

Data Mining Project

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Comparative Analysis of K-medoids and Hierarchical Clustering for House Pricing Dataset

In this project, we explore two popular clustering algorithms, K-medoids and hierarchical clustering, to analyze a house pricing dataset. Clustering is a fundamental unsupervised learning technique used to group similar data points together based on their characteristics. The aim of this project is to compare the performance of these two clustering algorithms in segmenting houses into distinct price categories.

Dataset Description: The house pricing dataset used in this project contains various features related to house characteristics such as area, number of bedrooms, city, level, etc., along with the corresponding prices.

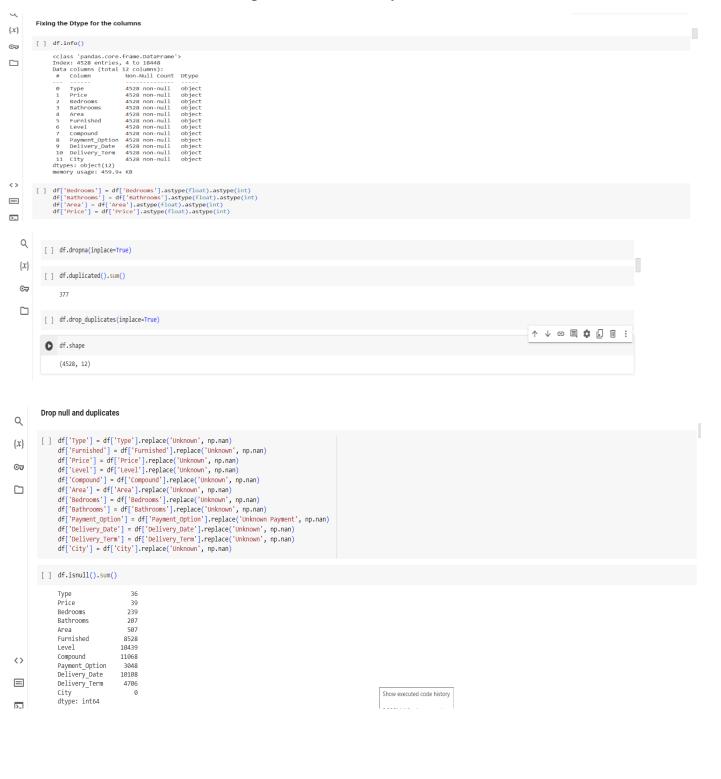
Methodology:

1. Download data "Egypt House Pricing"



Data Preprocessing:

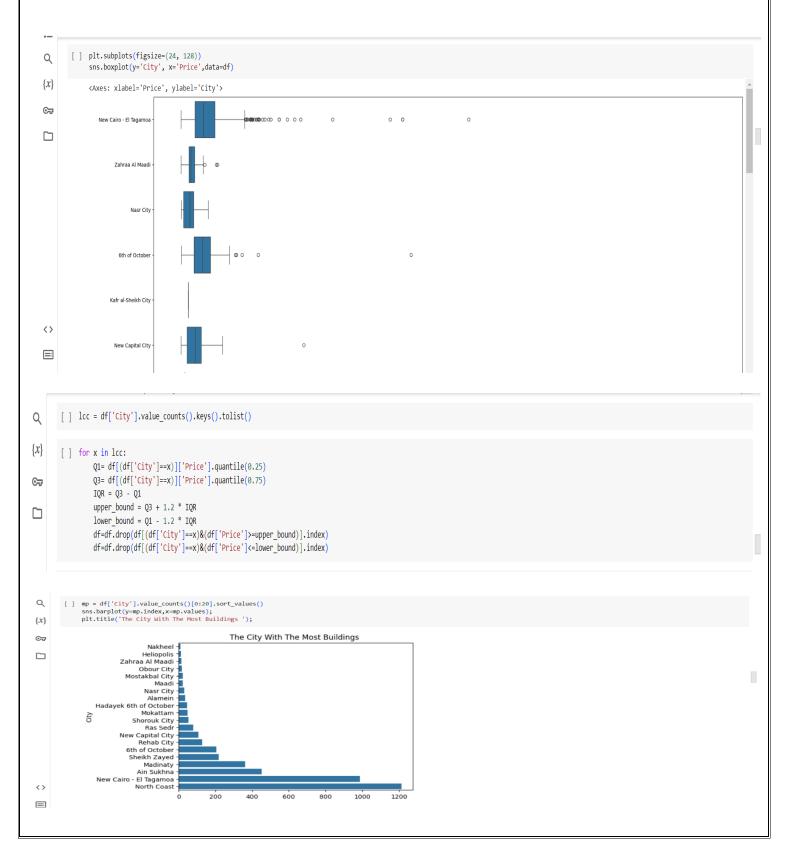
We removed rows with missing values and duplicate entries to ensure data cleanliness and integrity. This step involved checking for missing values in each column and dropping rows with null values. Additionally, duplicate rows were identified and eliminated to prevent redundancy in the dataset.

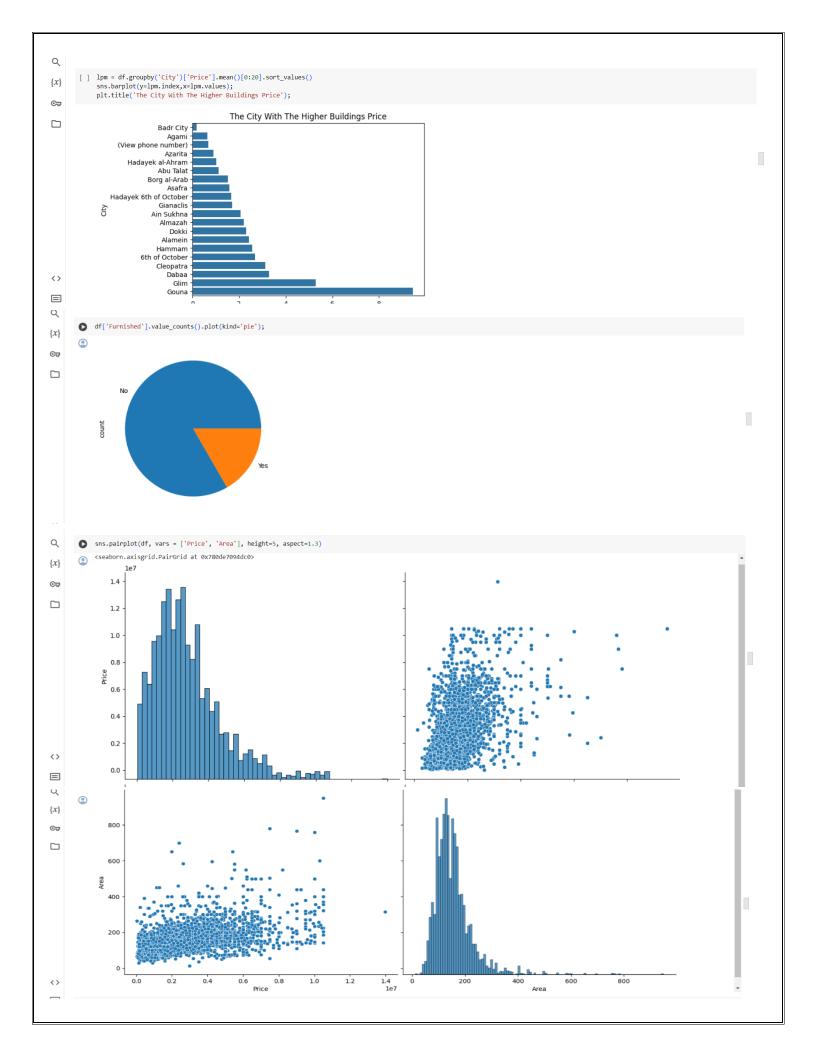


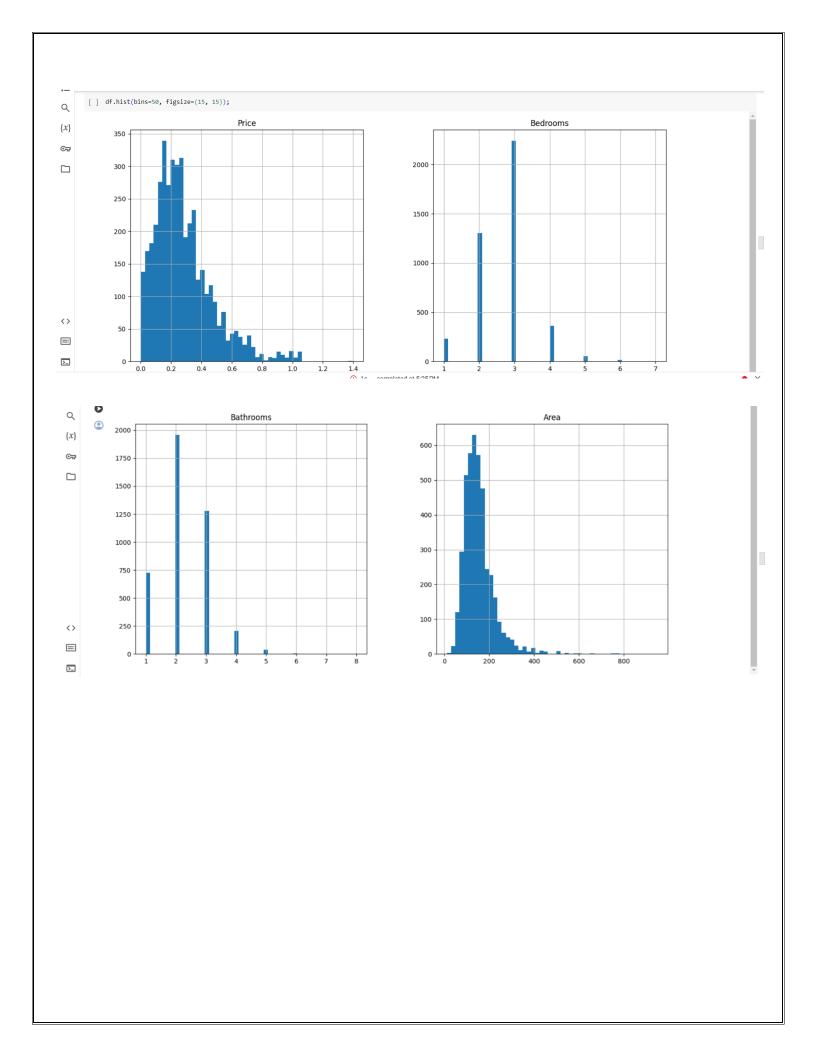


Visualization of Preprocessed Data:

- After preprocessing, we visualized the preprocessed data to gain insights into the distribution and relationships between features.
- Visualization techniques such as scatter plots, histograms, or pair plots were employed to explore the data's characteristics and identify any patterns or outliers.





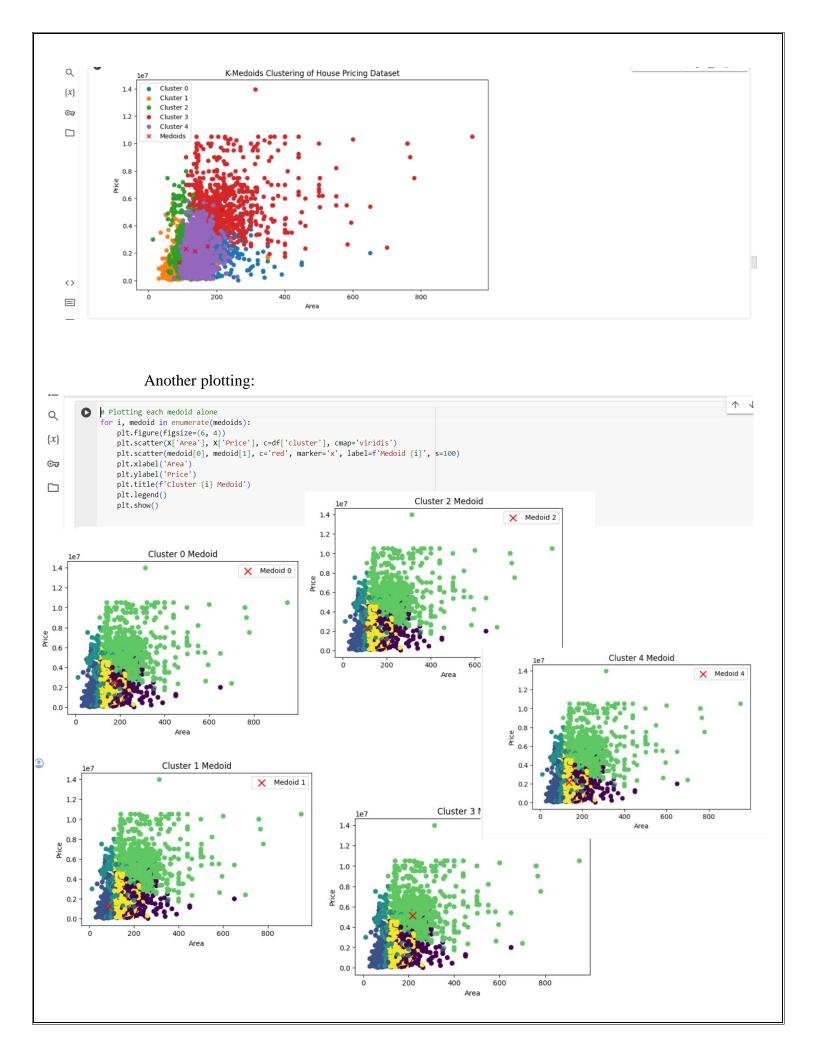


• K-medoids Clustering:

- We applied the K-medoids algorithm to the preprocessed dataset to cluster houses into K distinct groups based on their characteristics.
- The number of clusters (K) was determined using techniques like the elbow method or silhouette analysis.
- Visualization: We visualized the clustering results using scatter plots, where each point represents a house, colored according to its assigned cluster. Additionally, we marked the medoids of each cluster for better interpretation.

```
[ ] import pandas as pd
Q
              import matplotlib.pyplot as plt
              from sklearn_extra.cluster import KMedoids
{x}
              from sklearn.preprocessing import StandardScaler
             # Assuming 'df' is your DataFrame containing the dataset
⊙
             X = df[['Area', 'Price', 'Bedrooms', 'Bathrooms', 'Level']]
# Standardize the data
             scaler = StandardScaler()
             X_scaled = scaler.fit_transform(X)
             n clusters = 5
              \label{eq:kmedoids} \begin{tabular}{ll} $\mathsf{KMedoids}(n_{clusters} = n_{clusters}, \ random_{state} = 0).fit(X_{scaled}) \end{tabular}
              # Add cluster labels to the original DataFrame
             df['cluster'] = kmedoids.labels_
              # Print the cluster centers (medoids)
              medoids = scaler.inverse_transform(kmedoids.cluster_centers_)
              print("Cluster medoids:"
             print(pd.DataFrame(medoids, columns=X.columns))
             # Print the cluster sizes
             print("Cluster sizes:"
<>
             print(df['cluster'].value_counts())
             # Plotting the clusters
\equiv
              plt.figure(figsize=(10, 6))
             for cluster_label in range(n_clusters):
    cluster_data = X[df['cluster'] == cluster_label]
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```
plt.scatter(cluster_data['Area'], cluster_data['Price'], label=f'Cluster {cluster_label}')
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Q
          # Plotting the medoids
          plt.scatter(medoids[:, 0], medoids[:, 1], c='red', marker='x', label='Medoids')
\{x\}
          plt.xlabel('Area')
⊙
          plt.ylabel('Price')
          plt.title('K-Medoids Clustering of House Pricing Dataset')
          plt.legend()
plt.show()
       Cluster medoids:
                     Price Bedrooms Bathrooms Level
          0 174.0 2500000.0 3.0 3.0 2.0
          1 90.0 1300000.0
                                 2.0
                                          1.0
                                                 2.0
          2 110.0 2310000.0
                                2.0 2.0 1.0
          3 217.0 5150000.0 3.0 3.0 1.0
4 137.0 2150000.0 3.0 2.0 2.0
          Cluster sizes:
          cluster
          4 1104
               848
              810
               777
              669
          Name: count, dtype: int64
()
```



• Hierarchical Clustering:

- We also employed hierarchical clustering, a bottom-up approach that forms a hierarchy of clusters, to segment the houses based on their similarity.
- Different linkage methods such as complete, average, and ward linkage were explored to measure the distance between clusters.
- Visualization: We visualized the hierarchical clustering dendrogram, illustrating
 the hierarchical structure of the clusters. Additionally, we created dendrograms
 for each linkage method to compare the clustering results visually.

Hierarchical clustering

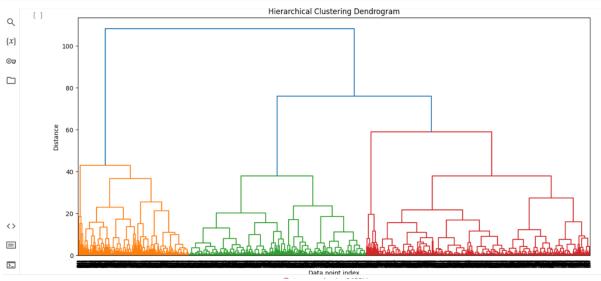
```
from sklearn.preprocessing import StandardScaler
from scipy.cluster.hierarchy import dendrogram, linkage

# Assuming 'df' is your DataFrame containing the dataset
X = df[['Area', 'Price', 'Bedrooms', 'Bathrooms', 'Level']]

# Standardize the data
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)

# Perform hierarchical clustering
Z = linkage(X_scaled, method='ward')

# Plot the dendrogram
plt.figure(figsize=(15, 7))
dendrogram(Z)
plt.title('Hierarchical clustering Dendrogram')
plt.xlabel('Data point index')
plt.ylabel('Distance')
plt.show()
```



Evaluation:

- We evaluated the clustering results of both algorithms using metrics such as silhouette score, Davies–Bouldin index, and visual inspection of cluster separation.
- The silhouette score measures the cohesion and separation of clusters, with higher scores indicating better-defined clusters.
- The Davies–Bouldin index assesses

```
from sklearn.metrics import silhouette_score, davies_bouldin_score
{\tt def\ evaluate\_clustering(clustering\_algorithm,\ data):}
    # Fit clustering algorithm
    cluster_labels = clustering_algorithm.fit_predict(data)
    # Silhouette score
    silhouette_avg = silhouette_score(data, cluster_labels)
    # Davies-Bouldin index
    db_index = davies_bouldin_score(data, cluster_labels)
    return silhouette_avg, db_index
# Example usage:
# Assuming 'kmedoids' is the K-medoids clustering algorithm and 'hierarchical' is the hierarchical clustering algorithm
# Assuming 'scaled_features' is the preprocessed and scaled feature data
# Evaluate K-medoids clustering
kmedoids_silhouette, kmedoids_db = evaluate_clustering(kmedoids, scaled_features)
print("K-medoids Silhouette Score:", kmedoids_silhouette)
print("K-medoids Davies-Bouldin Index:", kmedoids_db)
# Evaluate Hierarchical clustering
hierarchical_silhouette, hierarchical_db = evaluate_clustering(hierarchical, scaled_features)
print("\nHierarchical Silhouette Score:", hierarchical_silhouette)
print("Hierarchical Davies-Bouldin Index:", hierarchical_db)
```

Notebook Link:

https://colab.research.google.com/drive/15qBezbJfrh4F8cdN6ZFYPap-PGmfIzVU?usp=sharing

Conclusion:	
•	In conclusion, we compared the effectiveness of K-medoids and hierarchical clustering algorithms in segmenting a house pricing dataset.
•	Both algorithms provided valuable insights into the underlying structure of the dataset, enabling us to identify distinct groups of houses based on their characteristics and prices.
•	The choice of clustering algorithm depends on factors such as dataset size, structure, and the desired level of interpretability.