Discovery of Emerging Topics by Co-citation Graph on the Web

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Abstract. Discovering new topics covering profitable items and ideas (e.g., mobile phone, global warming, human genome project, etc) is important and interesting. However, since we cannot completely encode the world surrounding us, it's difficult to detect such topics and their mechanisms in advance. In order to support the detection, we show a method for revealing the structure of WWW by using the KeyGraph algorithm. Empirical results are reported.

1 Introduction

In our daily lives, a new topic sometimes become suddenly popular. The topic might seem insignificant at first sight, however, it turns out to match potential needs of us. The Tipping Point[1] describes this kind of phenomenon where a 'little' thing can make a big difference in the future. However, we cannot detect new topics and their mechanisms in advance since we cannot completely decode the world surrounding us. Detection of a Tipping Point, in face of this obstacle, could be a big chance for our various activities because competitors are not aware of such new topics. We here interpret 'topics' in the broad sense that cover new items, problems, ideas, and so on (e.g., mobile phone, global warming, human genome project, etc).

These topics were born when new collaborations of existing interests satisfy our potential needs or demands. Although the hidden factors might be 'submerged' in human mind, we believe that a few signs can be mined from a database on human behaviors reflecting human mind. For this purpose, the web is an attractive information source for its size and sensitivity to trends. The web consists of an abundance of communities [2, 3], each corresponding to a cluster of web pages sharing common interest. Since a community means a chunk of shared interest, it is considered that a web page supported (or linked) from some communities satisfies their interests, and shows the movement direction of the widen human world. From this point of view, we are expecting the structure of WWW might be a key to understand the real world.

In this paper, we show a method for revealing the structure of WWW by using KeyGraph algorithm[8] to inspect that WWW reflects the real world, and that the revealed structure of WWW supports our detection of new significant topics.

2 KeyGraph Algorithm

KeyGraph[8] is originally an algorithm for extracting assertions based on co-occurrence graph of terms from textual data. The strategy of KeyGraph comes from considering that a document is constructed like a building for expressing new ideas based on traditional concepts. The processes of KeyGraph are composed by four phases.

- **0)Document preparation:** Prior to processing a document D, $stop\ words[5]$ which have little meaning are discarded from D, words in D are stemmed[6], and phrases in D are identified[7]. Hereafter, a term means a word or a phrase in processed D.
- 1) Extracting foundations: Graph G for document D is made of nodes representing terms, and links representing the *co-occurrence* (term-pairs which frequently occur in same sentences throughout D). Nodes and links in G are defined as follow:
 - **Nodes** Nodes in G represent high-frequency terms in D because terms might appear frequently for expressing typical basic concept in the domain. High frequency terms (HF) are the set of terms above the 30th highest frequency.
 - Links Nodes in HF are linked if the association between the corresponding terms is strong. The association of terms w_i and w_j in D are defined as

$$assoc(w_i, w_j) = \sum_{s \in D} \min(|w_i|_s, |w_j|_s), \tag{1}$$

where $|x|_s$ denotes the count of x in sentence s. Pairs of high-frequency terms in HF are sorted by assoc and the pair above the $(number\ of\ nodes\ in\ G)$ - 1 th tightest association are represented in G by links between nodes.

2) Extracting columns: The probability of term w to appear if all the foundations in G are considered by the author is defined as key(w), and the key(w) is defined by

$$key(w) = 1 - \prod_{g \in G} \left(1 - \frac{\sum_{s \in D} |w|_s |g - w|_s}{\sum_{s \in D} \sum_{w \in s} |w|_s |g - w|_s} \right).$$
 (2)

Sorting terms in D by keys produces a list of terms ranked by their association with cluster, and the 12 top key terms are taken for $high\ key\ terms$.

3)Extracting roofs: The strength of column between a high key term w_i and a high frequency term $w_i \subset HF$ is expressed as

$$column(w_i, w_j) = \sum_{s \in D} \min(|w_i|_s, |w_j|_s).$$
(3)

Columns touching w_i are sorted by $column(w_i, w_j)$, for each $high\ key\ term\ w_i$. Columns with the highest column values connecting term w_i to two or more clusters are selected to create new links in G. Finally, nodes in G are sorted by the sum of column of touching columns. Terms represented by nodes of higher values of these sums than a certain threshold are extracted as the keywords for document D.

3 Our Approach

By focusing on the analogy between a document and other textual data, KeyGraph can be applied to a variety of topics. For example, KeyGraph has been adopted to

- find areas with the highest risks of near-future earthquakes from data of observed past earthquakes[9],
- get timely files from visualized structure of our working history [10],
- construct planning to guide concept understanding in WWW[11],
- make tools for shifting human context into disasters[12],
- discover potential motivations and fountains of chances [13].

In a document D, high-frequency terms are used for expressing typical basic concept, and term-pairs which frequently occur in the same sentences mean strong association throughout D (see Sect. 2).

In this paper, we extend the use of KeyGraph to another kind of data, i.e., Web-page set (corresponding to D, document in Sect. 2) including Web-pages (each corresponding to a sentence in Sect. 2) having URL-links, each corresponding to words in Sect. 2. That is, high-frequency links (which are the URLs pointing to other web pages) in a collection W of web pages show popular web pages, and link-pairs which frequently occur in the same web pages show strong relations in W[4]. Our fundamental hypothesis here is that the occurrence of a document and a collection of web pages have common causal structures, and our strategy for applying KeyGraph is based on this analogy.

Let us be more formal. A web page(which URL is u) is translated to a sentence as

$$u u_1 u_2 u_3 \cdots u_i \cdots u_n. \tag{4}$$

Where $u_i(i = 1, 2, 3, ..., n)$ are the URLs contained in the web page. A document is formed by combining sentence, shown in eq. (4), for each web page of a collection. By this translation, we can obtain the document reflecting the link structure of WWW.

4 An Example of Experiment

In this section, we report on our experiment where we applied KeyGraph to two sets of collections C_A and C_B , each of which contains 500 popular web pages obtained by Google for the input query 'human genome', to follow the changes of the communities with time. The difference between the collections is the date: C_A is obtained on November 26, 2000, and C_B is on March 11, 2001.

After C_A and C_B were translated into two documents like Sect. 3, for each document KeyGraph output URLs as roof(asserted) keywords. The URLs for C_A and C_B are shown in Table 1. Comparing the output URLs for C_A and C_B in Table 1, we can recognize the movement among them. For example,

http://www.ncbi.nlm.nih.gov http://www.nhgri.nih.gov http://www.sanger.ac.ul

Table 1: Output UR	L lists of KeyGraph for collections C_A and C_B .
Output URLs for C_A	Affiliation
www.ncbi.nlm.nih.gov	National Center for Biotechnology Information
gdbwww.gdb.org	The Genome Database
www.ornl.gov	Oak Ridge National Laboratory
www.nhgri.nih.gov	The National Human Genome Research Institute
www.gene.ucl.ac.uk	The Galton Laboratory
www.ebi.ac.uk	European Bioinformatics Institute
www.gdb.org	The Genome Database
lpg.nci.nih.gov	CGAP Genetic Annotation Initiative
www.sanger.ac.uk	The Sanger Centre
www.genetics.utah.edu	Human Genetics Department in University of Utah
Output URLs for C_B	Affiliation
Output URLs for C _B	Affiliation National Center for Biotechnology Information
www.ncbi.nlm.nih.gov	National Center for Biotechnology Information
www.ncbi.nlm.nih.gov www.nhgri.nih.gov	National Center for Biotechnology Information The National Human Genome Research Institute
www.ncbi.nlm.nih.gov www.nhgri.nih.gov www.ornl.gov	National Center for Biotechnology Information The National Human Genome Research Institute Oak Ridge National Laboratory
www.ncbi.nlm.nih.gov www.nhgri.nih.gov www.ornl.gov www.cnn.com	National Center for Biotechnology Information The National Human Genome Research Institute Oak Ridge National Laboratory CNN.com
www.ncbi.nlm.nih.gov www.nhgri.nih.gov www.ornl.gov www.cnn.com genome.wustl.edu	National Center for Biotechnology Information The National Human Genome Research Institute Oak Ridge National Laboratory CNN.com Genome Sequence Center in Washington University
www.ncbi.nlm.nih.gov www.nhgri.nih.gov www.ornl.gov www.cnn.com genome.wustl.edu onhealth.webmd.com	National Center for Biotechnology Information The National Human Genome Research Institute Oak Ridge National Laboratory CNN.com Genome Sequence Center in Washington University OnHealth Network Company
www.ncbi.nlm.nih.gov www.nhgri.nih.gov www.ornl.gov www.cnn.com genome.wustl.edu onhealth.webmd.com www.gdb.org	National Center for Biotechnology Information The National Human Genome Research Institute Oak Ridge National Laboratory CNN.com Genome Sequence Center in Washington University OnHealth Network Company The Genome Database

Table 1: Output URL lists of KeyGraph for collections C_A and C_B

appear in both the outputs. These are the most authorized research institutes in the area of human genomics. On the other hand,

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http://www.tigr.org
http://www.celera.com
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appear only in URLs for C_B . These are newly growing research organizations in the area. This movement reflects events/situations of the real world in the topic of human genome, and means that KeyGraph could detect the major changes in the society. However, we cannot see why these changes occurred, from these tables.

In the field of human genome, revolutionary events were occured in 2000 and 2001. Celera Genomics and the U.S. Human Genome Project has been entered into a keen competition for leadership. Both two rivals independently accomplished the analysis of most of the human genome at almost the same time, and their papers were appeared in *Science*[15] and *Nature*[14] respectively in February 2001.

Considering these real events/situations, the emerging URLs(http://www.tigr.org, http://www.celera.com) are considered to reflect the real society.

5 Conclusions

In this paper, we introduced a method for aiding human awareness on significant novel, i.e., emerging topics. Here, the algorithm of KeyGraph is extended to be a method for the analysis and visualization of cocitations between Web pages. Communities, each having members (Web pages, their authors/readers) with common interests are obtained

as graph-based clusters, and an emerging topic is detected as a Web page relevant to multiple communities. Experiments, an example of which is presented in this paper, show that the aimed effect of our method is realized.

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