# Gebze Technical University Computer Engineering CSE 454 Data Mining Fall 2020 Project Assignments

**Project Report** 

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# Running Clustering Algorithms and Visualizing Data on Fifa 2021 Dataset

# Summary

In this study, it provides information about the implementation of DBScan, Hierarchical and Mean Shift Clustering algorithms on Fifa 2021 data. The aim of the study is to classify the approximately 17000 players in the Fifa 2021 game according to their characteristics in 80 different categories and to identify players with anormal data, if any. In this study, it is also planned to investigate whether the same players in the specified clustering algorithms will be included in the same clusters or in different clusters.

#### Introduction

FIFA 2021 is a football simulation game developed by EA Sports company. Game data includes detailed information about the players and teams. There is a data set consisting of 80 columns for each player. The main ones are the salary, the transfer fee, which foot, etc. are information. The data set includes information of 17 126 players from 164 countries, 651 teams. It has sufficient content for data set classification algorithms and we need to do some preliminary studies before starting operations on the data set.

### PREPROCESS STEP

These clustering algorithms are algorithms that work on numerical data. Therefore, non-numerical data in the data set has been turned into numerical data. It has been decided which data will be effective in clustering the players. Most of these data are variables of the type of continuous variable and these data are turned into categorical variables because they increase the granules. Variables such as M, K used in the data actually mean millions and thousands, and the places where these values are corrected by multiplying millions and thousands. To use the data more efficiently, the standard scaling method was used to pull the data into a more meaningful range. Thus, data such as wage and transfer price (value) were included in a meaningful scale.

#### **MODELS**

# A) DBScan

DBSCAN algorithm was presented by Ester, Kriegel, Sander and Xu at the conference KDD'96. This algorithm calculates the distance of objects from their neighbors and performs clustering by grouping areas with more objects than a predetermined threshold in a given region. The DBSCAN algorithm has brought many new terms and approaches to data mining.

Since DBScan is a density base algorithm, it is affected by the proximity of data to each other, so it can discover randomly shaped clusters. It can find clusters that are completely surrounded by different clusters. It is sensitive in detecting outlier data. Since it is a density-based algorithm, it may not be able to cluster if the data is very sparse. Sampling affects density measures.

# B) Hierarchical Clustering

Hierarchical clustering is a clustering algorithm as the name suggests. It has two different variations as Agglomerative (from part to whole) and Divisive (from whole to part). In this work, the Agglomerative version was studied.

In this variation, first all the data is made into a set, that is, if there are N elements, N sets are formed. Then, clusters that are close to each other in distance merge to form a new cluster. This situation continues until the system is stable or the number of clusters the programmer enters.

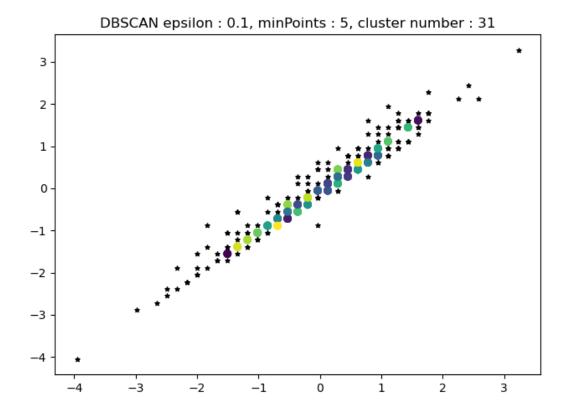
# C) Mean Shift Clustering

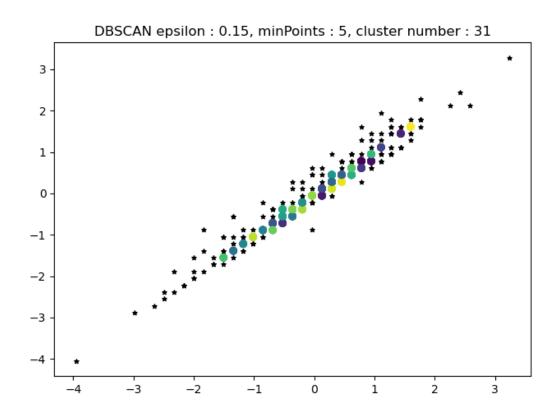
Meanshift falls under a category of clustering algorithms, unlike Unsupervised learning, which recursively allocates data points to clusters by shifting points towards mode (mode is the highest density of data points in the region in the context of Meanshift). Hence, it is also known as the Mod search algorithm. The average shift algorithm has applications in image processing and computer vision.

Given a set of data points, the algorithm recursively assigns each data point toward the nearest cluster center, and the direction to the closest cluster center point is determined by where most of the nearby points are. Therefore, each iteration, each data point will be closer to where the most points are located, which will reach the cluster center. When the algorithm stops, each point is assigned to a cluster.

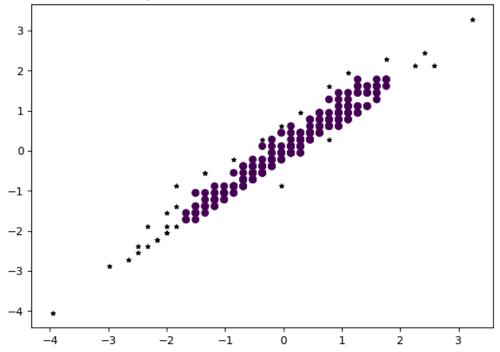
# **MODELS RESULTS**

# A) DBScan

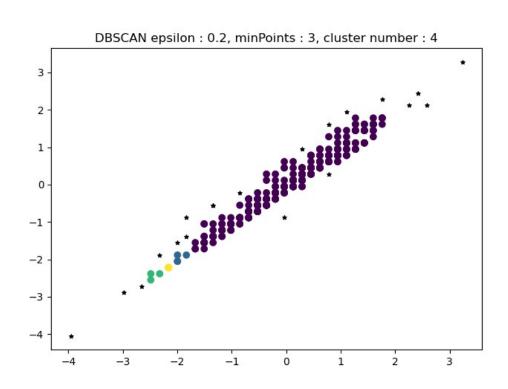




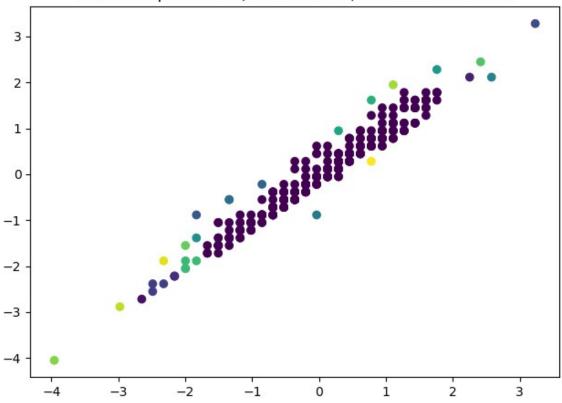




Now the minpoint variable will be kept constant at 5 and the epsilon value will be increased. As I value the Epsilon value relatively low, separate clusters were formed as seen in the plot. In the dataset I use, the data are collected on the middle side of the plot, so the clusters are on that side. Noise values have emerged on the right, where there is less data. And the areas where objects are relatively far from each other are these areas. I would like to point out that at 0.1 and 0.15 the same number of clusters is formed but different points are included in the cluster.

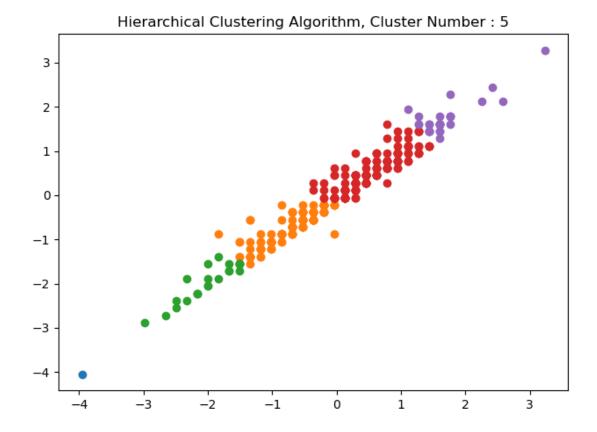


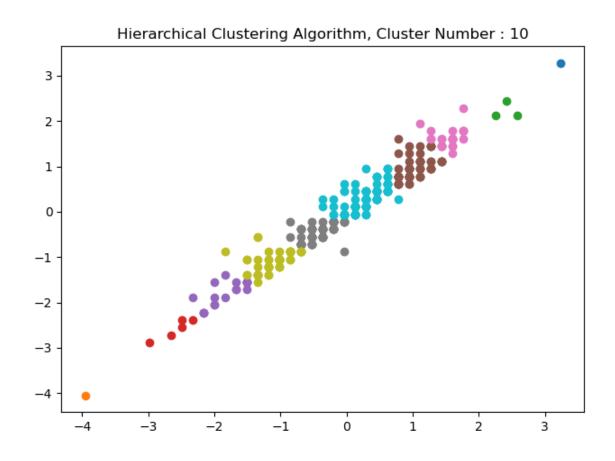
DBSCAN epsilon: 0.2, minPoints: 1, cluster number: 23



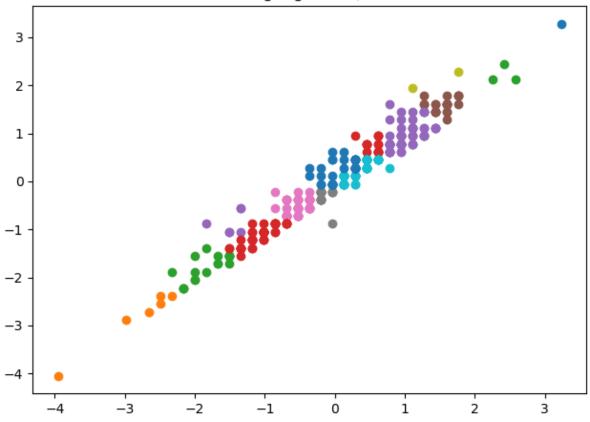
When I decreased the minpoints, I expect the number of clusters increased. Many clusters split. That's why it increased. Because now, theoretically, it is not provided that objects that have more distances to each other are in the same set. Likewise, it was observed that new clusters were formed on the right side where data was relatively less (distance between objects is relatively greater). Where objects are relatively close to each other, clusters are split, the number of clusters has increased.

# **B)** Hierarchical Clustering





Hierarchical Clustering Algorithm, Cluster Number: 15

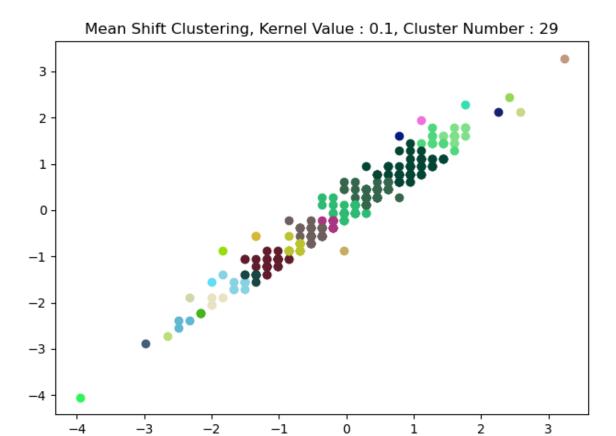


Single linkage is used in this algorithm. Single linkage is also sometimes called 'friends of friends' linkage method. The single linkage method always picks the minimum distance when it updates. Therefore, two points might be considered close under the single linkage scheme if they can be connected by a chain of points with small distances between any two consecutive points down the chain. The distance between the two endpoints of the chain is only as big as the longest link along the chain. The number of links along the chain does not matter. Therefore, it is expected that more clusters will be observed as the number of N increases. Players who play too well or those who play too bad are in a cluster.

#### C) **Mean Shift Clustering**

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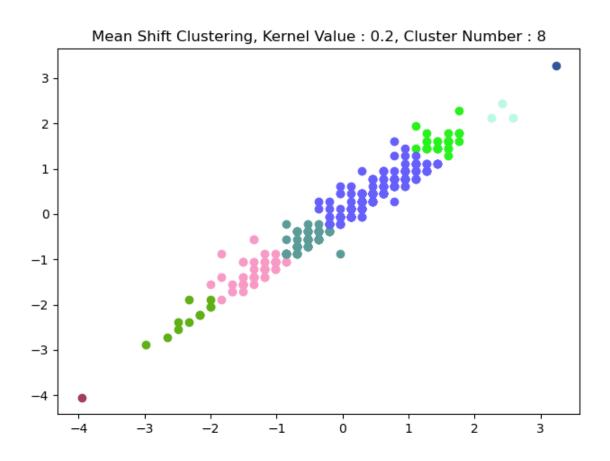
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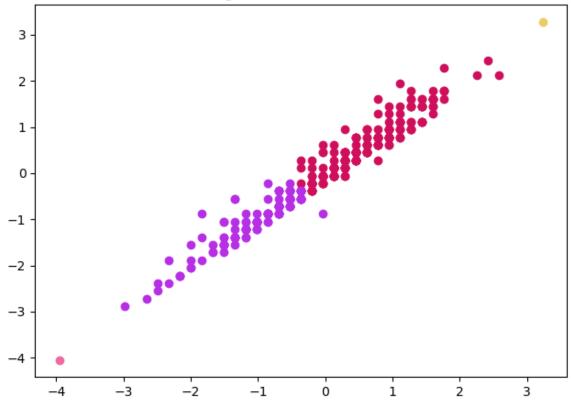
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Mean Shift Clustering, Kernel Value: 0.3, Cluster Number: 4



This algorithm starts from random data. And this data becomes the center of the kernel. The kernel is shifted as kernel value. This is the summary of the implementation of the algorithm. Here, it is observed that as you increase this value, the number of clusters decreases. Considering that the kernel has increased its hiccup, we can connect more easily. In this implementation I calculated according to the gauss distance.

### **GENERAL RESULT**

Thanks to the clustering algorithms applied on Fifa 2021 data, the data could be clustered. As a result of the applied algorithms, different results were obtained and similar results were obtained. The results were observed as expected and consistent with those observed in real life.

As a result of the algorithms applied, the data were classified successfully and consistently. As a result of the classification, samples that all algorithms gather in the same class were observed and it was concluded that this was a metric showing the accuracy of the study.

## **VISUALIZATION EXAMPLES**

Images of the images are available in the result\_pictures folder. It can be examined in detail from that folder.

