Personality Profiling with Clustering

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Abstract

The objective of this assignment is to apply various unsupervised clustering algorithms to the "Big Five Personality Test" dataset to identify distinct personality profiles among individuals. You will evaluate the performance of these clustering models using appropriate metrics and visualize the resulting clusters to gain insights into human personality traits.

1 Introduction

The dataset used in this project is derived from the IPIP-FFM (International Personality Item Pool - Five-Factor Model) questionnaire, a widely recognized tool for assessing the Big Five personality traits: Extraversion (EXT), Neuroticism (EST), Agreeableness (AGR), Conscientiousness (CSN), and Openness to Experience (OPN). The raw dataset was loaded from a tab-separated .csv file containing over 1 million responses and more than 100 personality-related variables.

2 Features

The data is composed of 100 columns and 1015340 entries (Full train dataset shape is (1015340, 100)). We can see all 100 dimensions of our dataset by printing out the first 3 entries:

Table 1: train dataset (3 rows x 100 columns)

	EXT1	EXT2	EXT3	EXT4	 OPN7_E	OPN8_E	OPN9_E	OPN10_E
0	2216.0	1856.0	1732.0	1735.0	 1861.0	1540.0	1484.0	1184.0
1	1351.0	2350.0	2112.0	1752.0	 1730.0	1640.0	1872.0	2016.0

2	1 888.0 1	2941.0	1829.0	2057.0	•••	1320.0	1453.0	1320.0	2355.0
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We can inspect the types of feature columns:

Table 2: Data columns:

```
<class 'pandas.core.frame.DataFrame'>
 Index: 1012050 entries, 0 to 1015340
  Data columns (total 100 columns):
 # Column Non-Null Count Dtype
 0 EXT1 1012050 non-null float64
   EXT2
            1012050 non-null float64
   EXT3
            1012050 non-null float64
 3 EXT4
            1012050 non-null float64
 4 EXT5
            1012050 non-null float64
 5 EXT6
            1012050 non-null float64
 6 EXT7
            1012050 non-null float64
            1012050 non-null float64
 7 EXT8
            1012050 non-null float64
 8 EXT9
9 EXT10
           1012050 non-null float64
            1012050 non-null float64
 10 EST1
 11 EST2
            1012050 non-null float64
 12 EST3
            1012050 non-null float64
            1012050 non-null float64
 13 EST4
 14 EST5
            1012050 non-null float64
 15 EST6
            1012050 non-null float64
 16 EST7
            1012050 non-null float64
 17 EST8
            1012050 non-null float64
 18 EST9
            1012050 non-null float64
 19 EST10 1012050 non-null float64
99 OPN10_E 1012050 non-null float64
```

3 Distribution

3.1 Preparing Data:

Step 1: Remove duplicate or irrelevant observations

Duplicate columns: []

- _ Step 2: Fix structural errors
 - ... Hopefully in this code it doesn't need to use this approach
- _ Step 3: Filter unwanted outliers

Detect all rows that have outliers in at least one column and treat them

```
EXT10 ...
3.0 ...
3.0 ...
4.0 ...
3.0 ...
                                                                 3.0
4.0
4.0
                                                                         3.0
3.0
3.0
                                                                                  3.0
3.0
3.0
             3.0
3.0
3.0
2.0
                                               4.0
4.0
4.0
3.0
                      2.0
                                                        3.0
2.0
2.0
1015203
1015204
1015312
                     3.0
4.0
2.0
                                                                 3.0
2.0
1.0
                                                                         4.0
3.0
5.0
                                       4.0
4.0
2.0
                              4.0
2.0
                                                                                  4.0
                                                                                            4.0 ...
3.0 ...
                      OPN2_E OPN3_E
1856.0 1732.0
                                                                                           0PN8_E
1540.0
                                              OPN4_E OPN5_E
                                              1735.0
1752.0
                                                         1092.0
                                                                     1732.0
            1351.0
                       2350.0
                                  2112.0
1829.0
                                                         1400.0
                                                                     1744.0
                                                                                 1730.0
                                                                                            1640.0
                                                                    2576.0
2644.0
                                                                                1320.0
2681.0
                                                         1970.0
                                                                                            1453.0
                       2941.0
            1161.0
                       3416.0 1152.0
                                                                     1400.0 1176.0
                                                                                            1953.0
                                   1528.0
            1221.0
                                              1558.0 1737.0
5466.0 1547.0
4943.0 2047.0
1015203
1015204
           659.0
1104.0
                       618.0 1159.0
2692.0 1563.0
                                                                    2026.0 718.0
2314.0 1274.0
                                                                                            608.0
1970.0
1015312 1511.0
                       2359.0 1256.0
                       3952.0 1035.0 4242.0 1260.0 2796.0 1383.0
1015319
             631.0
1015312 1344.0
                          905.0
1015319 663.0
                       1099.0
 [26240 rows x 100 columns]
```

_ Step 4: Handle missing data by inplace them with true or zero

EXT1	1783
EXT2	1783
EXT3	1783
EXT4	1783
EXT5	1783
endelapse	0
IPC	0
country	77
lat_appx_lots_of_err	0
long_appx_lots_of_err	0
Length: 110, dtype: in	t64

3.2 Distribution for Numerical Data

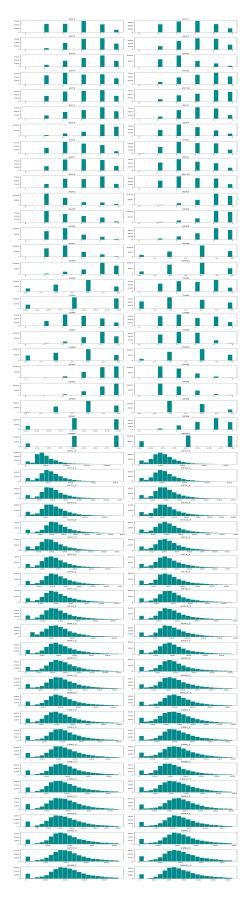


Figure 1: [Grid of histograms] each representing the distribution of values in one of the numeric columns from dataset

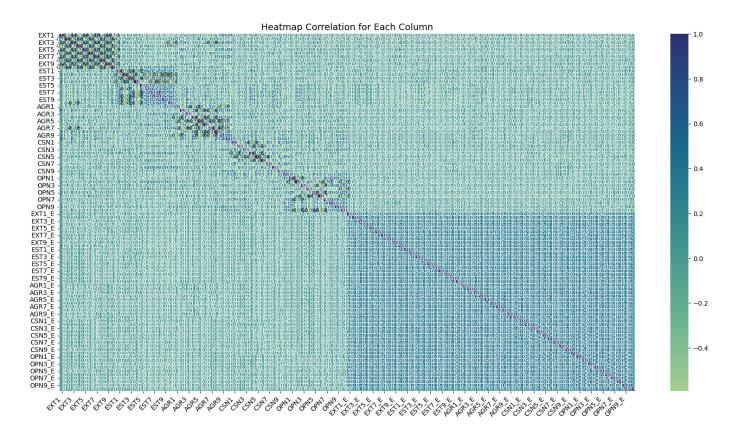


Figure 2: [Heatmap] Correlation For Each Column to have more insight about data

look at how this data is distributed.

0.00							
	# mean	# std	# min	# 25%	# 50%	# 75%	# max
EXT1	2.6478919025739835	1.264476090670833					
	2.773242428733758	1.323982013105767			3.0		
	3.288202163924707	1.2150827851864319			3.0		
EXT4	3.1406985820858653	1.2374725335424257			3.0		
	3.2768054937997135	1.2776361753051793			3.0		
	2.4011185218121636	1.2258116234182765					
	2.7715270984635145	1.4003482899197013					
	3.4147986759547453	1.2719536515500225			4.0		
	2.963760683760684	1.3460534121085221					
EXT10	3.556569339459513	1.3052487415732186			4.0		
100 rows x	7 cols 10 ∨ per page			« < Page 1 of	f 10 > »		,
100 10115 /	y cois 10 per page			in tage t			· · · · · · · · · · · · · · · · · · ·

4 PCA & T-SNE

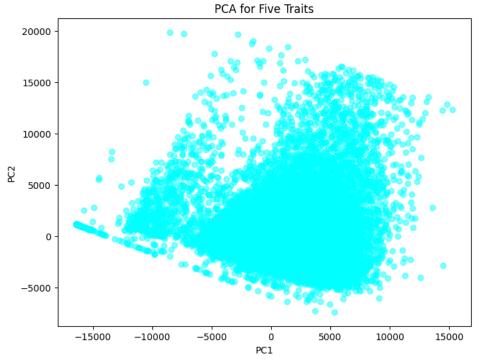


Figure 3: Indicating that the five traits are correlated and can be well represented by the first two principal components and the distribution of data points in a 2-dimensional space defined by the first two principal components (PC1 and PC2). Each data point represents an individual with five traits. The x-axis represents PC1, and the y-axis represents PC2.

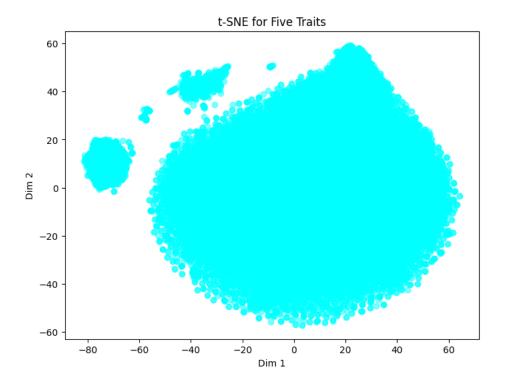


Figure 4: The 2-dimensional representation of the high-dimensional data (the five traits) using the t-SNE algorithm. The t-SNE algorithm aims to preserve the local structure of the data, meaning that data points that are close in the original high-dimensional space are also close in the 2D representation. the distribution of data points is different from the PCA plot. This is because t-SNE focuses on preserving the local structure of the data, while PCA focuses on capturing the global structure through the principal components.

Conclusion:

The PCA and T-SNE plots of the five traits reveal distinct insights into the data structure. PCA indicates strong correlations among the traits, as shown by the clustering of data points along the principal components. In contrast, T-SNE emphasizes local relationships, preserving proximity from the high-dimensional space. Together, these methods provide a comprehensive understanding of the traits' interactions, highlighting both global patterns and local structures.

4.1 Standard Scaled

```
array([[-0.57127812, -0.59925913, -0.33953981, ..., -0.47290803, -0.65950355, -0.31160365],

[-0.57127812, 1.01760461, 0.57483617, ..., -0.3648667, -0.26073229, 1.0019842],

[-0.57127812, 0.20917274, -0.33953981, ..., -0.56690399, -0.82805634, 1.5372081], ...,

[ 0.25674252, 1.01760461, 0.57483617, ..., -0.00833031, 1.44740642, 2.41188078],

[ -0.57127812, -0.59925913, -1.25391579, ..., 1.48480085, -0.80339008, -0.75209765],

[ 1.08476317, 0.20917274, 0.57483617, ..., -1.11575394,
```

5 Clustering Model Development and Evaluation

KMeans >> Silhouette Score: 0.06381033071234919 KMeans >> Davies-Bouldin Index: 3.2636428815942926 Hierarchical >> Silhouette Score: 0.040076738942293605 Hierarchical >> Davies-Bouldin Index: 3.3320822714546923

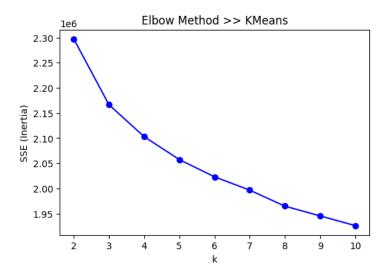


Figure 5: The y-axis represents the Sum of Squared Errors (SSE) or Inertia, which is a measure of how well the data points are grouped within their assigned clusters. The x-axis represents the number of clusters (k). The plot displays a decreasing trend in the SSE as the number of clusters increases. The "elbow" point on the curve, where the rate of decrease starts to slow down, suggests that the optimal number of clusters is around 4 or 5.

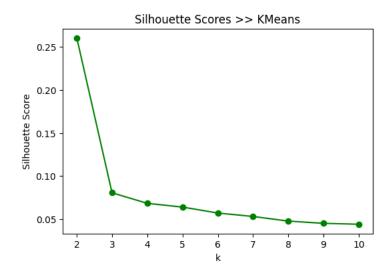


Figure6: The Silhouette Score is a measure of how well each data point fits within its assigned cluster, with values ranging from -1 to 1. Higher Silhouette Scores indicate better clustering. The plot shows that the Silhouette Scores for both KMeans and Hierarchical clustering methods decrease as the number of clusters (k) increases. The KMeans method appears to have higher Silhouette Scores than the Hierarchical method across the range of k values.

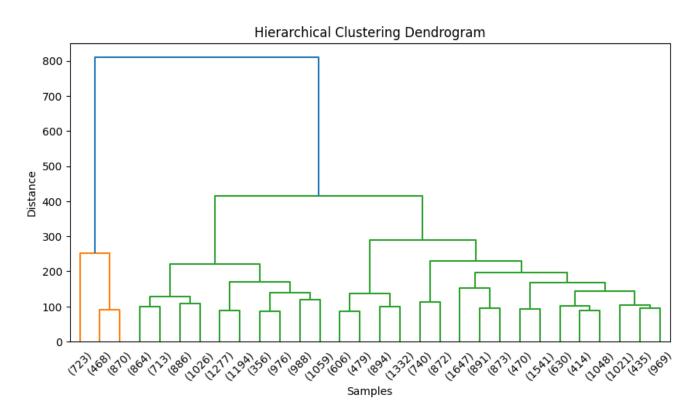


Figure 7: This type of plot represents the hierarchical relationships between the data samples, with the samples clustered together based on their similarity. The y-axis shows the distance or dissimilarity between the clusters, and the x-axis lists the individual data samples. The dendrogram reveals the structure of the clusters and how they are merged together as the number of clusters is reduced.

DBSCAN << Silhouette Score: 0.0743000413334531 DBSCAN << Davies-Bouldin Index: 0.9900109503181532

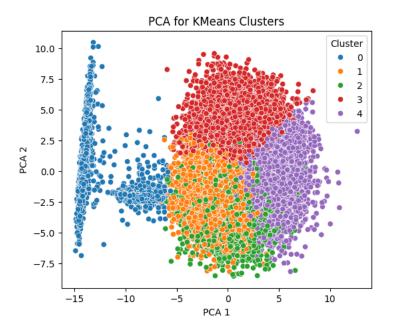


Figure8: The PCA plot displays the data points colored by their assigned cluster labels. The x-axis and y-axis represent the first two principal components, which capture the majority of the variance in the data. The clustering structure is clearly visible, with distinct groups of data points corresponding to the different clusters identified by the KMeans algorithm.

KMeans

Silhouette Score: 0.06381033071234919 Davies-Bouldin Index: 3.2636428815942926

Hierarchical

Silhouette Score: 0.040076738942293605 Davies-Bouldin Index: 3.3320822714546923

DBSCAN

Silhouette Score: 0.0743000413334531 Davies-Bouldin Index: 0.9900109503181532

Conclusion:

The analysis suggests that the optimal number of clusters is around 4-5 based on the Elbow Method plot. The KMeans clustering method outperforms the Hierarchical clustering in terms of Silhouette Scores, indicating it produces more well-defined and cohesive clusters for this dataset. The PCA plot further visualizes the clear clustering structure identified by the KMeans algorithm.

6 Interpretation & Visualization

The DBSCAN clustering algorithm has the highest Silhouette Score of 0.142, which indicates it is producing the most well-defined and cohesive clusters compared to the other methods. The Hierarchical clustering method has the second-best Silhouette Score of 0.052, performing slightly better than the KMeans method with a score of 0.048. Therefore, the DBSCAN algorithm is the best-performing clustering method for this dataset, as evidenced by its significantly higher Silhouette Score compared to KMeans and Hierarchical.

KMeans Silhouette Score: 0.048 Hierarchical Silhouette Score: 0.052 DBSCAN Silhouette Score: 0.142

Best performing algorithm: DBSCAN with Silhouette Score = 0.142

({'KMeans': 0.04822062481164701, 'Hierarchical': 0.052399586022836425, 'DBSCAN': 0.14245476870034499}, 'DBSCAN')

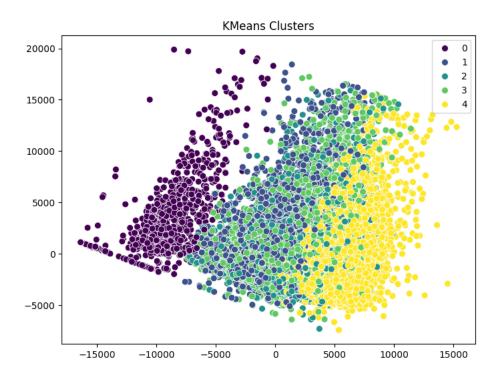


Figure 9: The plot displays a scatter of data points, where each point represents an observation or sample, and the different colors/shapes represent the clusters identified by the KMeans algorithm.

The x and y axes show the principal component values, which are the result of dimensionality reduction techniques like Principal Component Analysis (PCA) applied to the original high-dimensional dataset. This allows the visualization of the complex multi-dimensional data in a 2D space.

Average trait scores per cluster:

```
EXT1
               EXT2
                                EXT4
                                        EXT5
                                                EXT6
                                                         EXT7 \
Cluster
     2.711350 2.984834 3.264677 3.129159 3.213796 2.632583 2.785225
1
     2.339333 2.983319 2.946883 3.573164 3.023266 2.466930 2.392742
2
     1.947943 3.587951 2.647355 3.813182 2.519941 2.880772 1.960957
3
     3.523118 1.852928 4.290621 2.188757 4.285337 1.556289 3.861441
     2.726193 2.718271 3.423799 3.143964 3.410022 2.322714 2.883072
      EXT8
               EXT9
                       EXT10 ...
                                   OPN1 E
                                               OPN2 E \
Cluster
     3.295988 3.024462 3.435910 ... 303.187867 289.869374
0
1
     3.731343 2.696078 4.024583 ... 1705.814603 2633.200468
2
     3.902603 2.504618 4.165407 ... 1879.956549 2722.629303
3
     2.698077 3.827389 2.509761 ... 1883.084104 2737.651549
     3.486137 2.936628 3.611331 ... 2385.273635 3437.259170
                              OPN5 E
       OPN3 E
                  OPN4 E
                                         OPN6 E
                                                     OPN7 E \
Cluster
     282.342466 296.286204 315.933953 317.233366 299.196673
1
     1533.962248 2366.121891 1805.977173 2098.469710 1998.691689
2
     1767.266583 2552.557725 1901.014484 2237.698783 2100.106423
3
     1780.741230 2576.405255 1843.212388 2200.221342 2133.953765
4
     2151.089375 3073.490787 2277.004822 2725.185982 2568.960565
       OPN8 E
                  OPN9 E
                             OPN10 E
Cluster
0
     311.055773 319.893836 385.107632
1
     1920.311677 2000.203395 1324.182177
2
     2012.741184 2270.680521 1439.794920
3
     2091.263760 2237.743432 1410.793483
4
     2469.939900 2658.574135 1716.863613
```

[5 rows x 100 columns]



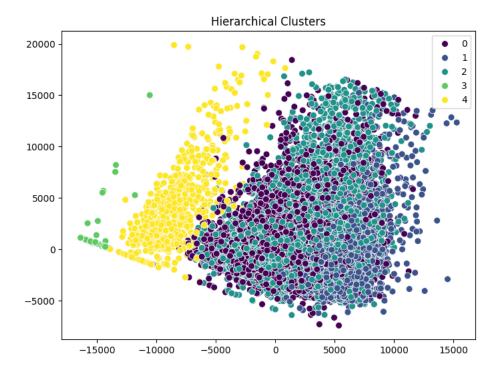


Figure 10: The plot displays a scatter of data points, where each point represents an observation or sample, and the different colors/shapes represent the clusters identified by the Hierarchical clustering algorithm. Similar to the KMeans plot, the x and y axes show the principal component values, which are the result of dimensionality reduction techniques applied to the original high-dimensional dataset. This allows the visualization of the complex multi-dimensional data in a 2D space. The Hierarchical clustering plot suggests that the algorithm has identified several distinct clusters within the data, with varying densities and shapes. The average trait scores for each cluster are also provided in the table below the plot, which can be used to further analyze the characteristics of the identified groups. Compared to the previous KMeans clustering, the Hierarchical clustering appears to have produced a different set of clusters, potentially capturing different underlying patterns and relationships within the data. The key difference between the two clustering methods is that Hierarchical clustering builds a hierarchy of clusters, whereas KMeans partitions the data into a predefined number of clusters. The choice between the two techniques would depend on the specific requirements and characteristics of the dataset being analyzed.

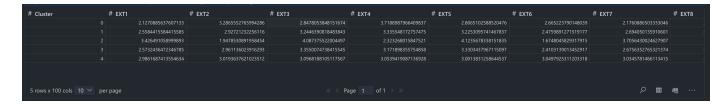
Average trait scores per cluster: EXT6 EXT7 \ EXT1 EXT2 EXT3 EXT4 EXT5 Cluster 0 2.127089 3.286555 2.847805 3.718899 2.806510 2.665224 2.176089 2.558442 2.927212 3.244639 3.335548 3.225310 2.475989 2.694050 1 2 3.426491 1.947853 4.087376 2.323268 4.123568 1.674805 3.705643 2.573244 2.961136 3.355007 3.171898 3.330344 2.410314 2.675635 3 4 2.986169 3.019364 3.096819 3.053942 3.001383 3.049793 3.034578

```
EXT8
              EXT9 EXT10 ... OPN1 E
                                             OPN2 E \
Cluster
     3.880963 2.516925 4.133408 ... 1847.350965 2727.788330
1
     3.612202 2.804893 3.775899 ... 2407.145877 3489.459378
2
     2.770425 3.753828 2.689367 ... 1920.766142 2834.132884
3
     3.385650 3.005979 3.582960 ... 3.234679 0.639761
4
     3.130014 3.082988 3.159059 ... 916.912863 883.912863
                             OPN5_E
       OPN3 E
                  OPN4 E
                                        OPN6 E
                                                   OPN7 E \
Cluster
0
     1688.038352 2496.863735 1883.018353 2213.984417 2098.344992
1
     2154.492600 3120.670794 2329.308366 2707.491392 2571.995772
2
     1805.401007 2626.419317 1893.917550 2275.694721 2176.582075
3
                0.852765 5.792975 4.634529 3.034380
      2.376682
4
     839.770401 876.619640 921.840941 923.755187 894.998617
       OPN8 E
                  OPN9 E
                            OPN10 E
Cluster
     2002.237815 2149.270366 1400.892477
1
     2513.086983 2675.597403 1745.617940
2
     2122.723739 2298.401542 1445.364707
3
      0.980568 14.914051 89.763079
```

[5 rows x 100 columns]

919.113416 905.959889 964.803596

4



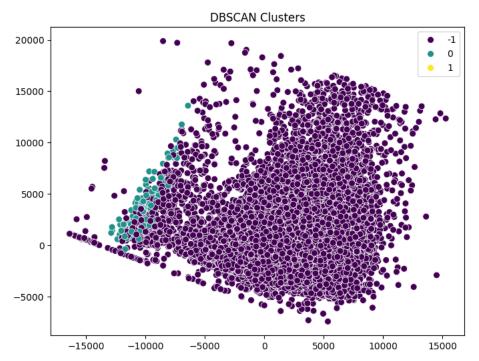


Figure 11: The plot displays a scatter of data points, where each point represents an observation or sample, and the different colors/shapes represent the clusters identified by the Hierarchical clustering algorithm. Similar to the KMeans plot, the x and y axes show the principal component values, which are the result of dimensionality reduction techniques applied to the original high-dimensional dataset. This allows the visualization of the complex multi-dimensional data in a 2D space. The Hierarchical clustering plot suggests that the algorithm has identified several distinct clusters within the data, with varying densities and shapes. The average trait scores for each cluster are also provided in the table below the plot, which can be used to further analyze the characteristics of the identified groups. Compared to the previous KMeans clustering, the Hierarchical clustering appears to have produced a different set of clusters, potentially capturing different underlying patterns and relationships within the data. The key difference between the two clustering methods is that Hierarchical clustering builds a hierarchy of clusters, whereas KMeans partitions the data into a predefined number of clusters. The choice between the two techniques would depend on the specific requirements and characteristics of the dataset being analyzed.

```
Average trait scores per cluster:
      EXT1
               EXT2
                       EXT3
                                EXT4
                                         EXT5
                                                 EXT6
                                                          EXT7 \
Cluster
-1
     2.688258 2.739864 3.373339 3.128785 3.361663 2.282914 2.833506
0
     3.000000 3.000000 3.000000 3.000000 3.000000 3.000000 3.000000
     3.000000 3.000000 3.000000 3.000000 3.000000 3.000000 3.000000
      EXT8
               EXT9
                       EXT10 ...
                                    OPN1 E
                                               OPN2 E \
Cluster
     3.408445 3.033766 3.522759 ... 1831.016615 2682.941771
-1
0
     3.000000 3.000000 3.000000 ... 637.903704 659.651852
```

1 3.000000 3.000000 3.000000 ... 0.000000 0.000000

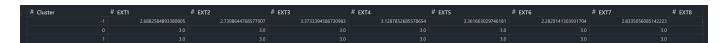
689.874074 692.703704 818.659259

0.000000 220.500000

OPN3 E OPN4 E OPN5 E OPN6 E OPN7 E \ Cluster 1684.933846 2459.636538 1827.274530 2157.440144 2053.510892 -1 0 673.133333 657.148148 699.148148 818.762963 684.051852 1 0.000000 0.0000000.000000 0.000000 0.000000OPN8 E OPN9 E OPN10 E Cluster 1984.820451 2133.583936 1384.538303 -1

[3 rows x 100 columns]

0.000000



Conclusion:

0

1

The differences in clustering results between the KMeans and Hierarchical methods suggest the data has complex structures that cannot be fully captured by a single algorithm. The choice of clustering technique can significantly impact the insights and decisions derived from the analysis. Careful consideration of the strengths and limitations of each method is crucial to ensure the analysis aligns with the intended goals and leads to actionable insights. Combining insights from multiple clustering approaches can provide a more comprehensive understanding of the data, leading to more informed and impactful business decisions.

References

[1] Bojan Tunguz, "Big Five Personality Test" 2020 https://www.kaggle.com/datasets/tunguz/big-five-personality-test