Advances in Intelligent Systems and Computing 553

Sanjiv K. Bhatia Krishn K. Mishra Shailesh Tiwari Vivek Kumar Singh *Editors* 

# Advances in Computer and Computational Sciences

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## Preface

The ICCCCS is a major multidisciplinary conference organized with the objective of bringing together researchers, developers and practitioners from academia and industry working in all areas of computer and computational sciences. It is organized specifically to help computer industry to derive the advances of next generation computer and communication technology. The invited researchers present the latest developments and technical solutions.

Technological developments all over the world are dependent upon globalization of various research activities. Exchange of information and innovative ideas are necessary to accelerate the development of technology. Keeping this ideology in preference, Aryabhatta College of Engineering & Research Center, Ajmer, India has come up with an event—International Conference on Computer, Communication and Computational Sciences (ICCCCS-2016) during August 12–13, 2016.

Ajmer is situated in the heart of India; just over 130 km southwest of Jaipur, a burgeoning town on the shore of the Ana Sagar Lake, flanked by barren hills. Ajmer has historical strategic importance and was ransacked by Mohammed Gauri on one of his periodic forays from Afghanistan. Later, it became a favorite residence of the mighty Mughals. The city was handed over to the British in 1818, becoming one of the few places in Rajasthan controlled directly by the British rather than being part of a princely state. The British chose Ajmer as the site for Mayo College, a prestigious school opened in 1875 exclusively for the Indian Princes, but today open to all those who can afford the fees. Ajmer is a perfect place that can be symbolized for demonstration of Indian culture, ethics and display of perfect blend of wide plethora of diverse religion, community, culture, linguistics, etc., all coexisting and flourishing in peace and harmony. This city is known for the famous Dargah Sharif, Pushkar Lake, Brahma Temple and many more evidences of history.

This is the First time Aryabhatta College of Engineering & Research Center, Ajmer, India is organizing International Conference on Computer, Communication and Computational Sciences (ICCCCS 2016), with a foreseen objective of enhancing the research activities at a large scale. Technical Program Committee and Advisory Board of ICCCCS include eminent academicians, researchers and practitioners from abroad as well as from all over the nation.

In this volume, selected manuscripts have been subdivided into various tracks named 'Intelligent Hardware and Software Design', 'Advanced Communications', 'Power and Energy Optimization', 'Intelligent Image Processing', Advanced Software Engineering', 'IoT', 'ADBMS & Security', 'Evolutionary and Soft Computing'. A sincere effort has been made to make it an immense source of knowledge for all and includes 147 manuscripts. The selected manuscripts have gone through a rigorous review process and are revised by authors after incorporating the suggestions of the reviewers.

ICCCCS 2016 received 429 submissions from around 729 authors of 12 different countries such as USA, Iceland, China, Saudi Arabia, South Africa, Taiwan, Malaysia and many more. Each submission has been checked with anti-plagiarism software. On the basis of plagiarism report, each submission was rigorously reviewed by at least two reviewers with an average of 2.45 reviewers per review. Even some submissions have more than two reviews. On the basis of these reviews, 140 high-quality papers were selected for publication in this proceedings volume, with an acceptance rate of 32.6%.

We are thankful to the speakers, delegates and the authors for their participation and their interest in ICCCCS as a platform to share their ideas and innovation. We are also thankful to the Prof. Dr. Janusz Kacprzyk, Series Editor, AISC, Springer and Mr. Aninda Bose, Senior Editor, Hard Sciences, Springer for providing continuous guidance and support. Also, we extend our heartfelt gratitude to the reviewers and Technical Program Committee Members for showing their concern and efforts in the review process. We are indeed thankful to everyone directly or indirectly associated with the conference organizing team for leading it towards the success.

Although utmost care has been taken in compilation and editing, however, a few errors may still occur. We request the participants to bear with such errors and lapses (if any). We wish you all the best.

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## Part I Intelligent Hardware and Software Design

## **Biogeography-Based Optimization** for Cluster Analysis

Xueyan Wu, Hainan Wang, Zhimin Chen, Zhihai Lu, Preetha Phillips, Shuihua Wang and Yudong Zhang

**Abstract** With the aim of resolving the issue of cluster analysis more precisely and validly, a new approach was proposed based on biogeography-based optimization (abbreviated as BBO) algorithm. (Method) First, we reformulated the problem with an optimization model based on the variance ratio criterion (VARAC). Then, BBO was presented to search the optimal solution of the VARAC. There are 400 data of four groups in the experimental dataset, which have the degrees of overlapping of three distinct scales. The first one is nonoverlapping, the second one is partial overlapping, and the last is severely overlapping. BBO algorithm was compared with three different state-of-the-art approaches. We ran every algorithm 20 times. In this experiment, our results demonstrate the maximum VARAC values that can be found by BBO. The conclusion is that BBO is predominant which is extremely quick for the issue of clustering analysis.

**Keywords** Biogeography-based optimization • Genetic algorithm • Cluster analysis

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#### 1 Introduction

In the context of a group, cluster analysis is defined as a case of substance with the mode that a lot of targets in a large cluster which are closer to another side than others in the additional clusters [1]. Cluster analysis is a way of unsupervised studying, and in lots of areas, it is also used for statistical data analysis in a common technique, consisting of breeding value [2], food quality monitoring [3], gene engineering [4], pediatric immunization distress [5], chronic rhinosinusitis [6], community analysis [7], etc.

Currently, various algorithms were proposed for cluster analysis. They can be basically classified into the following four categories: centroid-based clustering, distribution-based clustering, density-based clustering, and connectivity-based clustering.

In the research, the most attractive to us is centroid-based methods. In this type, there are two representative algorithms, one is fuzzy *c*-means clustering (FCM) [8], the other is *k*-means clustering [9]. These are iterative methods and affected by a lot of factors, for example, if the initial partition is not determined properly, they may stop at local best instead of global optimal solution.

To solve above the problems, Cotta and Moscato [10] proposed that the best way to detect the global optimum clustering is branch and bound algorithm. But it requires too much calculating time. In the past few years, to solve the problem of clusters, evolutionary algorithm has been proposed, because it is insensitive to initial values. Jarboui et al. [11] put forward an original clustering approach on a foundation of the combinatorial particle swarm optimization (CPSO) algorithm. Gelbard et al. [12] proposed cross-cultural research to cluster analysis with the methodology named Multi-Algorithm Voting (MAV). Niknam and Amiri [13]

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thought the k-means algorithm is extremely dependent on its initial state and integration of the local best solution. Hence, they put forward to a fresh hybrid evolutionary algorithm. Huo [14] raised chaotic artificial bee colony (CABC) as one method for the issue of cluster division. Abul Hasan and Ramakrishnan [15] underwent a survey on cluster analysis about hybrid evolutionary algorithms. They believed that optimization algorithms are required for cluster analysis to obtain better results. Zhang and Wu [16] proposed bacterial foraging optimization (BFO) for cluster analysis. Kuo et al. [17] put forward an original approach named dynamic clustering on the basis of PSO and GA (DCPG) algorithm. Firefly algorithm (FFA) was used by Zhang and Li [18]. They took 400 data and divided them into four groups for testing. Yang et al. [19] proposed and explored the idea of exemplar-based clustering analysis optimized by genetic algorithms (GA). Wan [20] made a full use of the assembling analysis and researched k-means with particle swarm optimization (KPSO). Palaparthi et al. [21] researched a multi-objective optimization (MOO) channel in relevance with cluster analysis to research the vocal folds' form-function connection. On the basis of the algorithms of improved genetic, Cao and Mu [22] put forward to a weighted k-means clustering algorithm. Ozturk et al. [23] proposed a modified binary artificial bee colony algorithm. Zhou and Zhang [24] proposed a very advanced algorithm called quantum-behaved particle swarm optimization (OPSO).

Nevertheless, those above approaches are vulnerable to several disadvantages: (1) They may converge to local optimal solutions, corresponding to suboptimal results. (2) They may cost too lengthy time and cost large memory. Hence, a new algorithm, viz., biogeography-based optimization (BBO) [25] was suggested in settling the issue with this research.

The remaining portion of this paper is as follows: In Sect. 2, we define the mathematical model of partitional clustering, as well as propose the clustering criterion and encoding strategy. In Sect. 3, we describe the novel BBO method. In Sect. 4, we describe the experiment, this section includes three types of artificial data which have different overlapping degrees. In Sect. 5, we discuss about the experiment results. Finally, in Sect. 6, we conclude this paper.

#### 2 Model Definition

**Hypothesis**: there are *n* samples  $Q = (q_1, q_2, ..., q_n)$  in a *d*-dimensional metric space. Every sample belongs to one of the *m* sets. Every  $q_i \in R^d$  indicates a particular feature vector that is composed of *d* dimension. The clusters are denoted as  $F = \{f_1, f_2, ..., f_k\}$ , where *k* represents the number of clusters. They should submit three declarations as follows:

$$f_i \neq \phi \quad \text{for} i = 1, 2, \dots, m$$
  

$$f_i \cap f_j = \phi \quad (i \neq j)$$
  

$$\cup f_i = \{q_1, q_2, q_3, \dots, q_n\}$$
(1)

The aim of this study is to look for the most appropriate partition  $F^*$  that corresponds to the best sufficiency among all possible solutions [26].

The search space is set to be *n*-dimension, which is consistent with *n*-objects. Every dimension not only expresses a sample but also expresses the *i*-th individual  $Y_i = \{y_{i1}, y_{i2}, \ldots, y_{in}\}$  that conforms with the affection of *n* samples, such as  $y_{ij} \in \{1, 2, \ldots, m\}$ , in which *j* indicates the *j*-th sample [27].

Suppose n = 6 and k = 2. Suppose the first cluster A contains the first, second, and third object, and the second cluster B contains the fourth, fifth, and sixth object. Figure 1 illustrates the encoded representation of the cluster solution.

Some criteria have been put forward to estimate the sufficient where a predefined dataset could be assembled. Among the partitional clustering strategies, "VAriance RAtio Criterion (VARAC) [28]" is one of the most popular used, which is defined as

$$VARAC = \frac{E}{R} \times \frac{n-d}{d-1},$$
(2)

where R and E, respectively, represents the transformation between intra-cluster and inter-cluster. Their definitions are as follows:

$$R = \sum_{j=1}^{d} \sum_{i=1}^{n_j} \left( q_j^i - \bar{q}_j \right)^T \left( q_j^i - \bar{q}_j \right)$$
(3)

$$E = \sum_{j=1}^{d} n_j (\bar{q}_j - \bar{q})^T (\bar{q}_j - \bar{q}), \qquad (4)$$

where (.)<sup>T</sup> represents the transpose operation,  $n_j$  stands for the cardinal of the cluster  $f_j$ ,  $q_j^i$  represents the *i*-th target allotted to  $f_j$ , which is one of the clusters,  $\bar{q}_j$  represents the center of the *j*-th cluster, and  $\bar{q}$  represents the center of the whole data, (d - 1) denotes the between-cluster variation freedom degree, while (n - d) represents the within-cluster variation freedom degree.

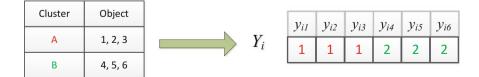


Fig. 1 Encoded representation

Generally speaking, it is expected that tight and divided clusters have important value of small R and large E. Therefore, if data partition result is much better, the value of VARAC is much larger. In order to avoid the rate to increase monotonically as the quantity of the clusters, the term (n - d)/(d - 1) is normalized, which make VARAC by means of an optimization of the clusters, it can achieve maximize criterion.

#### **3** Biogeography-Based Optimization

We gave an analogy in Fig. 2 with the well-known GA as a counterpart for making it easy to understand. As is shown in Fig. 2, the population of candidate solutions is represented as habitats in BBO [29–31]. The solution vector components are considered to be suitability index variables (SIVs). Those good solutions are considered habitat with high habitat suitability index (HSI), and vice versa [32–34]. BBO has been proven to give better performance than genetic algorithm (shorted as GA) [35], artificial bee colony (shorted as ABC) [36], firefly algorithm [37], bacterial chemotaxis optimization [38], particle swarm optimization (shorted as PSO) [39], and ant colony optimization [40]. Table 1 demonstrates the pseudocodes of BBO.

#### 4 Experiments

The experiments were performed on a computer with i7 processor and 16 GB memory, under Windows 7 operating system.

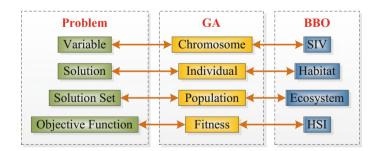


Fig. 2 The connection of optimization problem, GA, and BBO