching multiple review papers with their citations will be easier with in less of time. This is one of the great solutions to create the genealogy with different citations in yearly form. This proposed work gives efficient result and high performance to search the seminal papers as well as relevant papers year wise in different citations.

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