Clustering

Assignment Questions







Theoretical Questions:

- 1. What is unsupervised learning in the context of machine learning?
- 2. How does K-Means clustering algorithm work?
- 3. Explain the concept of a dendrogram in hierarchical clustering.
- 4. What is the main difference between K-Means and Hierarchical Clustering?
- 5. What are the advantages of DBSCAN over K-Means?
- 6. When would you use Silhouette Score in clustering?
- 7. What are the limitations of Hierarchical Clustering?
- 8. Why is feature scaling important in clustering algorithms like K-Means?
- 9. How does DBSCAN identify noise points?
- 10. Define inertia in the context of K-Means.
- 11. What is the elbow method in K-Means clustering?
- 12. Describe the concept of "density" in DBSCAN.
- 13. Can hierarchical clustering be used on categorical data?
- 14. What does a negative Silhouette Score indicate?
- 15. Explain the term "linkage criteria" in hierarchical clustering.
- 16. Why might K-Means clustering perform poorly on data with varying cluster sizes or densities?
- 17. What are the core parameters in DBSCAN, and how do they influence clustering?
- 18. How does K-Means++ improve upon standard K-Means initialization?
- 19. What is agglomerative clustering?
- 20. What makes Silhouette Score a better metric than just inertia for model evaluation?



Practical Questions:

- 21. Generate synthetic data with 4 centers using make_blobs and apply K-Means clustering. Visualize using a scatter plot.
- 22. Load the Iris dataset and use Agglomerative Clustering to group the data into 3 clusters. Display the first 10 predicted labels.
- 23. Generate synthetic data using make_moons and apply DBSCAN. Highlight outliers in the plot.
- 24. Load the Wine dataset and apply K-Means clustering after standardizing the features. Print the size of each cluster.
- 25. Use make_circles to generate synthetic data and cluster it using DBSCAN. Plot the result.
- 26. Load the Breast Cancer dataset, apply MinMaxScaler, and use K-Means with 2 clusters. Output the cluster centroids.
- 27. Generate synthetic data using make_blobs with varying cluster standard deviations and cluster with DBSCAN.
- 28. Load the Digits dataset, reduce it to 2D using PCA, and visualize clusters from K-Means.
- 29. Create synthetic data using make_blobs and evaluate silhouette scores for k = 2 to 5. Display as a bar chart.
- 30. Load the Iris dataset and use hierarchical clustering to group data. Plot a dendrogram with average linkage.
- 31. Generate synthetic data with overlapping clusters using make_blobs, then apply K-Means and visualize with decision boundaries.
- 32. Load the Digits dataset and apply DBSCAN after reducing dimensions with t-SNE. Visualize the results.
- 33. Generate synthetic data using make_blobs and apply Agglomerative Clustering with complete linkage. Plot the result.
- 34. Load the Breast Cancer dataset and compare inertia values for K = 2 to 6 using K-Means. Show results in a line plot.
- 35. Generate synthetic concentric circles using make_circles and cluster using Agglomerative Clustering with single linkage.
- 36. Use the Wine dataset, apply DBSCAN after scaling the data, and count the number of clusters (excluding noise)
- 37. Generate synthetic data with make_blobs and apply KMeans. Then plot the cluster centers on top of the data points.
- 38. Load the Iris dataset, cluster with DBSCAN, and print how many samples were identified as noise.
- 39. Generate synthetic non-linearly separable data using make_moons, apply K-Means, and visualize the clustering result.
- 40. Load the Digits dataset, apply PCA to reduce to 3 components, then use KMeans and visualize with a 3D scatter plot.



- 41. Generate synthetic blobs with 5 centers and apply KMeans. Then use silhouette_score to evaluate the clustering.
- 42. Load the Breast Cancer dataset, reduce dimensionality using PCA, and apply Agglomerative Clustering. Visualize in 2D.
- 43. Generate noisy circular data using make_circles and visualize clustering results from KMeans and DBSCAN side-by-side.
- 44. Load the Iris dataset and plot the Silhouette Coefficient for each sample after KMeans clustering.
- 45. Generate synthetic data using make_blobs and apply Agglomerative Clustering with 'average' linkage. Visualize clusters.
- 46. Load the Wine dataset, apply KMeans, and visualize the cluster assignments in a seaborn pairplot (first 4 features).
- 47. Generate noisy blobs using make_blobs and use DBSCAN to identify both clusters and noise points. Print the count.
- 48. Load the Digits dataset, reduce dimensions using t-SNE, then apply Agglomerative Clustering and plot the clusters.