Data Mining Project PT. 2 - K-Means and DBSCAN analysis and implementation

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Abstract

This report is on the development and analysis of two Data Mining clustering algorithms, namely K-Means clustering and DBSCAN, within an interactive graphical program written in Java programming language. This report is dived in three parts. In the first part we will present a usage section, explaining how to interact with the program. We will then introduce the architecture of the system, explaining our implementation in detail. In the third part we will present our analysis of the algorithms

1 Getting Started

1.1 Introduction

The DWDM Project part II consists in implementing some Data Mining algorithms to get a deeper understanding of their behavior, their strengths and their weaknesses. We choosed to implement the K-Means and the DBSCAN algorithms performing the same task and then we compared their behaviours.

1.2 Contents of the archive

When un-compressed, the archive is organized in the following structure:

dwdmproject.jar is the Java Archive of the program we implemented, the datasets sub-folder contains the two datasets on which we tested the application. The javadoc subfolder contains the exported documentation of the program.

1.3 Running the application for the first time

First, we launch the JAR file. This can be done either by double clicking on the JAR file or by executing the below command in a console window:

```
java - jar dwdmproject. jar
```

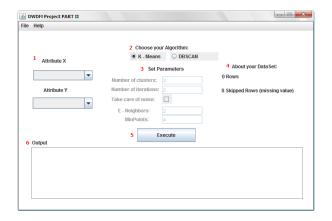


Figure 1.1: Quick overview

Our application is tested and supported for Windows XP and Windows 7. There are little unexpected behaviors when running it under Mac Os X but they are all related to the graphical implementation of the project: the functionalities seem working fine anyway. There are bigger issues with Gnu/Linux, all related to the graphical implementation: some functionalities will not run.

1.4 Quick overview

As the application starts all features are disabled. This is because there is not a dataset loaded yet. The application main window, represented in Figure 1.1, is composed by the following parts:

- 1. Attribute X + Attribute Y: we get the possibility to select the attributes on which the algoritms will perform their calculations;
- 2. Choose your algorithm: we select which algorithm to be run against the dataset
- 3. Set Parameters: in case of K-Means algorithm selected, the available parameters to be set are the number of clusters, the number of iterations and whether to take care of noise. If the DBSCAN algorithm is selected, we have to specify only the values for E-Neighbours and Min-Points;
- 4. About your Dataset: it displays the number of rows loaded into memory and the number of skipped rows because of missing values;
- 5. Execute: this bottom executes the selected algorithm;
- 6. Out: algorithms' outcome.

1.5 How to run K-Means algorithm

The steps for running the K-Means algorithm are the following:

• First we need to load our dataset into the memory. This can be done by clicking on menu File > Open, then select either "census-income.arff" or "blood-transfusion.arff" provided as attachment. Note, the loading window allows to import only arff file, otherwise our parser is not able to parse its content.

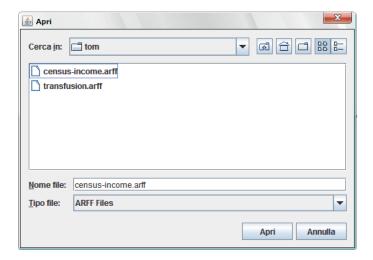


Figure 1.2: Open window

• Important: while importing the dataset, the application checks wheter missing values are present. This information will be diplayed later in the "DataSet information" section described before. In case of a dataset with missing values in every row, the application informs the user that it has no data for running the algorithms.



Figure 1.3: No data popup

• Once the dataset has been loaded, we can decide on which attributes to perform the K-Means algorithm. We selected as attributes "age" and "relationship", 2 clusters and 2 iterations. We also wanted to take care of the noise.

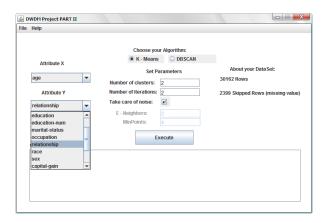


Figure 1.4: Selecting attributes

- We click the Execute button;
- In case of big dataset size, the execution of the K-Means algorithm takes a while. For this reason we implemented a progress bar to inform the user how long the execution is going to take.

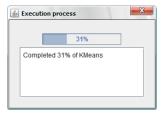


Figure 1.5: Progress bar

• Once the progress bar finishes, the user sees the outcome of the K-Means algorighm:

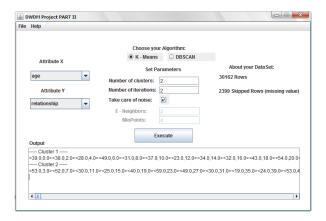


Figure 1.6: K-Means outcome

1.6 How to run DBSCAN algorigthm

The steps for executing the DBSCAN algorithm are basically the same of the K-Means algorithm. The only things that change are the parameters to be set. In fact, we have to specify the number of E-Neighbors and the number of MinPoints.

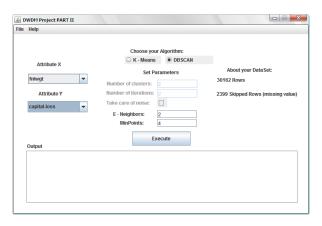


Figure 1.7: DBSCAN Algorithm

2 Architecture

For implementing the system we decided to follow a personalized Model-View-Controller design pattern, that we name "MVA", or Model-View-Algorithm, respectively mapped in the Java packages **types**, **gui** and **algorithms**. This is for a better separation of concerns. Figure 2.1 illustrates with a very simple scheme our general view of the architecture of the system. It is a personalized version of the MVC pattern because we further separated the

algorithms from the controller part, putting the non algorithm-related code in the **gui** package anyway. The user interacts only with the **gui** component that, based on user input, calls the components of the **algorithms** package that will do some computation on data defined in the **types** package and return the result.

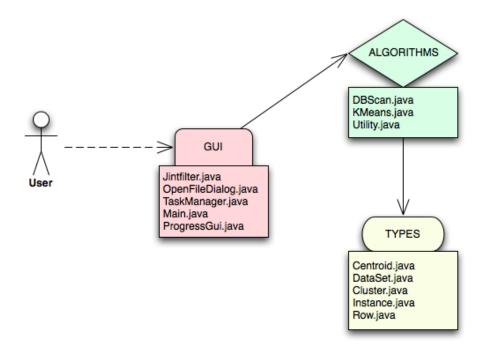


Figure 2.1: General Architecture View of the System

The **gui** package contains the Main class that is responsible for accessing the program and constructing the graphical user interface using Swing. The Jintfilter is a specialization of Swing's Document object and is used for filtering user input on algorithm parameters. The OpenFileDialog class is a personalized dialog for opening the datasets, in ARFF format. The ProgressGui class is responsible for creating a window containing the status and progress of the operations. The TaskManager class is a wrapper for creating threads that contain the algorithm tasks.

The **algorithms** package, better illustrated in Figure 2.1, contains the pure, type independant implementations of the clustering algorithms, K-Means and DBSCAN. They make use of generalized data types defined in the **types** package, in order to have an abstraction of the data that could be loaded from any datasource, being it an ARFF file, a CSV, and XML or a Database. The Utility class actually contains the method for loading the datasets and converting them in objects declared in types package.

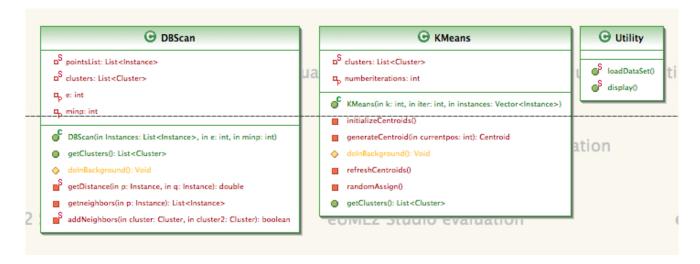


Figure 2.2: Algorithms Package

The **types** package, represented in Figure 2.3, contains pure Java representation of the data that can be loaded by the algorithms. They are an abstract representation of an ARFF file but they are completely independent from their structure. They act as wrappers for the algorithm and it would be a matter of changing the Utility.loadDataSet() method for loading any kind of data format. The DataSet class represents a data format, e.g. a dataset, that is composed by rows, defined in Row class. The other classes are related to the algorithm own data types, for example Cluster and Centroid.

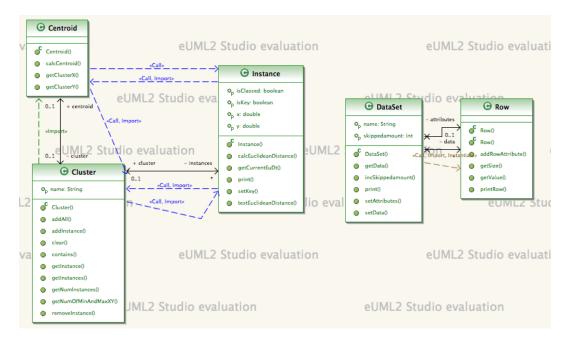


Figure 2.3: Types Package

All the code has been clearly documented using Javadoc documentation. For a deep understanding of the functioning of each method, please see the javadoc folder, containing the HTML exported documentation.

Regarding the additional method we created in order to achieve the results, they are all documented in the code and in the documentation, but we will mention here one of them. It is a method for removing noise from the datasets. It begins at line 143 of it.unibz.algorithms.KMeans. It is inspired by [Barca, J.C. Rumantir, G.] A Modified K-Means Algorithm for Noise Reduction in Optical Motion Capture Data.

In order to take care of noise, we don't reduce it before executing the algorithm, but we added an additional check during the execution. On every end of iteration, we try to find out how compact a cluster is. If instances

inside a cluster overcome the average compactness of the clusters in the dataset, this cluster is flagged as noise. In modification to the classical K-Means algorithm, we try to add an automatic determination of the optimum number of clusters we should have. A cluster is considered noise if it only has a few instances in it. The minimum number of instances in a cluster, or the cluster size, are set such that it minimizes the degree of false positives (i.e. data clusters incorrectly classified as noise) and false negatives (i.e. noise clusters incorrectly classified as data). The following is a code snippet that handles the problem:

3 Analysis

3.1 Comparison of the algorithms

Two clustering algorithms - namely the K-Means and the DBSCAN - were examined and their clustering performances were compared. First of all, in the DBSCAN algorithm the user does not have to specify the K value, which drastically affects the outcome in the K-Means algorithm. The DBSCAN deals with noise data and outliers. On the other hand, the K-Means algorithm is affected by noisy data. Additionally, in the DBSCAN the ordering of the points in the database is not so important. We found that the DBSCAN algorithm has a superior performance.

3.2 An algorithms analysis

3.2.1 K-Means Properties

- Pro: K-Means is relatively scalable and efficient in processing large data sets
- Pro: the computation complexity of the algorithm is O(objects*clusters*iterations)
- Con: it can be applied only when the mean of a cluster is defined
- Con: the user must specify a value for K parameter
- Con: it is not suitable for discovering clusters with nonconvex shapes or clusters of very different size
- Con: it is sensitive to noise and outlier data points (can influence the mean value).

3.2.2 DBSCAN Properties

- Pro: it does not force you to choose the number of clusters in the data a priori, as opposed to K-Means
- Pro: it can find arbitrarily shaped clusters. It finds clusters completely surrounded by a different cluster
- Pro: it considers noisy values
- Pro: it requires just two parameters for working
- Pro: it is mostly insensitive to the ordering of the points in the database
- Con: the distance measure in the function for finding neighbours is crucial. We used the euclidean distance measure and especially for high-dimensional data, this distance metric can be rendered almost useless
- Con: varying densities affects the outcome of this algorithm (hierarchical data sets)

3.3 Benchmarking the algorithms (AMD Phenom 9600 Quad Core $2.6 \mathrm{Ghz}$)

DataSet Type	K-Means (C=20, I=2)	K-Means (C=20, I=2, Noise removal)	DBScan (E=5, MinPoints=20)
BT (748 Rows)	4.5 ms	$5~\mathrm{ms}$	$7~\mathrm{ms}$
BT*2 (1496 Rows)	30 ms	33 ms	$3 \operatorname{Sec} 498 \operatorname{ms}$
CI/2 (15124 Rows)	3 Min 40 sec	4 Min 14 sec	2 Min 04 sec
CI (27763 Rows)	9 Min 19 sec	10 min 40 sec	4 Min 18 sec