Analyzing Subspace Clustering Approaches for High Dimensional Data

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1 Introduction

A huge body of knowledge can be exploited through clustering techniques. Clustering algorithms make use of knowledge present in input data. It uncovers the hidden patterns existing in data. Clustering is one of the effective data mining techniques for grouping similar data objects [\[1](#page-23-0)]. The grouping is made on the basis of attributes. With the tremendous growth in the technology of information sciences, huge data is being produced. Such data are sensor data, web data, bioinformatics data and many more. The large volume of data not only includes a large number of rows but also a large number of attributes. The datasets with many columns/features are termed as high dimensional data. Clustering high dimensional data poses a num-ber of computational challenges [[2\]](#page-23-1). Traditional clustering algorithms' efficacy degrades on high dimensional data. This problem is termed as "curse of dimensionality." This is because traditional clustering algorithms fnd clusters in all dimension space. However, clusters might exist in a few subsets of dimensions. All dimensions might not be important for all clusters. Considering all dimensions for clustering might hide the relevant dimensions. Another challenge is the distance measure. Distance between data objects becomes meaningless in high dimensional data [[3\]](#page-23-2). All objects appear equidistant from each other. Most of the traditional clustering algorithms use distance measure for grouping data objects. New techniques are developed to fnd relevant dimensions and overcome challenges of high dimensional clustering.

Relevant dimensions are selected through feature selection methods and, subsequently, selected features to participate in complete clustering method. However,

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clusters might exist in different subspaces. Dimensions participating in one cluster might be irrelevant for another cluster. Hence to fnd clusters in different subspaces as shown in Fig. [1](#page-1-0) (each circle represents one cluster), subspace clustering methods are employed [[4\]](#page-23-3). It is an extended version of traditional clustering algorithms. Subspace clustering algorithms confne the search in a certain manner, i.e., either top-down or bottom-up approach so that it is able to determine clusters existing in various subspaces. Further, subspace clusters can be non-overlapping or overlapping in dimensions/objects.

The signifcance of subspace clustering is intensifying from last two decades. Figure [2](#page-1-1) shows the percentage of papers published on various methods of high

Fig. 2 Papers published on different techniques of high dimensional clustering

dimensional clustering from the year 2000 onwards. The main techniques for handling high dimensional clustering $[5, 6]$ $[5, 6]$ $[5, 6]$ $[5, 6]$ are as follows:

- (a) Dimensionality reduction
- (b) Parsimonious models
- (c) Projected clustering
- (d) Soft projected clustering
- (e) Subspace clustering

It was observed that only 15% of work is done in high dimensional clustering through dimensionality reduction approach. Sixteen percent of papers are published for clustering high dimensional data using soft projected clustering and parsimonious models. Seventeen percent of the work is dedicated to projected clustering. The large amount of work for high dimensional clustering is being done by subspace clustering approach. This proves that subspace clustering methods are gaining considerable attention in current research.

This study is entailed to answer the critical research questions identifed below.

- (i) What are the major challenges faced by traditional clustering algorithms to cluster high dimensional data?
- (ii) What search techniques are being used in subspace clustering to determine subspaces?
- (iii) What are the evaluation measures for comparing subspace clustering algorithms?
- (iv) What is the current scenario of subspace clustering?
- (v) What are the research gaps in the literature and the future prospects of subspace clustering?

This chapter presents a combined review of approximately all subspace clustering algorithms belonging to different classes. Various evaluation measures required for comparing the clusters and subspaces are also presented. Additionally, statistical data of subspace clustering approaches published in different years and different repositories are provided. This chapter is targeted to researchers planning to work in subspace clustering area. It provides a roadmap of research in subspace clustering approach for high dimensional data. The chapter also presents the application areas, identifes gaps in present work, and suggests future opportunities for research in this feld.

The chapter is divided into following subsections: Sect. [2](#page-3-0) presents challenges in subspace clustering, Sect. [3](#page-3-1) gives various classifcations of subspace clustering approaches, Sect. [4](#page-6-0) presents evaluation measure for comparing subspace clustering algorithms, and Sects. [5](#page-9-0) and [6](#page-10-0) depict the literature survey and empirical assessment of subspace clustering algorithms, respectively. Applications and future prospects are illustrated in Sect. [7,](#page-20-0) and Sect. [8](#page-22-0) fnally concludes the chapter.

2 Challenges in Subspace Clustering

The major goal of clustering is not only to fnd similar groups of data points but to fnd high-quality groups within a reasonable time. In cases where clusters exist in different subsets of dimensions, it is essential for clustering algorithms to determine effective clusters along with relevant dimensions. Subspace clustering algorithms have proved to be efficient for extracting clusters from high dimensional datasets [\[7](#page-23-6)]. However, a number of challenges persist which needs to be focused before developing an effcient subspace clustering algorithm. These challenges are as follows:

- (a) It is hard to determine the subsets of dimensions where data points are similar. This is because a number of dimensions are large and possible combinations are huge.
- (b) It is quite diffcult to determine the distribution of data within subspaces. If data is near the cluster center and far from another subspace center, then clustering is easy; otherwise it is difficult to cluster within subspaces.
- (c) There may be overlapping subspaces that mean few dimensions may be common in few subspaces. Clustering becomes even more complicated in case of overlapping subspaces.
- (d) There are possibilities of noisy data in the dataset. That means some data points might not belong to any subspace or if they are part of any subspace, they are not part of a cluster of particular subspace. Handling of such data becomes a challenge for subspace clustering.
- (e) It is diffcult to understand which clustering algorithm and subspace strategy are appropriate for a given problem. As the number of subspaces and the dimension of each subspace is unknown, the problem becomes intricate.

Due to the above challenges encountered in subspace clustering approaches, there has been a scope of improvement in these algorithms. Subspace clustering not only determines the clusters in the dataset but also the subspaces in which these clusters are present. The next section presents the review of subspace clustering classifcation.

3 Classifcation of Subspace Clustering Approaches

In order to determine a group of similar data points in different subspaces, subspace clustering algorithms are employed. This section discusses the various categories of subspace clustering approaches on the basis of different parameters. Figure [3](#page-4-0) depicts the categories of subspace clustering algorithms on the basis of parameters required in algorithms for forming clusters from data objects [[8\]](#page-24-0). These are as follows:

Cell-based Subspace Clustering: This approach is also called a grid-based approach. It makes use of an approximate number of cells required to form a cluster.

The cluster description starts with a minimum width "w" of a number of cells. Each cell contains a minimum threshold number of objects. The cells of a cluster are either of fxed grid size or variable in number forming hypercube of width "w." The cells participating in clustering uses subsets of dimensions of datasets. Hence the relevant dimensions to a particular cluster are determined. Irrelevant dimensions, not participating in clustering, expand on other cells. Few cell-based algorithms are CLIQUE, DOC, MINECLUS, SCHISM, etc.

Density-based Subspace Clustering: This approach is able to determine the clusters of arbitrary shapes. It is dependent upon the density of data objects lying in datasets. This approach separates the dense region from sparse region. Density is determined through distance measure. The parameters used in algorithms are the least number of points "minpts" required to form a cluster and "epsilon" distance among the neighboring points. The dense region is formed by counting the number of points "minpts" within "epsilon" neighboring distance. Any region not satisfying the "minpts" and "epsilon" properties is not able to form a cluster and is termed as sparse region. Some density-based subspace clustering algorithms are FIRES, INCY, SUBCLU, etc.

Clustering-Oriented-Based Subspace Clustering: This approach does not provide any requirements for cluster formation. It is not dependent on any cluster defnition or on input parameters to form a cluster. As the name suggests it gives statistical orientation properties of total clusters formed. It means it defnes properties of resultant clusters formed like a number of clusters formed, average dimensionality per cluster, etc. Clustering oriented approach is more suitable to datasets of varied distributions. Some algorithms of this approach are STATPC, P3C, PROCLUS, etc.

Subspace clustering approaches confne their search in such a manner that clusters existing in different subspaces are extracted [\[3](#page-23-2)]. The search either proceeds from single-dimensional to full dimensional dataset (bottom-up) or full dimensional to single dimensional dataset (top-down). Each search method is defned as follows:

Bottom-Up Subspace Search Method: It is a grid-based method which starts from single dimension. This method follows an a priori approach [[1\]](#page-23-0) to determine relevant dimensions. The method begins with forming similar groups in single dimensions on the basis of density threshold parameters and grid size. The data objects participating in single dimension will also participate in multi-dimensional

grids. This approach detects the noise and also determines overlapping subspace clusters. However, it may fnd redundant subspaces or clusters across the dataset. Some algorithms of bottom-up approach are CLIQUE, ENCLUS, DOC, etc. (Fig. [4\)](#page-5-0).

Top-Down Subspace Search Method: It is an iterative method which starts from entire dimensions of dataset. Initially, each dimension is assigned equal weights and clustering begins. After clustering, the weights of each dimension for each cluster are updated. In the next iteration, updated weights are used, clustering proceeds, and again weights are updated. Dimensions with highest weights for a cluster are relevant dimensions. This is expensive method which requires many iterations. Input parameters used in this approach are number of clusters and size of subspaces. Both the parameters are diffcult to decide before clustering. This method fnds nonoverlapping clusters. Some algorithms are PROCLUS, FINDIT, MAFIA, etc.

Data objects of clusters determined from subspace clustering approaches may or may not be aligned along the axis [\[9](#page-24-1), [10\]](#page-24-2). On this basis, subspace clustering approaches are divided into two categories as shown in Fig. [5](#page-5-1).

Axis Aligned: Clusters determined along the parallel axis to data space are axisaligned. These subspace clusters could be determined with low computational complexity. Number of subspaces determined from this approach are fxed in number, e.g., CLIQUE and DOC.

Non-axis Aligned: the subspace clusters determined in the arbitrary orientation of data space are non-axis aligned. Clusters may be expressed in a better way using

Fig. 4 Classifcation of subspace clustering approaches on basis of search methods

Fig. 5 Classifcation of subspace clustering approaches on basis of axis alignment

this approach. However, computation complexity is quite high in fnding clusters in arbitrary orientation. Subspaces may go infnite in number, e.g., Orclus.

4 Evaluation Measures for Subspace Clustering Algorithms

This section describes the various systematic evaluation measures used for comparing objects and subspaces of clusters formed. However, there are no standard criteria defned for comparing the subspaces or clusters formed from subspace clustering approach. In literature, researchers have employed different evaluation measures for performance assessment of subspace clustering algorithms. A common ground of comparing subspace clustering algorithms is lacking. This is because true cluster labels along with relevant dimensions are lacking in datasets. The paper presents thorough evaluation measures shown in Fig. [6,](#page-6-1) required for comparing subspace clustering algorithms [\[8](#page-24-0), [9](#page-24-1)].

Object-Based Validation Measures: This type of validation considers the data objects participating in the clustering process. The various measures are as follows:

F1 Measure – It is the harmonic mean of precision and recall values. This measure ensures that actual cluster (found cluster from the algorithm) should mask maximum objects of true cluster (already given class in dataset) and unmask the objects of different clusters. This is computed from the confusion matrix [[1\]](#page-23-0). Let TP is true positive, i.e., objects of actual cluster are same of true cluster; TN is true negative, i.e., objects of different true clusters are mapped to different actual

Fig. 6 Evaluation measures for subspace clustering approaches

clusters; FN false negative, i.e., objects of same true cluster belong to different actual cluster; FP false positive, i.e., objects of different true cluster belong to same actual cluster. Precision is minimum mapping of objects from other clusters, while recall is maximum mapping of objects from same true cluster. It is given by Eqs. [1](#page-7-0), [2,](#page-7-1) and [3:](#page-7-2)

$$
Precision = \frac{TP}{TP + FP}
$$
 (1)

$$
Recall = \frac{TP}{TP + FN}
$$
 (2)

$$
F1_Measure = \frac{2 * Precision * Recall}{Precision + Recall}
$$
 (3)

Accuracy: Accuracy is the ratio of a number of objects of the actual cluster correctly mapped with objects of true cluster by total objects. It can be calculated using confusion matrix through following Eq. [4](#page-7-3):

$$
Accuracy = \frac{TP + TN}{TP + TN + FP + FN}
$$
 (4)

Purity: This measures the purity or homogeneity of actual clusters determined from clustering algorithm with respect to true clusters. It can be calculated as follows:

$$
Purity = \frac{TP}{TP + TN + FP + FN}
$$
 (5)

Object and Subspace-Based Validation Measures: This type of validation considers not only the data objects but the relevant subsets of dimensions participating in the clustering process. These measures actually evaluate how well clusters are formed in various subspaces. In these measures, an object of a dataset is assumed to be divided into number of sub-objects which spans across all the dimensions of datasets. Hence a subspace cluster consists of relevant dimensions along with subobjects. An object shared between the subspace clusters with disjoint dimensions will have different sub-objects. Thus, subspace clusters having same objects with different relevant dimensions do not overlap. The various measures are as follows:

Relative Non-Intersecting Area (RNIA): This measure ensures the maximum number of sub-objects found in actual subspace cluster maps the true subspace cluster. A confusion matrix is formed from actual subspace clusters versus true subspace clusters [[9\]](#page-24-1). The matching sub-objects with each subspace clusters are flled in the matrix. Let "U" is the total number of sub-objects participating in true or actual subspace clusters. Same sub-objects participating found in both true and actual cluster (with the same dimension) is counted once. Let "I" is the intersecting

subjects of true and actual subspace clusters, or it is the sum of all elements of confusion matrix. Then RNIA is calculated as follows:

$$
RNIA = \frac{U - I}{U}
$$
 (6)

Clustering Error (CE): RNIA gives the same result in a case when several actual clusters match one true cluster or one actual cluster matches true cluster. Clustering error maps one actual cluster to almost one true cluster and vice versa. CE is also calculated from confusion matrix formed by above method. Let D is the sum of all principal diagonal elements, then CE is given as:

$$
CE = \frac{U - D}{U} \tag{7}
$$

Rand Index: This index is measured on the basis of counting the pair of subobjects that do or do not participate in clustering. All sub-objects that do not overlap are considered as single clusters. A confusion matrix is created considering all subspace clusters including singleton clusters. There are four labels important, i.e., N11, pair of sub-objects that are same in both actual and true subspace clusters; N10, pair of sub-objects that are same in a true cluster but not in actual cluster; N01, pair of sub-objects that are same in actual cluster but not in true cluster; and N00, pair of sub-objects that are different in both actual and true subspace clusters.

$$
1 - \text{Rand}_{\text{index}} = \frac{\text{N10} + \text{N01}}{\text{N}} \tag{8}
$$

where $N = \frac{U*(U-1)}{2}$, U is the union of all sub-objects participating in clustering.

Scalability: Scalability is the measure which could be used to visualize and analyze the behavior of any algorithm. In subspace clustering algorithms, scalability is measured in terms of dimensionality. It depicts the performance of an algorithm with the increase in a number of dimensions of dataset. The graphs are plotted which could be shown for any evaluation measure with respect to dimensions. The X axis shows the dimensions and Y axis depicts evaluation metric like F1_Measure or accuracy etc. Scalability of subspace clustering algorithms is shown in the latter part of the chapter.

Time Complexity: Run time complexity measures the time taken by an algorithm to cluster a given dataset. It is represented in the form of graphs to compare subspace clustering algorithm with respect to dimensions. Few examples of representation are shown in [[8\]](#page-24-0).

Ranking: Subspace clustering algorithms can be ranked on the basis of average ranking (AR) and success rate ratio ranking (SRR) [\[11](#page-24-3)]. The average rank of an algorithm is computed by taking the mean of ranks on all dataset on basis of any evaluation measure. Let r_j^i be the j^{th} algorithm rank for i^{th} dataset. The average rank of each algorithm on total "*n*" datasets is computed using following Eq. [9:](#page-9-1)

$$
r_j = \frac{\sum_{i=1}^n r_j^i}{n} \tag{9}
$$

Success rate ratio rank (SRR) computes the ratio of success rates in a pair of algorithms. This method is useful in determining the signifcant differences in algorithms. In SRR ranking, an algorithm and dataset are taken and accuracy (any evaluation measure) ratio is calculated with respect to other algorithms. This ratio is computed by the following Equation:

$$
\text{SRR}_{j,k,j\neq k}^{i} = \frac{\text{acc}_{j}^{i}}{\text{acc}_{k}^{i}}
$$
 (10)

where "*i*" is the dataset, "*j*" is the algorithm for which SR is calculated, and "*k*" is the compared algorithm different from *j*. Hence SRR is calculated for algorithm "*j*" with reference to algorithm "*k*" on an *i*th dataset. Likewise, with the same pair of algorithms, SRR is calculated on all datasets. Subsequently, the mean of SRR is computed for all *n* dataset using Eq. [11.](#page-9-2)

$$
SRR_{j,k,j\neq k} = \frac{\sum_{i=1}^{n} SRR_{j,k,j\neq k}^{i}}{n}
$$
 (11)

In the same way, the algorithm "*j*" is paired with all other algorithms on all datasets and overall SRR for algorithm "*j*" is given by:

$$
\text{SRR}_{j} = \frac{\sum_{k} \text{SRR}_{j,k,j \neq k}}{m-1} \tag{12}
$$

where "*m*" is a number of subspace algorithms used for comparison. Similarly, SRR for all algorithms against each and every algorithm is calculated and ranked.

The further sections discuss the literature survey outline with analysis of subspace clustering algorithms on a few evaluation metrics.

5 Literature

There is a number of good surveys made by researchers on high dimensional clustering but very few are on subspace clustering. Assent [\[2](#page-23-1)] made a brief survey of high dimensional clustering on different types of datasets. Authors have also shown different methods adopted for clustering high dimensional datasets. Clustering in high dimensions is proposed through various models by [\[6](#page-23-5)]. A survey on models of high dimensional clustering datasets is given by [\[12](#page-24-4)]. Kriegel et al. [\[5](#page-23-4)] presented a survey on clustering high dimension data through subspace, correlation, and patternbased clustering methods. Steinbach et al. illustrated the challenges of high

dimensional clustering in detail and proposed a concept-based model for handling dataset with large attributes [[13\]](#page-24-5). Fahad et al. made a survey describing various classes of clustering algorithms along with different evaluation metrics [\[14](#page-24-6)]. A comparative analysis of different swarm intelligence-based clustering algorithm is depicted in [[15\]](#page-24-7). A new clustering algorithm with modifed fower pollination algorithm is shown in [[16\]](#page-24-8). However, the various studies discussed above mainly focused on clustering high dimensional datasets; limited studies have been performed on subspace clustering approaches. Parson [\[3](#page-23-2)] made a comprehensive survey on subspace clustering approaches portraying different subspace search methods. Müller [\[8](#page-24-0)] depicted the different evaluation measures as well as different categories of subspace clustering algorithms. The work [\[17](#page-24-9)] also provided a WEKA platform for evaluating various subspace clustering algorithms. Subspace clustering through evolutionary technique is proposed by [\[18](#page-24-10)]. Table [1](#page-11-0) presents the review on various subspace clustering algorithms along with evaluation metrics used, type of dataset applied, and maximum dimensions evaluated and drawbacks.

It can be observed from Table [1](#page-11-0) that maximum dimension evaluated in high dimensional dataset is 5920, but 6144 is also there. Many algorithms have been proposed on subspace clustering to handle high dimensional datasets. A Monte Carlo-based subspace clustering algorithm [[44\]](#page-25-0) is proposed by Olson et al. and evaluated against subspace and projected clustering algorithms on real and synthetic datasets. The signifcance of subspace clustering approaches is increasing in this big data era. Figure [7](#page-16-0) depicts the number of papers published in a span of 5 years (shown on the x-axis).

Subspace clustering approaches also gained importance in various repositories of publications. Figure [8](#page-16-1) shows the percentage of subspace clustering papers deposited in different repositories.

Figure [9](#page-17-0) gives the percentage of papers published in conference or journals. 53% of subspace clustering techniques was published in conferences and 47% of papers were published in journals.

The next section illustrates the empirical comparison of subspace clustering algorithms on real and artifcial datasets.

6 Empirical Assessment

Subspace clustering algorithms are mainly classifed as cell, density, and clusteringoriented-based algorithms [[8\]](#page-24-0). The algorithms belonging to these categories cover almost all subspace clustering algorithms. Present section shows the empirical assessment of subspace algorithms on the basis of two evaluation measure i.e. F1_ Measure and accuracy. The algorithms are compared on the basis of ranking and scalability.

(i) Ranking of Subspace Clustering Algorithms: Rank of algorithms [\[11](#page-24-3)] on real and synthetic datasets are made independently on accuracy and F1_Measure.

Table 1 Survey of various subspace clustering algorithms **Table 1** Survey of various subspace clustering algorithms

(continued)

Table 1 (continued)

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Table 1 (continued) **Table 1** (continued)

Fig. 7 Year-wise number of papers published on subspace clustering

Fig. 8 Subspace clustering in different repositories

Real and synthetic datasets are obtained from [\[17](#page-24-9)]. Actual values of accuracy and F1_Measure of various subspace clustering algorithms on real datasets are extracted from [[8\]](#page-24-0). While subscale algorithm is implemented in MATLAB R2013a platform adopting same parameter values given in [[27\]](#page-24-19). Tables [2](#page-17-1) and [3](#page-17-2) shows the average rank and SRR rank of subspace clustering algorithms on real and synthetic datasets respectively. CLIQUE emerged to be on the frst rank on accuracy, while MINECLUS best performs on F1_Measure.

Tables [4](#page-18-0) and [5](#page-18-1) present average and SRR ranks of subspace clustering algorithms on synthetic datasets in terms of F1_Measure and accuracy, respectively. Actual values of subspace algorithms on accuracy and F1_Measure on synthetic datasets were not available. Hence each algorithm is implemented on WEKA toolbox of subspace clustering provided by [\[17](#page-24-9)] with best parameter settings. It is observed that the subscale algorithm depicts better accuracy while DOC is best performer in terms of F1_Measure on synthetic datasets.

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Accuracy		
	AR	SRR
SUBSCALE		$\overline{2}$
DOC	5	3
MINECLUS	6	$\overline{4}$
SCHISM	3	⇁
FIRES		
INCY	4	8
PROCLUS	8	6
STATPC	9	$\mathbf Q$
P ₃ C		

Table 4 Accuracy on synthetic datasets

F1_Measure		
AR	SRR	
6	5	
$\overline{4}$	\overline{c}	
7	ℸ	
$\overline{2}$	3	
5	6	
3	$\overline{4}$	
9	9	
8	8	

Table 5 F1_Measure on synthetic datasets

(ii) Scalability: Scalability of cell-based, density-based, and clustering-orientedbased subspace algorithms is shown in Figs. [9](#page-17-0), [10,](#page-19-0) [11](#page-19-1), [12](#page-20-1), [13,](#page-20-2) and [14](#page-21-0). Graphs are represented in terms of data dimensionality. Scalability depicts the performance of the algorithm with increasing number of dimensions. It is shown on synthetic datasets as number of records/objects are constant and dimensions vary from 10 to 75. Synthetic datasets are given in [[17\]](#page-24-9). The X axis of graphs represents data dimensionality, and Y axis shows accuracy or F1_Measure.

Figures [10](#page-19-0), [11,](#page-19-1) and [12](#page-20-1) represent the performance of cell-based, density-based, and clustering-oriented-based algorithms, respectively, in terms of accuracy. It is observed from cell-based and density-based algorithms that SCHISM and INCY could not cope up after 25 dimensions and hence give results till D25 dataset only. MINECLUS accuracy varies randomly from dimensions to dimensions. DOC gives highest accuracy at 15 attribute dataset and then shows downfall. However, its accuracy shows slight improvement from D20 to D25 but again its performance declines. For density-based algorithms, FIRES shows improvement till D25, and then with

Fig. 10 Scalability of cell-based algorithms (accuracy)

Fig. 11 Scalability of density-based algorithms (accuracy)

slight downfall, its performance becomes stagnant. SUBSCALE shows improvement in accuracy after D25 dimensional dataset. SUBSCALE and FIRES give highest and approximately same accuracy at 75-dimensional dataset. For clustering-oriented-based algorithms shown in Fig. [12,](#page-20-1) STATPC shows highest accuracy at D75 dataset. PROCLUS performance falls after 25-dimensional dataset. P3C shows low performance for overall datasets.

Figures [13](#page-20-2), [14](#page-21-0), and [15](#page-21-1) depict the scalability of algorithms in terms of F1_ Measure. It has been noticed that subscale gives highest F1_Measure on the 75-dimensional dataset. However, the effcacy of DOC, FIRES, and P3C shows downfall in performance with an increase in dimensions.

It can be concluded from given experiments that on synthetic datasets, FIRES and DOC depict the best performance on the basis of accuracy and F1_Measure, while CLIQUE and MINECLUS are best performers on accuracy and F1_Measure on real datasets. That means cell-based and density-based algorithm is more appropriate to opt for subspace clustering.

The next section illustrates the various application areas along with future prospects.

Fig. 12 Scalability of clustering-oriented-based algorithms (accuracy)

Fig. 13 Scalability of cell-based algorithms (F1 Measure)

7 Applications and Future Prospects

In previous sections, it has been shown that the trend of using subspace clustering algorithms for high dimensional problems is rising. This section discusses the application areas where subspace clustering algorithms are suitable. Additionally developing amalgamated subspace clustering algorithms are suggested. Following are the application areas of high dimensional clustering:

(i) Collaborative Filtering: The other name of collaborative fltering is a recommendation system. It is a social fltering technique where information is defned on basis of recommendations given by people [[5\]](#page-23-4). People who like certain item in past are more likely to purchase in the future. People may like the items recommended by friends, neighbor, family, colleagues on social media, etc. Recommendations can be user-based or item-based. High dimensional clustering algorithms play an important role in such systems. The dataset is matrix of users and products. Clustering in such dataset retrieves the group of users liking the same product or group of items with relevant users. Hence subspace

Fig. 14 Scalability of density-based algorithms (F1_Measure)

Fig. 15 Scalability of clustering-oriented-based algorithms (F1_Measure)

clustering could be applied [[45,](#page-25-15) [46\]](#page-25-16). Some examples are Customer Recommendation System, Movie Recommendation System, etc.

- (ii) Computer Vision: This feld is based on mining useful information from single image or video to attain automatic visual understanding. The attributes extracted from image or video are large in number. An example of such dataset is image segmentation data [[47\]](#page-25-17). The dataset contains 2310 data objects and 19 attributes. Clustering techniques are applied to cluster in either shape group or RGB group [[33\]](#page-25-4). Similarly there a number of datasets with large attributes where clusters may exist in different subspaces [\[48](#page-25-18)]. Field of computer vision where subspace clustering can be applied is Facial Recognition, Gesture Recognition, etc.
- (iii) Biological Dataset: Gene expression dataset is the most widely used dataset where subspace clustering can be applied [\[5](#page-23-4)]. Microarray DNA is a technology which measures a large amount of genes expressions under different circumstances. In order to understand the various types of diseases caused by genetic disorder at different levels, subspace clustering is required [\[49](#page-25-19)].
- (iv) Text Documents: Clustering in text documents is an important task for web mining. Web pages are clustered on the basis of frequency of terms occurring in the page. Text documents are represented as high dimensional feature vector, where each feature is a frequency of term in document. Each document is represented by a data record/object. Hence the dataset formed from text document is high dimensional data [\[50](#page-25-20)]. The cluster of related document may exist on basis of similar word count, themes, etc. Hence subspace clustering techniques are applicable in such datasets.
- (v) Distributed Databases: Massive amount of data is being dissipated by a number of sources like social media, microarray DNA, etc. Nowadays such a large amount of data is stored in different physical locations named distributed databases. The massive data is fragmented so that it can be distributed over multiple servers and parallel queries can be executed. Fragmentation can be row-wise (horizontal) or column-wise (vertical). Subspace clustering techniques can be applied to aid the fragmentation process. However, no work has been done yet on this application using subspace clustering approach.
- (vi) Social Network: The data produced from social media is large in volume. Clustering can be performed on the social network to fnd infuential groups in the network. This task is achieved by analyzing the linkages (edges) and nodes of the network. Clustering can be done on bases of various attributes associated with each node depending upon the objective in hand. Determining influential groups on the basis of various topics can be helpful in promotional activities, market segmentation problem, community detection problem [[51\]](#page-26-0), etc. Hence subspace clustering can be applied to fnd the groups in subsets of dimensions. However, a little amount of work is done on this application using subspace clustering approach.

In recent years, subspace clustering is being used with metaheuristic approaches. Nature-inspired algorithms [\[52](#page-26-1), [53\]](#page-26-2) are the prominent metaheuristic techniques that are used for determining the near-optimal solution to complex and hard problems. The frst hybrid approach of subspace clustering with an evolutionary algorithm is proposed in [\[54](#page-26-3)]. With the advent of new metaheuristic algorithms, subspace clustering results could be improved [\[30](#page-25-1), [36](#page-25-7)]. Some other algorithms can be developed amalgamating artifcial bee colony algorithm, grey wolf algorithm, etc., with various subspace clustering algorithms. The hybrid algorithms developed were not applied to the applications described above. This will give a new direction of future work to researchers in the feld of subspace clustering of high dimensional data.

8 Conclusion

Subspace clustering fnds the clusters existing in various subsets of dimensions. The chapter presents a comprehensive survey on subspace clustering approaches, evaluation metrics, and application areas. The chapter also reveals the signifcance of subspace clustering in literature by presenting the statistical data. Comparison of conventional subspace clustering algorithms is also depicted through average ranking and success rate ratio ranking. Performance assessment of algorithms is made through scalability on basis data dimensionality. The chapter answers the following research questions:

(i) What are the major challenges faced by traditional clustering algorithms to cluster high dimensional data?

ANSWER: The problem of the curse of dimensionality is described in the frst paragraph of Sect. [1](#page-0-0).

(ii) What search techniques are being used in subspace clustering to determine subspaces?

ANSWER: Top-down and bottom-up search techniques used to fnd subspaces and are described in Sect. [3](#page-3-1).

- (iii) What are evaluation measures for comparing subspace clustering algorithms? ANSWER: Different evaluation measures are described in Sect. [4.](#page-6-0)
- (iv) What is the current scenario of subspace clustering? ANSWER: Literature survey on subspace clustering with statistical data is illustrated in Sect. [5.](#page-9-0)
- (v) What are the research gaps in the literature and the future prospects of subspace clustering?

ANSWER: Research gaps are given in Table [1](#page-11-0), and future prospects are presented in Sect. [7](#page-20-0).

Thus, the chapter is useful to the researchers planning to work in the feld of subspace clustering. Additionally, it suggests the algorithms to develop in the future along with application areas.

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