# Document Clustering based on Genetic Algorithm using D-Individual

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**Abstract.** Document clustering using genetic algorithm shows good performance. However the genetic algorithm has problem of performance degradation by premature convergence phenomenon. In this paper, we proposed the document clustering based on Genetic Algorithm using D-Individual (DIGA) to solve this problem. Genetic algorithm is based on the diversity of population and the capability to convergence. Success of genetic algorithm depends on these two factors. If we use these factors efficiently, we can get a better solution in reduced execution time. We apply DIGA to Reuter-21578 text collection and demonstrate the effect of our clustering algorithm. The results show that our DIGA has better performance than traditional clustering algorithms (K-means, Group Average and genetic algorithm) in various experiments.

### **1** Introduction

The researches of document clustering are actively progressing for analysis and application of mass information. The document clustering is to group the documents which are similar in a set of documents without prior information.

The clustering algorithms are classified to the hierarchical clustering algorithm and the non-hierarchical clustering algorithm based on grouping method [1],[2]. Kmeans clustering algorithm is one of the non-hierarchical clustering algorithms [3]. Group Average clustering is one of the hierarchical clustering algorithms [2]. These two clustering algorithms are fast in execution and easy to understand but do not achieve the highest results [1],[2]. To solve it, genetic algorithm which is one of artificial intelligent algorithms has been applied to the document clustering and it shows good results [4],[5].

Genetic algorithm is optimization algorithm using natural selection and evolution operation. Genetic algorithm depends on the diversity of individuals and the capability to convergence [4],[5],[6]. Capability to convergence of genetic algorithm has weakness that is premature convergence phenomenon. The premature convergence phenomenon is cause of depress the performance of genetic algorithm [7].

In this paper we propose the document clustering based on genetic algorithm using the individuals which have the feature information of document clusters. The algorithm has better performance and shorter execution time than the general Genetic Algorithm. This paper is organized as follows. The next section describes the document clustering using genetic algorithm. Section 3 presents the principle of DIGA. Section 4 explains experiment setting, evaluation approaches, results, and analysis. Section 5 concludes and discusses future work.

### 2 Document Clustering using Genetic Algorithm

Genetic Algorithm(GA) is based on the principles of the natural selection and the survival of the fittest [6],[8]. Basic components of GA are gene, chromosome, individual and fitness function. Individual is set of chromosomes characterized by genes. Basic operations of GA are selection, crossover and mutation. GA makes the better solution by these operations and components [4],[5].

Fig. 1 shows that the structure of individual which is one of basic components in Genetic Algorithm for Document Clustering(GADC). As shown in Fig. 1 each gene present the cluster number for one document in the document data set. In population Initialization step, the cluster numbers generated randomly from 1 to k (where k is the number of clusters).

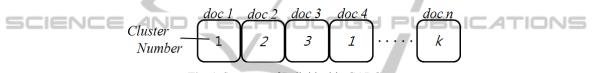


Fig. 1. Structure of Individual in GADC.

GADC consists of Population Initialization, Fitness Function and basic operations (selection, crossover, mutation). Fig. 2 shows that the GADC process. Each module in the process is explained as follows

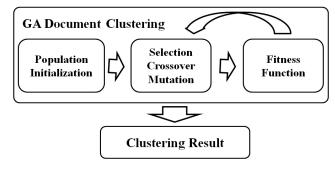


Fig. 2. GADC process.

**Population Initialization.** Suppose GADC has *P* individuals. Each individual has the information of the cluster which a document belongs to. In population initialization each document in assigned to a cluster randomly. The number of individuals *P* influences the performance and the execution time of GADC.

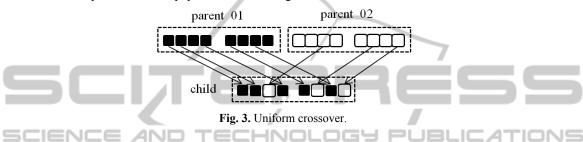
Selection Operation. Selection operation of GADC is quality proportion roulette wheel selection one. (1) shows the definition of quality proportion roulette wheel

selection. After each individual is evaluated the best individual has *t* times more survival chance than the worst individual.

$$C_{i} = (f_{w} - f_{i}) + \frac{(f_{w} - f_{b})}{t - 1}, t > 1$$
(1)

where  $C_i$  means selection probability of current individual,  $f_i$  the fitness of current individual,  $f_w$  the worst fitness of set of individuals and  $f_b$  the best fitness in individuals. In our experiment, *t* equals to 3.

**Crossover Operation.** Crossover operation uses gene information of parent populations to generate new population. Crossover operation of GADC uses uniform crossover operation in this paper as shown in Fig. 3.



In the uniform crossover scheme, genes in the chromosome are compared between two parents. There genes are swapped with a fixed probability. We use 0.5 as the fixed probability value.

**Mutation Operation.** Mutation operation produces the features which are not at the parent features. The failure mutations are select out and the successful mutations contribute to the improvement the quality of population. Mutation operation of GADC is typical variation. In the typical variation scheme the information of gene is randomly changed with a fixed probability (0.015).

**Fitness Function.** The individuals are evaluated by fitness function if they are good or not. The fitness value is calculated by in the GADC by the average similarity of all documents in a cluster [9],[10]. (2) shows that the formula of the Fitness Function.

Fitness = 
$$\frac{1}{K} \sum_{i=0}^{k} \text{AveSim}_i$$
, where  $\text{AveSim}_i = \sum_{j=0}^{NC_i-1} \sum_{k=j+1}^{NC_i} \text{Sim}(d_{ij}, d_{ik})$  (2)

where k is the number of clusters and NC<sub>i</sub> the number of documents in *i*th cluster.  $d_{ij}$  is *j*th document in *i*th cluster. The similarity function  $Sim(d_{ij}, d_{ik})$  is cosine value between  $d_{ij}$  and  $d_{ik}$ . In this paper we use cosine similarity for  $Sim(d_{ij}, d_{ik})$  [9],[10].

# **3** Document Clustering based on Genetic Algorithm using D-Individual

Sometimes genetic algorithm makes the result not enough to solve the problem. That is cause of low performance and called premature convergence phenomenon [7]. In

this paper proposed the genetic algorithm using D-Individual (DIGA) to solve the phenomenon. D-Individual is generated from GADC. (It is first introduced in this paper). Fig. 4 shows the flowchart of DIGA. The process of DIGA is almost like that of GADC. The only difference is using the D-individual in the Population Initialization. The other operations of DIGA, selection, crossover and mutation are the same in the operations of GADC.

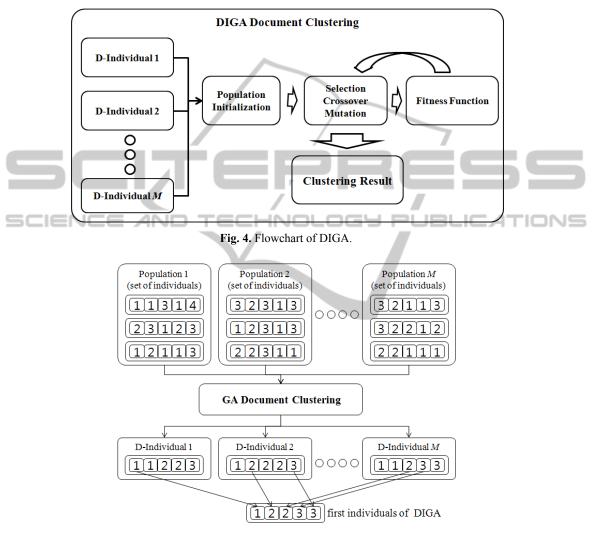


Fig. 5. Generating process of D-individual in DIGA algorithm.

**Population Initialization of DIGA.** DIGA generates the first individuals by using the D-Individuals. D-Individuals have the feature information of document clusters. Using D-Individuals we can make individual which has a higher possibility of evolution. Fig. 5 shows the generating process of D-Individual in DIGA. DIGA decides M(number of D-Individuals) and P(number of individuals). M and P influence DIGA

performance and execution time. DIGA can have better performance in short time compare with traditional GA depends on the *M* and *P*. An experiment and result analysis about this are covered in more detail in Chapter 4.

#### 4 Implementation and Results

This paper proposes the document clustering based on the genetic algorithm using D-Individual. For estimating its performance, the Reuter-21578 text collection set is used. Three Topic-Sets are experimented and four subjects were allocated to each Topic-Set. Each subject has 50 documents, so that a Topic-Set has 200 documents. Fig. 6 shows the Topic-Sets and subjects.

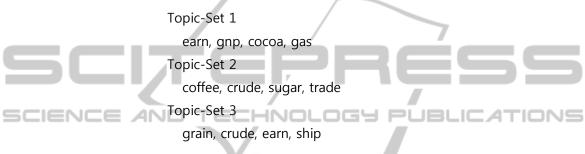


Fig. 6. Topic-Sets and Subjects.

As shown Fig. 6, 4 subjects in a Topic-Set are less related so that each subject clearly distinguished each other. To evaluate the clustering performances, F1-measure defined as in (3) is used

$$F1 - Measure = \frac{2 \times Pricision \times Reccall}{Pricision + Recall}$$
(3)

The performances and execution time of GADC depends on the number of individuals as well as those of DIGA are depends on the number of individuals and the number of D-Individuals. Fig. 7 and 8 shows the difference of the GADC and DIGA performances and execution times depending upon P(number of individuals) and M(number of D-Individuals).

From result of Fig. 7 we can see that the DIGA shows a better performance than GADC when P is the same. But these results are not enough because DIGA has longer execution time than the GADC as shown in Fig. 8. So we need more comparison and analysis.

Fig. 9 shows the comparison of performance of DIGA and GADC. From Fig. 9 we can see DIGA has better performance than GADC in same execution time.

The K-means clustering algorithm, the Group Average clustering algorithm and GADC are compared to evaluate the proposed clustering algorithm, DIGA. Fig. 10 demonstrates the performance of clustering algorithms on each Topic-Set. It clearly shows that DIGA has better performance than others. GADC shows better performance than both K-means and Group Average clustering algorithms.

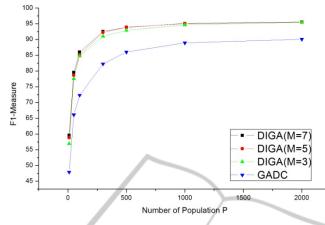


Fig. 7. GADC and DIGA performance depending upon P and M.

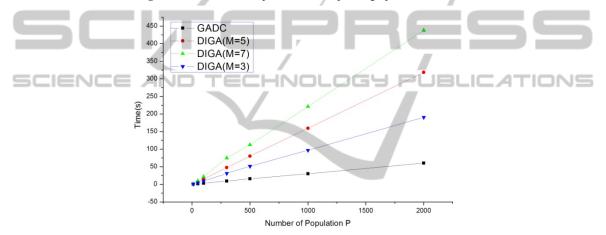


Fig. 8. GADC and DIGA execution time depending upon P and M.

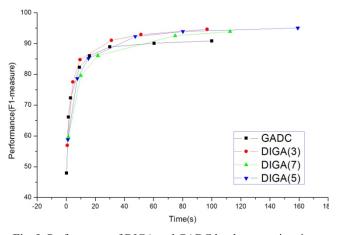
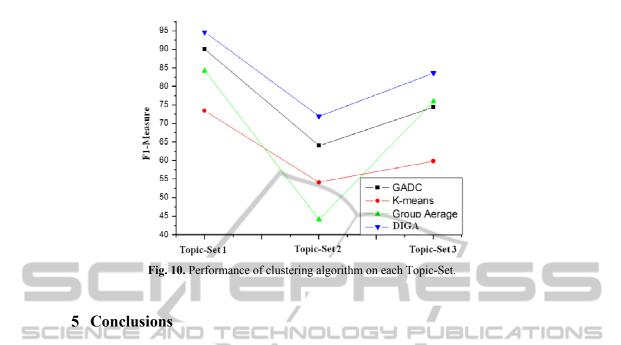


Fig. 9. Performance of DIGA and GADC by the execution time.



In this paper we have proposed the document clustering based on the genetic algorithm using D-Individual. Our DIGA demonstrates the best performance among the currently used clustering algorithms such as K-means clustering and Group Average clustering. Moreover, DIGA has the higher performance and shorter execution time than the general GADC. In this research, we find some meanings about D-individual which have the feature information of document clusters, that helps traditional genetic algorithm become more fast and accurate. DIGA has weakness of taking a long execution time due to the complexity still. Regardless of execution time, our experiments convince that DIGA shows the most accurate result in document clustering.

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