Research paper

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Threshold Based K-Mean Adaptive Clustering for High Dimensional Data Analysis

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Abstract.

Due to rapid growth of data sources exponential have evolved the necessity in developing new techniques for gathering useful information. To gain and gather knowledgeable information clustering is an investigable technique which is used in finding hidden homogeneous patterns from the data sources. Partition based clustering techniques are employed to acquire clustering information from user specified parameters such as similarity threshold value and number of clusters. For effective clustering, we suggest a deterministic algorithm called ad Threshold based k-means Adaptive Clustering (TAC). Artificial and real data sets have been tested by it. The algorithm k-means has also been contrasted. A parameter, neighborhood distance is used to cluster data items in this suggested TAC algorithm. The user doesn't specify neighborhood distance, but it is calculated automatically, and it is also an adaptive parameter. The minimum support value of the tiny clusters is another parameter in TAC algorithm. TAC's performance is also evaluated by using real and artificial datasets, which have found outliers can be detected and overlapped and non-overlapped clusters are generated. The results show that TAC algorithm produces clusters of distinct dimensions while k-means generates clusters of almost identical dimensions.

Keywords: Adaptive clustering, Hidden and homogeneous patterns, Neighborhood distance, and Similarity index.

1. Introduction

Clustering is the method by which unlabelled data objects segments are found based only on information found in the data that describes the objects and their relationships. In all application fields k-means algorithm is commonly recognized. But it is nondeterministic

Research paper © 2012 IJFANS. All Rights Reserved, UGC CARE Listed (Group -1) Journal Volume 8, Issue 3, 2019 because it needs the initial centroids supplied by the user, it is suspectable to outliers and produces overlapped clusters. However, the use of this algorithm was not limited by these aspects of k-means rather data scientists encouraged to improve its functionality. The kmeans algorithm depends on initial centroids or the data objects being randomly selected. This means that the findings of these algorithms are different in sequence of executions and they are deemed to be non-deterministic algorithm. So, in order to make k-means as deterministic algorithm an effective Threshold based Adaptive Clustering (TAC) has been suggested in this proposed work. TAC produces both overlapped and non-overlapped clusters. It uses an adaptive threshold value for similarity and detects outliers.

2. PROPOSED TAC ALGORITHM

Provide sufficient detail methods to allow the work to be reproduced. Methods already published should be indicated by a reference: only relevant modifications should be described.

A. Parameters used in TAC algorithm

Clustering parameters play an important role. The clustering algorithms functionality depends on them, so they should be carefully specified by the user. Two parameters were introduced in the TAC algorithm: neighborhood distance threshold and minimum support.

Neighborhood distance threshold: This is the first parameter used in the proposed TAC algorithm and is used to identify all the neighbours of a data object. Partitioning based algorithms are also employ this parameter but as non-adaptive and user specified. In this proposed TAC algorithm, it is an adaptive parameter, the value this parameter increases during the formation of a cluster, as well as user doesn't specify it. TAC algorithm needs neighborhood distance value (Nd) in order to form a cluster.

Mathematically Nd is defined as

Nd = max (minp, minq) (1)

Where minp = min $[|p-r| : r \in set of un-clustered data objects and Minq = min <math>[|q-r| : r \in set of un-clustered data objects.$ Here P and Q are the farthest data objects in the un clustered data objects. Nd is not fixed value, rather it is increased by a small value (Δ) during the formation of a cluster. The small increment to be made in Nd is the distance between the closest member of a cluster. It helps in increasing the size of a cluster. It is defined as

$$\delta = \min \{ [\mathbf{u} - \mathbf{v}] : \mathbf{u}, \mathbf{v} \in \text{cluster} \}$$
(2)

Research paper © 2012 IJFANS. All Rights Reserved, UGC CARE Listed (Group -I) Journal Volume 8, Issue 3, 2019 Moreover, value of Nd is initialized every time a new cluster is being formed.

Minimum Support Parameter: The proposed TAC algorithm uses this parameter to improve the result of the clustering. This parameter specifies the minimum number of data objects in a cluster. The usage of minimum support has been adopted by the Association Rule mining technique. Its value can be specified by the user after clusters have been generated. It improves the result of clustering by removing the small clusters

B. TAC algorithm

TAC algorithm consists of the following steps:

Step 1: Initially the farthest data objects are identified from the given dataset.

Step2: Now, the distance of the closest data object to one of the farthest objects and the distance of the closets data objects to other farthest data objects are calculated. The formation of a new cluster starts from one of the farthest data objects whose closest data object is closer than its counterpart.

Step 3: Initialize the value of Nd and assign the selected farthest data object and all other data objects within the neighborhood distance as a member of cluster being formed.

Step 4: Increase the value od Nd by a small value Δ and assign more data objects as the members of cluster being formed. This step is repeated till new data objects are added to cluster.

Step 5: If more than one data objects are left un clustered then again find the farthest data objects from un clustered data objects and repeat steps 2,3, and 4.

TAC uses a function new cluster(R,Nd,K,Adj[n][n]) to generate clusters. This function generates overlapping clusters and membership of overlapping data objects by using the above-mentioned algorithm. The size of each cluster generated is compared with given minimum support value and small size clusters to be considered insignificant. This algorithm always produces the same result for given dataset and minimum support value on successive runs [15].

3. Results and Discussion

The experiments were performed on artificial and real data sets respectively in the first and second subsections. The TAC algorithm is compared to the K-means algorithm in the third subsection

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3.1. Experiment on Artificial Datasets

Five artificially generated data sets used to assess TAC algorithm efficiency. Table 1 Presents the result of TAC worked on Artificial Datasets

	objects covered 77.2% of
	77.2% of
AD_2K_1 34 1.5%, i.e. 14 249, 36, 88, 7	//.2/0 01
Minimum 30 data 47, 40, 202, D	Dataset
objects/cluster 68, 48, 270,	
61, 33, 50, 66	
AND 286	
AD_1.5K_2 29 1.5%, i.e., 14 249, 88, 40, 80	80.6 % of
minimum 23 data 48, 100,53, D	Dataset
objects/cluster 171, 64, 28,	
14679, 94, 22	
AND 28	
AD_1.5K_3 31 1.5%, i.e. 13 36, 47, 29, 75	75 % of
minimum 23 data 202, 68, 33, D	Dataset
objects/cluster 296, 31, 60,	
41, 181, 29	
AND 72	
AD_1.5K_4 33 1.5%, i.e. 15 249, 88, 40, 78	78.5% of
minimum 23 data 48, 58, D	Dataset
objects/cluster 28,162, 124,	
29, 41, 4178,	
24, 131 AND	
37	

Table 1 Results of TAC algorithm

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Research paper	Research paper © 2012 IJFANS. All Rights Reserved, UGC CARE Listed (Group -I) Journal Volume 8, Issue 3, 2019				
AD_2K_1	34	1.5%, i.e., 14	249, 36, 88, 77.2% of		
		minimum 30 data	47, 40, 202, Dataset		
		objects/cluster	68, 48, 270,		
			61,33, 50, 66		
			AND 286		

Wisconsi-Original breast cancer (Bcw-O), Ecoli, Glass Identification (Gi), Haberman's Survival, Iris, Seed, Wine and Yeast were used to evaluate TAC algorithm efficiency. These datasets are taken from the repository of UCI. For these datasets, the minimum support value is 5% of the dataset size.

4. COMPARISON OF TAC ALGORITHM WITH K-MEANS ALGORITHM

To Compare Clustering Results of TAC algorithm on Artificial and Real Datasets, the portion of dataset which lies within the ambit of significant clusters is taken as an input to the k-means algorithm. The value of k required in the k-means algorithm is taken from the dataset clustered using TAC algorithm. It is equal to the number of significant clusters generated. in this section, the results of this comparison are presented. clustering results of k-means

algorithm on artificial datasets Ad_2k_1, Ad_1.5k_2, Ad_1.5k_3, Ad_1.5k_4, Ad_1.5k_5 And Ad_1.5k_6 are shown respectively. From these figures, it is evident that different partitions of a given dataset are generated. Clustering results, including comparison between k-means algorithm and tac algorithm are summarized in the following table . For these datasets K-means algorithm is executed for the same eight real datasets as used by the TAC algorithm. The clustering results obtained for these datasets are presented in table 3 and it has been observed that k-means algorithm partitions the dataset into K clusters such that the range of the number of data objects in a cluster is narrow whereas the proposed algorithm can handle a wider range. It indicates that k-means algorithm generates clusters of almost same sizes whereas tac algorithm generates clusters of different sizes.

 Table 2. COMPARISON OF TAC WITH K-MEANS

Ī	#	TOTAL	ALGORITH	
			М	

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DATASE	CLUSTERS	DATA		
Т				
	(K)	OBJECTS	# DATA	# DATA OBJECTS IN
			OBJECTS IN	CLUSTERS
	-			
			CLUSTERS	
BCW-O	4	617	179, 79, 94	40, 449, 89 AND 39
			AND	
			265	
ECOLI	2	276	204 AND 72	55 AND 221
GI	3	203	89, 23 AND	18, 174 AND 11
_			91	- 7
HABER	4	295	31, 74, 81	20, 200, 31 AND 44
MAN			AND	
			109	
IRIS	3	149	49 39 AND	31, 96 AND 22
			61	51, 7011112 22
SEED	7	174	25 18 24	26, 21, 38, 20, 25, 21,
JEED	/	1/4		20, 21, 30, 20, 23, 21,
			26,27,	AND 22
			25 AND 29	AND 23

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WINE	4	159	70, 56, 25	20, 78, 32 AND 29
			AND 8	
YEAST	5	1253	341, 214,	150, 718, 81, 143 AND
			124,	161
			208 AND	
			366	

5. Conclusions

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From the experiments carried out in the job described in this article the following observations were observed. TAC algorithm automatically clusters the datasets without knowing the number of clusters. Many clusters generated using this algorithm, however the parameter minimum support plays an important role to obtain significant clusters. The proposed algorithm can generate both overlapped and non-overlapped clusters. Since the algorithm is deterministic in its nature it gives same result on successive runs for a given dataset and given minimum support value it can detect outliers. It doesn't generate no spatial clusters as it is based on centroid based clustering. It requires adjacency matrix as the prerequisite, which increases the time complexity of the algorithm. Overlapped and non-overlapped clusters. It has an advantage over the single pass and modified single pass clustering algorithms as it does not depend on the order of the selection of the data objects. Here, it has also been observed from the experiments that tac algorithm generates clusters of different sizes whereas k-means algorithm generates clusters of almost same sizes. Therefore, the proposed TAC algorithm can be used as an alternate clustering algorithm.

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