

Improving Focused Crawling with Genetic Algorithms

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ABSTRACT

The Web, containing a large amount of useful information and resources, is expanding rapidly. Web crawlers are one of the most crucial components in search engines and their optimization would have a great effect on improving the searching efficiency. Focused Crawlers can selectively retrieve Web documents relevant to a specific domain to build collections for domain-specific search engines. In this paper, we use a genetic algorithm with focused crawling for improving its crawling performance. Expands initial keywords by using a genetic algorithm for focused crawling. The results showed that our approach could build domain-specific collections with higher quality than traditional focused crawling techniques.

General Terms

Focused crawling, keyword enhance

Keywords

Crawling, focused crawling, Genetic Algorithm, web crawler

1. INTRODUCTION

A Web crawler is a key component inside a search engine [1]. It can traverse the Web space by following Web page's hyperlinks and storing the downloaded Web documents in local repositories that will later be indexed and used to respond to the user's queries efficiently [2]. However, with the huge size and explosive growth of the Web, it becomes more and more difficult for search engines to provide effective services to end-users. Moreover, such a large collection often returns thousands of result documents in response to a single query. It is impossible for major search engines to update their collections to meet such rapid growth. As a result, end-users often find the information provided by major search engines not comprehensive or out-of date.

To address the above problems, focused crawler were introduced. A focused crawler or topical crawler is a web crawler that attempts to download only web pages that are relevant to a pre-defined topic or set of topics. Focused crawling was first introduced by Chakrabarti et al.[7]. Most focused crawlers use the content of traversed pages to determine the next hyperlink to crawl. They use a similarity function to find the most similar page to the initial keywords that is already downloaded and crawl the most similar one in the next step. These similarity functions use information retrieval techniques [20] to assign a weight to each page so that the page with the highest weight is more likely to have the most similar content.

2. Related Works to Focused Crawlers

Chakrabarti et al. seem to introduce focused crawling for the first time. In the crawler described in their article [2], the user picks a subject from a pool of hierarchically structured example documents. The program learns the subjects by studying the examples, and generates subject models. These

models are used to classify web pages. The link structure is also considered by the crawler to discover hubs. Hubs are described by Kleinberg as high-quality lists that guide users to recommended authorities, and authorities are prominent sources of primary content on a topic. Links from hubs can be relevant even though the text on the hub page itself does not appear to be relevant. Semi automatic web resource discovery using ontology-focused crawling [20]

The crawler described by Chakrabarti et al. [2] uses example documents and machine learning principles. One difference with Diligenti's [1] crawler is that it generates a context graph that describes the link structure around all the seed documents. Diligenti's crawler only focuses on web pages. Ester's crawler and other crawlers that use a static initial set of example documents for classification are very dependent on the quality of the initial training data. Sizov et al. have built a focused crawler that aims to overcome the limitation of the initial training data there are some other experiments which measure the similarity of page contents with a specific subject using special metrics and reorder the downloaded URLs for the next crawl [3] or even evaluate a learning scheme for identifying which URL the spider should crawl next in order to increase the efficiency in topic specific web resource discovery [25]. Bing Liu et al combined the crawling strategy with clustering concepts [14]. For each topic they first retrieve a specific number of top weighted retrieved pages from Google for that topic and then extract some other keywords from them.

3. Genetic Algorithm

Genetic algorithms (GA) are search algorithms based on the principle of natural selection and genetics. GA operates on a population of potential solutions applying the principle of the survival of the fittest to produce better and better approximation to the solution of the problem that GA is trying to solve. At each generation, a new set of approximations is created by the process of selecting individuals according to their level of fitness value in the problem domain and breeding them together using the operators borrowed from the genetic process performed in the nature, i.e. crossover and mutation. This process leads to the evolution of populations of individuals that are better adapted to their environment than the individuals that they were created from, just as it happens in natural adaptation.

3.1 Genetic Algorithms in Information Retrieval

Genetic algorithms (GA) are search algorithms based on the principle of natural selection and genetics. GA operates on a population of potential solutions applying the principle of the survival of the fittest to produce better and better approximation to the solution of the problem that GA is trying to solve. Until 1980 there were no serious attempts in the field of applying genetic algorithms in information retrieval. Raghavan and Aggarwal were the first ones who introduced

the subject by trying to optimize document clustering by the means of genetic algorithms [26]. One year later, Gordon proposed a new approach for improving the document descriptors. In his experiments documents were represented by an array of keywords which evolved over time by natural selection and genetic operators, and the final results were generally proved to be the best string of keywords describing the document [27]. The trend was followed by Yang et al. for improving the weights of keywords associated with the document topic [28]. Petry et al. applied genetic algorithms to increase the functionality of retrieving data from a weighted indexed collection of documents, by modifying weights of query terms.

3.2 Fitness Evaluation

Fitness function is a performance measure or reward function which evaluate how good each solution is. Result from fitness functions are interval 0 to 1. By 1.0 means document and query is sameness. Values near 1.0 mean documents and query are more relevant and values near 0.0 mean documents and query are less relevant. We use Jaccard coefficient as fitness function for our genetic algorithm. We use binary term vector of Jaccard coefficient.

3.3 Chromosome Representation

Both documents and queries are represented by vector. A document vector (Doc) with n keywords and a query vector with m query terms can be represented as

Doc = (term1, term2, term3, ..., termn)

Query = (qterm1, qterm2, qterm3, ..., qtermm)

We use binary term vector. For example, user enters a query into Google search engine that could retrieve 10 documents. These documents are

Query = {Khap, Panchayat, Honour, Killing}

Doc1 = {Honour, Khap, Killing, Panchayat}

Doc2 = {Killing, Decide, Government}

.

.

.

Doc10={Killing, Honour, Government, Session, Monsoon}

All keywords of these documents can be arranged in the ascending order as

Against, Boys, Caste, Couples, Court, Decide, Family, Girls, Government, Haryana, Honour, Intra, Khap, Killing, Law, Marriage Monsoon, Panchayat, Session, Union, and Village.

Encode in the chromosome representation as

Doc1={0 0 0 0 0 0 0 0 0 0 1 0 1 1 0 0 0 1 0 0 0}

Doc2={0 0 0 0 0 1 0 0 1 0 0 0 0 1 0 0 0 0 0 0 0}

.

.

Doc10={00 0 0 0 0 0 0 1 0 1 0 0 1 0 0 1 0 1 0 0}

These chromosomes are called initial population that feed into genetic operator process. The length of chromosome depends on number of keywords of documents retrieved from user

query. From our example the length of each chromosome is 21 bits.

3.4 Selection

After we evaluate population's fitness, the next step is chromosome selection. Selection embodies the principle of 'survival of the fittest'. Satisfied fitness chromosomes are selected for reproduction. Poor chromosomes or lower fitness chromosomes may be selected a few or not at all.

3.5 Crossover

Crossover is the genetic operators that mix two chromosomes together to form new offspring. Crossover occurs only with some probability Pc (crossover probability). GA's construct a better solution by mixture good characteristic of chromosomes together. Higher fitness chromosomes have an opportunity to be selected more than the lower ones, so good solution always alive to the next generation.

For example, two chromosomes are crossover between position 6 and 13.

Doc1={0 0 0 0 0 0 0 0 0 0 1 0 1 1 0 0 0 1 0 0 0}

Doc2={0 0 0 0 0 1 0 0 1 0 0 0 0 1 0 0 0 0 0 0 0}

The resulting crossover yields two new chromosomes.

Doc1={0 0 0 0 0 1 0 0 1 0 0 0 0 1 0 0 0 1 0 0 0}

Doc2={0 0 0 0 0 0 0 0 0 0 1 0 1 1 0 0 0 0 0 0 0}

3.6 Mutation

Mutation involves the modification of the values of each gene of a solution with some probability Pm (mutation probability). In accordance with changing some bit values of chromosomes, give the different breeds. For example randomly at position 10 apply mutation.

Doc1={0 0 0 0 0 0 0 0 0 0 1 0 1 1 0 0 0 1 0 0 0}

Result {0 0 0 0 0 0 0 0 0 1 1 0 1 1 0 0 0 1 0 0 0}

4. Empirical Results

Table 1 depicts initial keywords and the terms added given by genetic algorithm, written in bold. We entered the old and new keywords (old keywords plus the new term added by genetic algorithms functions) into Google, and calculated the average relevance based on the 10 first pages returned for each instance. As it is shown, the new set of keywords achieved a higher relevance score. For the first sample, the task is to search for news about khap panchayat. After downloading about 10 pages, Genetic algorithm added "marriage" to its initial set. This word was chosen because in most of the downloaded pages there was about marriage in Haryana.

This experimentation tests for 10 queries with fitness functions Jaccard coefficient. Average relevance is defined by the following equation

$$fitness(d_j) = \frac{1}{n} \cdot \sum_{k=1}^n \left[\frac{|d_j \cap d_q|}{|d_j \cup d_q|} \right]$$

Jaccard coefficient

5. Conclusions

As the size of the Web keeps growing, it has become increasingly important to build high-quality domain specific search engines. This research has proposed a new crawling technique to build domain-specific collections for search engines that incorporate a global search algorithm, Genetic Algorithm, into the crawling process.

In our user study, our proposed Genetic Algorithm with focused crawling built collections with significantly higher quality than did a traditional best-first crawler. Our proposed work apply a genetic algorithm to the focused crawling process to find more relevant Web resources in order to overcome the problems faced by traditional focused crawlers. Our goal is to extend the keyword set for the focused crawling. In this way searching made easy, we find more relevant document or web pages. Results shows that average relevance of document increase upto 50%. When focused crawler having key set with more relevancy then retrieved data also more relevant for local collection of a search engine. It improves the crawling performance.

Table 1: A.R=average relevance

Old Keyword	New Added Term	A.R With Old Keyword	A.R With add new Keyword
Khaph Panchayat honour killing	Marriage	0.3081	0.3911
Fiber optics technology information	Light	0.4175	0.4729
Genetic algorithm optimization softcomputing	Fitness function	0.2646	0.3466
Micheal Jackson music mp3	Download	0.3616	0.5208
Mouse Disney movie	Walt mickey	0.3515	0.5400
Economy finance stock market financial	News	0.2232	0.3567
Health medicine Medical disease	Symptoms	0.2304	0.4197
Sql server dbms database	Data	0.3144	0.4321
Artificial intelligence neural network	Applicatio n	0.4136	0.5596

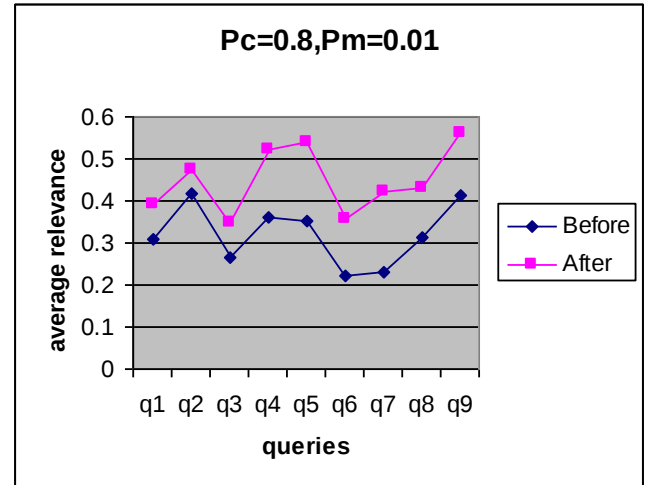


Fig 1

6. Future Work

There is still a lot of work to do for improving the efficiency of the focused crawling. There is also a downside to the algorithm. Some websites repeat one word their abbreviated name frequently. In this case, when the page content is really close to the search subject, the extracted keyword might be irrelevant to the initial keyword set. For example, when we were extracting keyword from a web page for news about the Khaph panchayat, we chose the google.com as the initial website and since the term was repeated so many times in almost all the pages that are not relevant to the query but related to that site like news, HT etc. This term was returned by the crawler as the keyword. This problem may be cumbersome to manage but attempts could be made to find solutions. Also, only one of the bits of the fittest chromosome would be added to the initial set and the other bits will be neglected. We intend to implement some simple learning procedures in order to find out which of these bits might be useful in the future. Work of selecting keyword using text analyzer tool is manual that is also a problem. It is also a future work make it automatically and online.

In conclusion, although the initial results are encouraging, there is still a long way to achieve the greatest possible crawling efficiency.

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