Analysis of correlation relation between pollution and petroleum consumption using algorithms (spectral clustering and K-mean)

by

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DEDICATION

To my father and mother

And to my wife thank you for helping me

And everyone who encouraged me to go to the sea of knowledge

And thanks to all those who support me

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ABSTRACT

Analysis of correlation relation between pollution and petroleum consumption using algorithms (spectral clustering & K-mean)

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Clustering can be considered the most significant unsupervised learning methods that reveal similar behaviors (sets) on large sets of data.

Clustering is the process of organizing objects into groups that are similar in some way to their members.

Spectral clustering data points as nodes of a connected graph and clusters are found by partitioning this graph, based on its spectral decomposition, into sub graphs.

K - means assembly: Divide objects into k groups so that some measurements are minimized relative to the middle points of the clusters.

In this thesis, we have applied both of algorithms (spectral clustering and k-mean) on data from the life process and these data are statistical data. The field of data is the quantities of pollution and petroleum consumption. The spectral clustering algorithm was applied several times and

k-mean was applied several times and the best result was compared and the best classification was also obtained.

Keywords: Clustering ,Spectral Clustering ,K-mean ,pollution, petroleum.

Table of content

LIST OF ABBREVIATIONS

1. INTRODUCTION

Constant advance in science and technology makes collection of data and storage much easier and very inexpensive than ever before. This led to form large data sets in science and government Industry, which should be processed or sorted for useful information.

For example, if we consider the results generated by a search engine for a specific query, the user has to clear through the long lists and find them the solution is required. But this task can be very difficult for the user if there is Millions of WebPages have been shown as a workaround for some queries. Etc. and so on Partition techniques can be very useful in a closely related compilation.

Specific query solutions and display results in the form to set so that even documents that have no relationship to each other can be uninstalled without.

That behind compiling any set of data is to find the authentic Structure in the data, interpret this structure as a set of groups, Data objects within each group must appear very high of the similarity known as the similarity within the mass, while the similarity.

1.1 CLUSTERING

Clustering is a Data collection is a way to make similar objects in groups Thus, same objects are in the same group and different objects are also in different groups Data clustering is considered as an unsupervised learning technique in which objects are grouped in unknown predefined clusters. On the contrary classification is a supervised education in the objects are set to predefined classes clusters.

Figure. 1.1 clustering

1.2 Basic Concepts Of Clustering

The problem of data clustering can be formulated as follows: given a dataset **D** that contains n objects x1, x2, xⁿ (data points, records, instances, patterns, observations, items) and each data point is in a d-dimensional space, i.e. each the data points has A dimensions (attribute, feature, variable, and component that is Can be expresses in the form of a matrix as allow.

$$
A = \begin{bmatrix} x1.1 & \cdots & x1.n \\ \vdots & \ddots & \vdots \\ xn, 1 & \cdots & xn.n \end{bmatrix}
$$

Data clustering is based on the similarity or dissimilarity (distance) measures between data points. Hence, these measures make the cluster analysis meaningful [28]. The high quality of aggregation is obtained in strong similarity in mass and weak similarity between groups as shown in Figure 1.2. In addition, when we use the dissimilarity (distance) concept, the latter sentence becomes: the high quality of clustering is to obtain low intra-cluster dissimilarity.

Figure 1.2 . Inter-clustering and Intra-clustering similarities of clusters.

1.3 Type Of Clustering

We can conduct groups in a number of different ways. Each assembly technique produces several different types of cluster.

Some take input parameters from the user like number clusters to be formed etc., but some decide on the type and amount of data given. The main developments have been the introduction to density based and grid based clustering methods. We can classify cluster algorithms into five different types.

- \checkmark Partitioning methods
- ✓ Hierarchical methods.
- \checkmark Model-based methods.
- \checkmark Density based methods.
- \checkmark Grid based methods.

1.3.1 Partitioning Methods.

The easiest and most important version of cluster analysis is the division that organizes group objectives in many exclusive groups or clusters. To keep the problem specification concise, we can assume that the number of clusters is given as background knowledge. This parameter is the first point of division methods.

The data set comes out of n objects formally and k is the number of groups to form a partitioning algorithm that arranges objects in k kn sections where. cluster is configured to improve an objective partitioning criterion such as a difference function based on the distance so that the objects are within a block, So the objects include the group "identical". If a database containing n data objects is given, then a partitioning

method constructs k clusters of the data where k<=n and k is that the input parameter provided by the user. That is, it classifies the data into groups which should satisfy the following conditions: (1) each group must contain at least one data object and (2) each data object ought to belong to just one group. The second demand becomes easy in fuzzy k mean agglomeration at intervals that one object square measure typically resembled by two or lots of teams. With k as the given number of partitions to be MADE the partitioning technique creates an initial partition methodology creates an initial partition method MADES an initial partition. Then more number of iterations are followed in which objects are moved from one group to other making sure that in-cluster similarity is more than similarity with Objects in another cluster.

Popular Partitioning methods k means and k-means the most well-known and usually used partitioning methods are k-means proposed by (Mac Queen 1967) and k-means proposed by (Kaufman and Rousseau 1987).

Figure 1.3. Original Points and **A partitional clustering**

1.3.2 Hierarchical Methods

The set of given data objects are partitioned in form of a tree like structure or nested clusters in hierarchical clustering. The hierarchical They were classified into two kinds.

Agglomerative

LE Divisive

In agglomerative method also known as bottom-up approach, Forms of all objects separate group. It successively merges the groups close tone another by checking the similarity function, until all the groups are merged into one, that's until the top most level of hierarchy is reached or until a termination condition holds. In divisive clustering also known a stop-down approach, initially all the objects are grouped into a single cluster which can also be called as parent. The repetition process is performed each time the object is divided into group until the object becomes in one group or it may be alone.

1.3.2.1 Agglomerative Method

This method begins by treating each object as an individual cluster and then proceeds by groping two nearest clusters. The distance between any two clusters m and n is defined by a metric $D_{m,n}$. Metrics can be single-link, complete-link and group average etc. A general class of metrics was given by Lance and Williams [1]. If Dk , ij be the distance between cluster k and the union of cluster *i* and cluster *j*, then: Dk , $ij = iDk$, $i + jDk$, $j + \beta Di$, $j + \gamma |Dk$, $i - Dk$, $j|$ The agglomerative method is as follows:

Consider each object to be an atomic cluster. The $(n \times n)$ distancematrix represents the distance between all possible pairs of clusters. Find the smallest element in the matrix. This corresponds to the pair of clusters that are most similar. Merge these two clusters, say and n, together. Measure the distances between the newly formed cluster and the other remaining clusters using a distance function. Delete the rowan column of m and overwrite row and column of cluster n with the new values. If the current number of clusters is more than k then go to step 2. otherwise stop. The merging process can continue until all the objects are in one cluster. The advantages of hierarchical methods are that It is therefore easy to implement computationally. They can tackle larger datasets than the k-means method and we can run the algorithm without providing the input k (the number of clusters to be formed). The drawbacks of agglomerative method are:

The algorithm has $O(n3)$ time complexity. Even though the order of the distance matrix decreases with each iteration, the cost of Step2 on iteration k is $O((n - k)2)$, and we are guaranteed $(n - k)$ iterationsbefore we get to k;The clusters produced are heavily dependent on the metric Di , j. Different metrics can produce different clusters. For instance, the completelink metric tends to produce spherical clusters, whereas the single-link metric produces elongated clusters [1].

1.3.2.2 Divisive Method

The contrast procedure of agglomerative clustering is the divisive method. Initially all the data objects are considered in one cluster. Then for each object the degree of irrelevance is measured and the most irrelevant data object is split from the main cluster and a new cluster is formed with only that data object in it. The highest degree of irrelevance of an object corresponds to the one that is most distant from all other objects in that cluster. Let the average distance between object *i* and then cluster *Cj* be defined as $[1]:Di, Cj$ =The most irrelevant object splits off and forms a new cluster. This is equivalent to splitting the cluster with the bigger diameter. The process will continue until it perfect the specific termination requirement or the required crusts above a certain threshold distance. These methods face the difficulty of making a right decision of splitting at a high level. The algorithm for divisive method is [1]:

- $\ddot{}$ Choose the cluster that has the most remote pair of objects. Thesis the cluster with the largest diameter.
- In this cluster, we find the object with an average distance greater than other objects. Remove the object from the cluster, allowing it to form a new atomic cluster.
- For object h in the cluster being split, calculate the average distance between it and the current cluster; and the average distance between the object and the new cluster. If the distance to the new cluster is less than the distance between it and the16-current cluster, move the object h to the new cluster. Loop over althea objects in the cluster.
- If no objects can be moved, but the current number of clusters is greater than k, go to step1. Otherwise stop. The drawbacks of divisive method are:
- \downarrow The time complexity of algorithm is O(n3), O(n2) on the step1 of the algorithm for each iteration. Moreover, there are expensive calculations that may take place in step3 of the algorithm.
- In step 3 the group averages between an object and the new and existing clusters need to be recalculated after an object is moved.

This will be costly in terms of number of calculations and the amount of storage required.

 \pm The method only searches one of the $N(n, k)$ possible partitions.

6

In hierarchical clustering once a split or merge is done, it cannot be undone. This fact acts as both key to success and drawback for hierarchical clustering. The firmness of hierarchical method leads to less computational cost without a combinatorial number of choices but the main problem with it is invalid decisions cannot be corrected.

Hierarchical clustering methods are simple but encounter problems at making critical decisions for selection of correct merge are split points. Such a decision is critical because once a group of objects is merged or split, the process at the next step will work on the newly generated clusters. It will ne'er undo what was done antecedently nor perform object swapping between clusters therefore merge or split if not done wise might end in caliber clusters. Thus, merge or split if not done wise may result in low quality clusters. These ways have scaling downside since the choice of merge or split must examine and measure a decent variety of objects or clusters Hierarchical clustering can be improved Processing By integration this technique with alternative cluster techniques for multiple part cluster. One such technique known as birch 1st partitions objects hierarchically use tree structures then applies different cluster techniques to provide refined clusters.

Figure 1.4. Agglomerative and Divisive

1.3.3 Model-based methods

Each component is described by a density function and has an associated probability or "weight" in the mixture. In precept, we will able to} adopt any probability model for the parts however usually we'll assume that parts are p variety normal distributions. (This does not necessarily mean things are easy: inference in tractable, however).

1.3.4 Density Based Methods

Density-based clustering ways are supported a local cluster criterion. Clusters area unit assumed as regions within the data area during which the objects area unit dense and therefore the clusters area unit separated by regions of low object density. These regions have a spot shape and the data points inside a cluster may be spot distributed.

1.3.5 Grid Based Methods

A grid based mostly clustering technique takes inside the thing space and quantizes it into affinity range of cells forming a grid structure. Then the method performs all the operations on it grid structure. The main advantage of this methodology is its quick interval that is freelance of the quantity of objects, and dependent solely on the quantity of cells in each dimension within the quantized area. Grid-based ways use one uniform grid mesh to partition the whole downside domain into the cells and the data objects settled within a cell area unit represented by the callusing a set of statistical attributes from the objects. cluster is performed on the grid cells, instead of info itself. Since the scale of the grid is far but the quantity of data objects, the process speed will be significantly improved.

1.4. Importance of Clustering

Data clustering is one of the main tasks of data mining [1] and pattern recognition [2]. Moreover, it can be used in many applications such as:

- 1. Data compression [3].
- 2. Image analysis [5].
- 3. Bioinformatics [6].
- 4. Academics [9].
- 5. Search engines [79].
- 6. Wireless sensor networks.
- 7. Intrusion detection [81].
- 8. Business planning [82].

9. Add another application to use the monastery These are social and statistical studies of real life data Such as the rate of crime and unemployment in a statistical study as well as pollution and its relationship to oil consumption, pollution and the production of oil. This is the focus of our study on this proposal using dye in the classification of countries in terms of pollution, oil consumption, pollution and oil production.

2. DATA CLUSTERING

data clustering may be a crucial obstacle with many applications in

- $\textcolor{red}{\blacktriangleleft}$ Machine learning.
- \leftarrow computer vision.
- $\frac{1}{2}$ Signal method.

the item of clustering is to divide a dataset into natural groups such as: Points among constant cluster area unit similar. Points incurious groups area unit dissimilar to

each different. Clustering methods can be: Hierarchical: Single Link, Complete Link, etc.

Partitioned or flat: k-means, Gaussian Mixture, Mode Seeking, Graph partitioning.

Figure 2.1. classification of clustering algorithms

2.1 Clustering Algorithms

There are many types of cluster algorithms, we will focus on only two types, which are used in thesis. K- means cluster and spectral cluster. But the study was applied to spectral cluster**.**

2.1.1 K-means cluster Algorithms

Some objects are separated into groups such as K cluster. Some groups are reduced to centroids of the clusters is minimized.

Figure 2.2. k-mean Clustering

2.1.1.1 Pros

- It's easier to understand.
- comparatively active.
- We get a better result when they are separate. So that each group is separated by a special center.
- You only need to calculate distances between the center only until its center is configured. The distance between the inner points is not measured.
- The following figure illustrates the concentration of data and the composition of the class.

2.1.1.2 Cons

- The education algorithm order apriority specialization of the number of cluster centers.
- Euclidean, space menstruation can unequally be weighing underlying factors.
- The education, algorithm supply the local optima of the square mistake function.
- The learning algorithm provides the native optima of the square error function.
- Random configuration of the cluster center may not lead us to a useful result.
- The application is only -used when finding and defining the average any failure in the class data Unable to handle extreme data the algorithm failed to set nonlinear data.

2.1.2 Spectral Clustering Algorithms

It is considered one of the best and most important algorithms in the extraction of data, and useful in the search for a large amount of data, and this is through the process of linking the analysis of these data and methods of artificial intelligence to become the best efficiency in the search process the Clustering is a process of data from similar clustering, a branch of data mining. The aggregation algorithm divides data sets into several groups, where the similarities between points within a group are greater than the similarities between two points within the two different communities. The idea of simple data collection in nature and very close to the human in his way of thinking where we whenever we deal with a large amount of data tend to a huge amount of data to summarize a few groups or categories, to facilitate the analysis process. Not only used to organize and classify the data, but used in data compression and build the

sample system algorithms to collect data widely. If data sets are found, it is possible to build

a problem model based on those groups. The clustering, in data extraction, can detect clusters and identify interesting distributions in basic data. It is illustrated by this figure (1). The data points are also nodes of connected.

It studies the relationship between the data points themselves and then distributes them to groups where data in the same group must be connected and coherent.

Figure 2.4. spectral clustering

2.1.2.1 Cons

- ❖ Strong assumptions cannot be obtained from cluster statistics.
- ❖ It features easy implementation.
- ❖ Reasonable clusters results.

2.1.2.2 Pros

- ❖ Some of the times are sensitive to the selection of parameters.
- ❖ When large data is made, it costs a lot to calculate.
- ❖ The data set that can be able to execute is not common and in real life.

2.2 ALGORITHMS (k-means and spectral)

2.2.1 K-mean clustering algorithm

These centers should be placed in a boring manner because of a different location causing a different result. Therefore, the best option is to put them as far as possible apart from each other. The next step is to take each point belonging to a particular data set and connect it to the nearest center.

To apply the algorithm correctly, it should be positioned as far as possible apart from each other. The next step is to take each point belonging to a particular data set and connect it to the nearest center.

When no point is outstanding, the primary step is completed and also the group is aged early. At this time, we'd like to calculate the new C-centroid because the Paris center of the clusters ensuing from the previous step. when we've these new centroid k. a new binding must be done between an equivalent data set points and the closest new center. A loop has been generated. As a result of this loop we could notice that the k centers amendment their location step by step till no additional changes are done or in alternative words centers don't move to any extent further. Finally, this algorithmic rule aims at minimizing AN objective function know as square error function given by

$$
J(V) = \sum_{i=1}^{c} \sum_{j=1}^{ci} (||x_i - v_j||^{2})^2
$$

Where:

 $||x_i - v_j||'$ is the Euclidean space between x_i and v_j ' c_i ' is the number of data points in I the cluster. $'c'$ is the number of cluster centers.

2.2.1.1 Algorithms steps for k-means clustering

Let $X = \{x_1, x_2, x_3, ..., x_n\}$ be the set of data points and $V = \{v_1, v_2, ..., v_c\}$ be the set of centers.

1. Random selection of 'C' cluster centers.

- 2. Calculate the distance between every data point and cluster centers.
- 3. Let the data point to the cluster center whose distance from the cluster center is minimum of all the cluster centers.
- 4. Assign the data point to the cluster center whose distance from the cluster center is minimum of all the cluster centers.
- 5. Repeat the new cluster center account using:

$$
v_i = (1/c_i) \sum_{j=1}^{c_i} x_i
$$

where, $'c_i'$ represents the number of data points in it cluster.

- 6. Recalculate the distance between every data point and new obtained cluster centers.
- 7. If no data point was reassigned then stop, otherwise repeat from step 3).

Figure 2.5. Showing the result of k-means for *'N'* **= 56 and** *'c'* **= 7**

Figure 2.6. k-mean Clustering 1-5

2.2.2 Algorithms steps Spectral Clustering

supposed data from, read it from outer file to the matrix.

 A=

Find affinity matrix A $[n, n]$ from matrix $A[n, 2]$ ${\it Affinity}\ =\ \frac{1}{\sqrt{N}}$ $e^{dis/2\sigma^2}$

where d is the distance between every Element in and others in the same matrix A by the low

dis
$$
(p_i, p_j) = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2}
$$
 Where $p_i = (x_i, y_i) \& B_j = (x_j, y_j)$.

Considering **K** is a parameter express a number of a suitable, vectors will be select from eigenvector in common, similar, to number of clusters k+1 to apply neigvec.

The scaling parameter σ^2 has a control over how fast this approximation falls with the distance between p_i , p_j Two Local Scaling is a measure of two similar points. This provides an intuitive way to determine possible values for σ. The chosen of σ is generally done manually. propose, choosing σ automatically by running their clustering algorithm frequently for a number of values of σ and selecting the one which supply lower distorted clusters.

We note. In addition, the set of values to be tested must still be determined manually. Furthermore, when input data includes groups with different local statistics, there may note a single value σ that works well for all data. The significant effect σ on the cluster. When the data contains multiple scales, even using optimization σ fails to provide a good set, the local scaling effect. The affinities across clusters are now safely, lower than the affinities within any single cluster. To set the value of σ , we calculate it for finding the Affinity of each two points *i* and *j* First we must analysis σ 2 to σ_i , $\sigma_j \sigma_i = \textit{dis}(s_i - s_k)$.

We considers k is K th neighbor of sipoints. The choosing of K is not dependent or independent of the scale and is the dimension function of data from the embedding area. However, in most experiments (both on synthetic data and on images) one value of $K = 3$ was used, which yields good results even for high-dimensional data $\sigma_i = dis(s_i - s_k)$.

Figure 2.1 Affinity Matrix

computation the degree matrix the grade of matrix Degree is the same as that of matrix Affinity Matrix taking into consideration. Calculate the sum of each row of matrix Affinity in a position equal to row and column Degree matrix.

Degree [*i*, *j*] =
$$
\sum_{j=1}^{n}
$$
 Degree [*i*, *j*] If *i* = *j if i*=*j*
Degree [*i*, *j*] =0 *if i* ≠ *j*

Consider n is a grade matrix with all elements of zeros except the main diameter.

Sum 1st row			
	Sum 2st row		
		Sum 3st row	
			Sum n _{st} row

Figure 2.2 .Degree Matrix

Find an affinity matrix for Laplacian normalization through these low levels and is used to find

each element in the N.Lap matrix Normalized L.N= AFFINIT[I,J] $\sqrt{degree[i,i]} * \sqrt{degree[j,j]}$

N.laplacian[1.1]	N.laplacian[1.2]	N.laplacian[1.3]	0.0.0	N.laplacian
N.laplacian[2.1]	N.laplacian[2.2]	N.laplacian[2.3]	0.0.0	N.laplacian
N.laplacian[2.3]	N.laplacian[3.2]	N.laplacian ^[3.3]	0.0.0	N.laplacian
000	0.0.0	0.0.0	0.0.0	000
N.laplacian[1.n]	N.laplacian[n.2]	N.laplacian[n.3]	0.0.0	N.laplacian

Figure 2.3. Normalized Laplacian Matrix

• **Decomposition(** linear algebra)

E.vectors [1.1]	E.vectors $[1.2]$	E.vectors $[1.3]$		E.vectors $[1.4]$
E.vectors [2.1]	E.vectors [2.2]	E.vectors [2.3]	\cdots	E.vectors [2.4]
E.vectors [3.1]	E.vectors [3.2]	E.vectors $[3.3]$	\cdots	E. vectors[3.4]
	\cdots	\cdots	\cdots	
E.vectors [n.1]	E.vectors [n.2]	E.vectors [n.3]	\cdots	Eigenvectors [n.n]

Figure 2.4. Eigen values, Eigenvectors

Vn	Vn-1	$Vn-2$	$Vn-3$
v(1,n)	$v(1, n-1)$	$v(1, n-2)$	$v(1, n-3)$
v(2,n)	$v(2, n-1)$	$v(2, n-1)$	$v(2, n-3)$
\cdots	\cdots	\cdots	
v(n,n)	$v(n, n-1)$	$v(n, n-2)$	$v(n, n-3)$

Figure 2.5. Eigenvalues, Eigenvectors

Figure 2.6. Rearrange Neigenvec

Figure 2.7 .Normalized Matrix

Ξ

3- METHODLOGY

Overview: This search is described by applying this algorithm. All data is real, which is one of the advantages of this study. Real and realistic data were obtained from the European Economic Association website. a world-class statistical data center. In this study, we will focus on the pollution data from the European statistical site EEA and its correlation relation with oil consumption in 31 EU countries from 2006 to 2014.Throughout this study, we will identify the countries most affected by pollution and petroleum consumption by applying the algorithm (**Spectral**, **K-means**)and phases of implementation.

3.1 General stages of implementation.

First phase.

Obtaining data from the warehouse which mostly unworkable. After the data are obtained, it is processed and prepared using a number of data mining techniques.

Second phase.

The spectral and k-means algorithm are implemented more than once on the algorithms to obtain a good result.

EXECUTE PHASE.

Results: We obtain a graphical pattern showing the distribution of the cluster in the two algorithms(spectral and k-means).

Final phase.

The results are checked in terms of data classification and as to whether there is a correlation relation.

Figure 3.1 .phases Of implementation

3.2 Preprocessing Data Phases.

The following figure, in general, illustrates a number of data mining techniques**.**

Figure 3.2 .phases preprocessing data.

3.2.1 Data Selection

The data to be analyzed is specified to arrive at a logical classification.

3.2.2 Data Clean

Exclude incomplete data and data not compatible with the remaining data.

3.2.3 Data Set 1

Data Set 1 includes the quantities of emissions of pollution in tons in 31 countries in the European Union for the 2006 to 2014 period **(**in Belgium, Bulgaria, Czech Republic, Denmark, Germany, Estonia, Ireland, Greece, Spain, France, Croatia, Italy, Cyprus, Latvia, Lithuania, Luxembourg Hungary, Malta, Netherlands, Austria, Poland, Portugal, Romania, Slovenia, Slovakia, Finland Sweden, United Kingdom, Norway and Turkey). This group shows the amount of pollution and the amount of oil consumption in tons.

Table. 3.1 .Data set 1

3.2.4 Data Set *2*

Quantities of oil consumption per ton. And 31 countries - in the European Union and for the period 2006 to 2014 **States are**: (Belgium, Bulgaria, Czech Republic, Denmark, Germany, Estonia, Ireland, Greece, Spain, France, Croatia, Italy, Cyprus, Latvia, Lithuania, Luxembourg Hungary, Malta, Netherlands, Austria, Poland, Portugal, Romania, Slovenia, Slovakia, Finland Sweden, United Kingdom, Norway .and Turkey) from 2006 to 2014. In this group shows the amount of oil, and the amount of oil consumption by tons.

Table. 3.2 .Data set 2

3.2.5 Data Transformation

At this stage, the data is converted into custom templates to suit the search and retrieval procedures. Through the compilation process, in our study, the years are converted into columns and placed in a sequential order from 2006 to 2014. with 31 countries in the European Union, maintaining the sequence. The following figure shows pollution data.

Countries	YEARS	POLLUTION
Belgium	2006	133,599
Bulgaria	2006	765,131
\cdots	\cdots	\cdots
Belgium	2007	124,092
Bulgaria	2007	820,614
Belgium	2010	60,615
Norway	2014	16,633
Turkey	2014	2,147,499

Figure. 3.3 .Data transformation

The following figure shows oil consumption data for these countries and for the same years starting from **Belgium** to **Turkey** between **2006** and **2014,** as follows.

Figure. 3.4 .Data transformation

3.2.6 Data Integration

At this stage, similar and relevant data are collected from multiple data sources and combined. Data were collected on petroleum consumption and pollution emissions while maintaining the order of countries and the sequence of years (from 2006 to 2014, which are relevant to our study).

figure. 3.5 .data integration

3.3 Data Mining

Combine the values associated with the study.With the exception of year and country and maintain sequence to indicate data- Oil consumption - tons of pollution. As in the following figure So that we get a good pattern. Be in three columns - Sequence - Petroleum - Pollution. While maintaining the previous sequence of data. To extract a knowledge-based pattern of petroleum-pollution.

Figure 3.6 . Data Patterns

This is the final form of the modification of the previous data.

Figure3.7 . Dataset (output dataset)

3.4 Implantation

After completion of data mining and data filtering, the spectral algorithm is ready to be implemented. The implementation phase of the algorithm is as follows:

1- Mathematical steps to calculate the value of matrix A (a square matrix is created from the original data). Also at this stage, the matrix of Affinity is calculated, in which we find the values of the distance between the points).Matrix D is also calculated. The diagonal of matrix A is used to find an N.L matrix.

2- Steps of Linear Algebra. >In this phase, the eigenvalue value is calculated. The matrix is also calculated by finding vector v.Apply the k-means algorithm to the U matrix.

3- Details of the implementation phase mathematical steps.

3.4.1 Load Dataset

Data is represented in the figure of a two-dimensional matrix. In our study, the matrix was arranged in**2** columns and in **270** rows, which are the data of pollution emission points and oil consumption $[2 \times 270]$.

Figure3.8 . Dataset

3.4.2 Running the Algorithm.

- ➢*Preprocessing.*
- ➢ *Decomposing.*

➢*Running k-means algorithm.*

To run the algorithm, there are three operations which execute sequentially, as follows:

3.4.2.1 Preprocessing

1) Calculate Affinity matrix:

Find affinity matrix A [270,270] from matrix A [270,2]

Affinity = $1/e$ **dis/** $2\sigma2$

where d is the distance between every element in and others in the same.

Matrix A by the low

\dot{a} **is** $p1, p2 = \sqrt{x1 - x2}$ + $(y1 - y2)$ 2

Where $p_1 = (x_1, y_1)$ & $p_2 = (x_2, y_2)$ The trial and error value were calculated to obtain a satisfactory result.

The value was k=3.

2) Calculate the Row Matrix

Calculate the sum of each row of matrix A in a position equal to the rows and columns of

Matrix **D**.

$$
D[i, j] = \sum D[i, j] \, nj = 1 \, \text{ if } i = j
$$
\n
$$
D[i, j] = \text{Zero} \qquad \text{ if } i \neq j
$$

where $n = 270$ is the matrix degree

All other elements in matrix D will be equal to zero.

3) Calculation to normalize the Laplacian affinity matrix

$$
L = \frac{affinty \{i,j\}}{\sqrt{d\{i,j\} * \sqrt{d\{j,j\}}}}
$$

3) Calculation is to normalize Laplacian affinity matrix

$$
L = \frac{affinty \{i,j\}}{\sqrt{d\{i,j\} * \sqrt{d\{j,j\}}}}
$$

3.4.2.2 Decomposing.

We used linear algebra techniques to find both V and the eigen value **λ (* V = λ * v)**,where v is the eigenvector of N.lap corresponding to λ. We have 270 values of eigenvectors and eigenvale, so for each λ, there is V (λ1. V1, λ2 ... V2,, λ270. V270). From the eigenvectors, we choose $(k + 1) = 2$ similar to k from any group. We select the largest integers according to k which correspond to the same number of the largest eigenvalues because k = 3 equals 4.

We construct a normalization matrix [U] from the Nig matrix obtained [270,2] to find each element of this matrix using the code.

$$
U\{I, J\} = \frac{v(i, j)}{\sqrt{v(i, 1)2 + v(i, 2)2 + v(i, 3)2}}.
$$

3.4.2.3 Running k-means Algorithm.

The K-means algorithm is run with the [U] matrix. Then, the data will be broken down into groups according to spectral clustering.

3.4.2.4 Results Analysis

Data is represented in the two dimensions. All cluster data are colored to distinguish one from the others in order to determine whether these results are acceptable. The results cannot be accepted in one way or another in the parameter change models for the k and σ values, or only one of them. For meaningful results, the properties of each group must be distinguished between the groups that support the end outputs of spectral clustering algorithm implementation.

Table 3.8 Outputs of Implementation S.C.A k=3,4,5,6

3.5 Result and Dissociation

3.5.1 Results of the Spectral Algorithm (*data set of oil and pollution***)**

When the dataset (pollution, oil) was performed using MATLAB language and using the A.S.C. algorithm, we produced the following pattern: a drawing that shows several clusters such that each color is a different color. Each color represents a different classification.

The following figure illustrates this classification**.**

Figure 3.9. Represent Dataset Points in 2D

Application of the algorithm to data related to the production of oil energy and the amount of pollution combines the result of implementing the algorithm and the original data set.

For best results. Therefore, we implemented the program several times with the change K value as the cluster number each time, and then we studied these results for the best.

Result S. C.A with k=3

And through follow-up and observation When the algorithm is executed for the **k = 3** value we get a logical relationship. The higher the consumption of the average of the oil, the lower the average of the pollution. As shown in Figure 3.5,when **k = 3** is executed, the following was obtained. Figure 3.11, average of pollution, average of oil and average standard deviation. Graph 3 shows the number of clusters.

Note. We used standard deviation, which is a measure of the [dispersion](http://www.investopedia.com/terms/d/dispersion.asp) of a set of data from its [mean.](http://www.investopedia.com/terms/m/mean.asp) It is calculated as the square root of the [variance](http://www.investopedia.com/terms/v/variance.asp) by determining the variation between each data point relative to the mean. If the data points are further from the mean, there is higher deviation within the data.

No. Clusters	AVG.Pollution	AVG. Oil	Avg.str
Cluster1	1286.125	5630.6	4129.073
Cluster2	10204.45	7407.96	1772.304
Cluster ₃	17299.76	18375.74	10408.96

Figure 3.14 two Avg. Values (pollution and oil) for S.C.A (k=3)

Result S. C.A with k=4

 The fourth cluster may not give a correlation relation. It gives only a classification. The remainder of the clusters gives a correlation relation and classification (clusters 1,2,3),as shown Figure 3.14.

Figure 3.14 two Avg. Values (pollution and oil) for S.C.A (K=4)

 Figure 3.15 . Representing S.C.A result with 4 Clusters Figure 3.16. Graph result for S.C.A with k=4

Result S. C.A with K= 5

We note that the representation is not integrated. The reason is the fourth CLUSTER, where the percentage of pollution has increased. The increase in pollution may be caused by other factors, as in Figure 3.17.

No. Clusters	AVG.Pollution	AVG Oil	Avg Str
CLUSTER 1	1550	14254	3046.294
CLUSTER 2	846.55	1761.75	1384.536
CLUSTER3	15036.6	12255.51	8572.648
CLUSTER 4	4574.183	5884.38	1341.554
CLUSTER5	3525	25902.21	1822.175

Figure 3.17 two Avg Values (pollution And oil) for S.C.A (K=5)

 Figure 3.18. Representing S.C.A result with 5 Clusters Figure 3.19 Graph result for S.C.A with k=5

Result S. C.A with k= 6

 figure 3.21. Representing S.C.A result with 6 Clusters figure 3.22. Graph result for S.C.A with k=6

3.5.2 Results of K-mean Algorithm (Data set of Oil and Pollution)

K-means will be executed more than once to obtain the best result.

Result k-mean with *K=3*

Through implementation, we obtained good classification only. However, there was no logical correlation relation between them, as in Figure **3.23.**

Figure 3.23. Avg. Values (Pollution and Oil) for k-mean (K=3)

 figure 3.24. Representing k-mean with 3 Clusters figure 3.25. Representing k-mean result with k= 3

Result k-mean with K=4

Through implementation, we obtained good classification and correlation relation, as in Figure **3.26.**

figure 3.24. Representing k-mean with 4 Clusters figure 3.25. Representing k-mean result with k= 4

Result k-mean with k=5

Through implementation, we obtained a good classification. However, there was no logical correlation relation between them, as in Figure **3.26**.

No. clusters	AVG.Pollution	AVG oil	Avg.str
CLUSTER 1	678.3257	1802.369	1192.084
CLUSTER 2	24609.23	22037	2021.371
CLUSTER3	3107.55	24244.42	2962.641
CLUSTER4	1291.49	12655	2484.558
CLUSTER5	7131.5	5669.5	2715.822

Table 3.26 (Pollution &Oil) for k-mean (K=5)

figure 3.27. Representing k-mean with 5 Clusters figure 3.28. Representing k-mean result with k= 5

Result k-mean at k=6

Through implementation, we obtained good classification only, but there was no logical correlation relation between them, as in Figure **8**.

No. clusters	AVG.Pollution	AVG oil	Avgstr
CLUSTER 1	342.7118	970.72	505.1815
CLUSTER 2	1350.39	12965.68	2431.155
CLUSTER3	24609	11037	2021.371
CLUSTER4	7965	6231	2599.006
CLUSTER5	1635	3690	1599.214
CLUSTER6	3107	24244	2962.641

Table 3.29 Avg. Values (pollution and Oil) for k-mean (K=6)

figure 3.27. Representing k-mean with 6 Clusters figure 3.28. Representing k-mean result with k= 6

3.6 Evaluation of Results.

After the algorithm is applied to several attempts, we acquired the best results when **k = 3.** The classification is good and there is also a correlation. It also follows the average standard deviation Table 3.19. We find the lowest value for cluster 3 and cluster 5. It is known that the lower the value of the standard deviation, the better the grouping of data and the better the classification.

figure 3.29 . number of k and stander deviation Figure 3.30 . number of k and stander eviation

When k-mean, k=3

We note that through our implementation of data (pollution and petroleum), the best results were obtained with k-means in many ways:

1 - in terms of the relationship between quantities of emissions and quantities of petroleum consumption when they were $k = 3$.

petroleum consumption, when they were $k = 3$, the relationship is the greater, The consumption of petroleum as pollution increases, as we have noticed in the following figure:

- 1. Shapes: One of the defects of k-means that this algorithm favors circular shapes. We note that the clustering has not been organized.
- 2. Size: We note a clear difference in the size of the cluster.
- 3. The average standard deviation for the value 4057 is greater than the remainder of the cluster. This indicates that the data representation is not strong.
- 4. Density: The density of cluster is different from one cluster to another; some clusters have higher densities and other shave lower densities.

In brief, when it was applied to the data (pollution, petroleum) on the k-means algorithm, the value of $k = 3$ was very logical and the cognitive pattern was good.

The main problem was the classification. The classification was not the best of solutions or attempts because of the lack of this algorithm, as mentioned earlier.

When the k-means **k=6**, in terms of classification and data representation, the best is k = 6 due to the average standard deviation of 2019.76 being Smaller than the remainder of the cluster. Moreover, it decreases the disadvantages of k-means of density and size, as in Figure 3.20 and higher values for the k-means. This makes the results better on this data.

Figure 3.33 Optimal Clusters

4- CONCLUSION

The purpose of this study is to clarify the pros and cons of spectral clustering and to compare results with the real world and apply the truth data and process them before use. We attempted to find a good classification and intelligent style in one of the algorithms and compare the performance of the algorithms. Through our analysis, we see useful results for using the spectral clustering algorithm where determining and arranging sets of any particular statistics or studies in many fields, such as the economy, education, people activities, social studies, criminal fields, finance, industry, health, pollution, environment and oil energy. We also saw their effects on each other. By doing so, we observe through the application of an algorithm, or technique such as regression and correlation that it is difficult to determine the independent or dependent variable, or determine which one is linear or nonlinear. The thesis summarizes the application of spectral and quantification of four variables based on real data from an official source in the 31 countries in the European Union in the seven years from 2006 to 2014. The first study ranked between tons of pollution and oil production or oil energy, according to quantities of harmonization of pollution and oil. The second study was between the quantities of pollution and the level of consumption of clean energy and oil, which was classified as quantities of pollution and the amount of energy consumed.

This algorithm makes it easier for us to classify countries in terms of quantities of pollution over the years, the magnitude of the oil consumption, and the extent to which these countries develop in terms of oil or oil production.

The result was that there was a direct effect of all these variables on each other by dividing these data into four different groups in their characteristics and specifications.

Analysis and study of the application of the S.C. algorithm showed a strong correlation between the four variables, as follows:

Classification of countries in terms of quantities of pollution and production of clean energy or oil energy to seven groups according to the average of both pollution and oil energy; and

The classification of countries in terms of quantities of pollution and consumption of petroleum also into seven groups according to the average of both pollution and consumption of petroleum.

40

The first study is the pollution and oil energy production of the EU countries for 2006-2014.

5. Future Recommendations

While executing the algorithm(spectral and k-means) on a large amount of data, we obtained different data and when comparing around, we found that the spectral clustering was better thanks to the k-means ,because k-means depends on the center (centroid).Therefore, in any future study, the best comparison can be made between the spectra clustering with the Bisecting k-means algorithm for clustering, such as Bisecting k-means, which results in less damaging defects.

We also have difficulty determining the sigma value and k clusters, which is heavily influenced by the data (depending on the resolution).

Note. Bisecting k-means a combination of k-means and hierarchical clustering.

It starts with all objects in a single cluster.

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APPENDIX A

SOURCE CODE FOR spectral cluster algorithm

We used Mat lab version 14 to implement the algorithm consists of the 4 programs

```
1. main program (Jordan_Weiss1) 
2. read data from an external file (Gen_Data1) 
3. Calculate Affinity matrix (Cal_Affinty1) 
Firstly, main program (Jordan_Weiss1) 
clear all; 
close all; 
dim=2 
data = Gen_Data1; 
affinty1 = Cal_Affinty1(data1, dim); 
for i=1: size(affinty1,1) 
D(i,i) = sum(affint y1(i,:));end 
for i=1: size(affinity,1) 
for j=1: size(affinty1,2) 
NL1(i,j) = affinty1(i,j) / (sqrt(D(i,i)) * sqrt(D(j,j)));end 
end 
[eigVector, eigValues1] = eig1(NL1); 
k = 7nEigVec = eigVector (:(size(eigVector,1) -(k-1)): size(eigVector,1)); 
for i=1: size(nEigVec,1) 
n = sqrt (sum (nEigVec(i, :).^2));
U (i, :) = nEigVec(i,:). / n;
end 
[IDX, C, sumb, d] = kmeans(U,3);
```

```
%data = display1(IDX,data,dim); 
if dim==2 
figure, plot3(data (:1), data (:2), data(:,3),'r+'), title('Original Data Points'); grid on;
figure, plot3(data (1,1), data (1,2), data(10,3),'w+'); grid on; 
hold on 54 
for i=1:size(IDX,1) 
if IDX(i,1) == 1plot3(data(i,1),data(i,2),data(i,3),'y+'); 
data(i,4)=1elseif IDX(i,1) == 2plot3(data(i,1),data(i,2),data(i,3),'g+'); 
data(i,4)=2elseif IDX(i,1) == 3plot3(data(i,1),data(i,2),data(i,3),'r+'); 
data(i,4)=3elseif IDX(i,1) == 4plot3(data(i,1),data(i,2),data(i,3),'b+'); 
data(i,4)=4elseif IDX(i,1) == 5plot3(data(i,1),data(i,2),data(i,3),'m+'); 
data(i,4)=5
elseif IDX(i,1) == 6plot3(data(i,1),data(i,2),data(i,3),'k+'); 
data(i,4)=6end 
end 
elseif dim==2 
figure,plot(data(:,1), data(:,2),'b+'), title('Original Data Points'); grid on;shg 
figure
```

```
hold on 
for i=1:size(IDX,1) 
if (IDX(i,1) == 1)plot(data(i,1),data(i,2),'r+'); 
data(i,3)=1elseif IDX(i,1) == 2plot(data(i,1),data(i,2),'k+'); 
data(i,3)=2elseif IDX(i,1) == 3plot(data(i,1),data(i,2),'m+'); 
data(i,3)=3elseif IDX(i,1) == 4plot(data(i,1),data(i,2),'r+'); 
data(i,3)=4elseif IDX(i,1) == 5plot(data(i,1),data(i,2),'m+'); 
data(i,3)=5 
else 
plot(data(i,1),data(i,2),'y+'); 
data(i,3)=6 55 
end 
end 
end 
xlswrite('D:\CLUTUR AND CRIM\NEW.xlsx',data,'sheet6');
hold off; 
grid on; 
tt=1; 
secondly read data from an external file (Gen_data) 
function [data] = GenerateData()
```

```
45
```

```
data=xlsread('E:\kemerburgaz\thises\data_try\input_data.xlsx','sheet1','a:c'); 
end 
thirdly calculate Affinity matrix (Cal_Affinity) 
function [affinity] = Cal_Affinity(data,di) 
sigma =0.572 
if di==3 
for i=1:size(data,1) 
for j=1:size(data,1) 
for K=1:size(data,1) 
dist = sqrt((data(i,1) - data(j,1) \binom{2}{} + (data(i,2) - data(j,2))^2+ (data(i,3)- data(j,3) \binom{2}{} );
affinity(i,j) = exp(-dist/(2*sigma^2));else 
for i=1:size(data,1) 
for j=1:size(data,1) 
dist = sqrt((data(i,1) - data(j,1) \bigwedge^2 + (data(i,2) - data(j,2)\bigwedge^2);
affinity(i,j) = exp(-dist/(2*sigma^{-2}));end
```
APPENDIX B

The table below consists of data representing the first study conducted using algorithms and their results.

