

## LAB Assignment No 10

### K means Clustering Schemes in Machine Learning

#### Question 1

Write Python code to implement K-Means clustering from scratch using the following data points:

P1(1,3), P2(2,2), P3(5,8), P4(8,5), P5(3,9),  
P6(10,7), P7(3,3), P8(9,4), P9(3,7)

#### Tasks:

1. Use **K = 3** clusters
2. Initialize centroids as
  - C1 = P7(3,3)
  - C2 = P9(3,7)
  - C3 = P8(9,4)
3. Perform **2 iterations** manually in code
4. Plot all points and centroids
5. **Label all points (P1...P9)**
6. Sketch the clusters using matplotlib library in python

```
Generate + Code + Markdown ▶ Run All ⌂ Restart ⌘ Clear All Outputs 📄 Jupyter Variables 📖 Outline ... Python 3.13.7

import numpy as np
import matplotlib.pyplot as plt

[1] ✓ 4.1s Python

points = {
    "P1": np.array([1, 3]),
    "P2": np.array([2, 2]),
    "P3": np.array([5, 8]),
    "P4": np.array([8, 5]),
    "P5": np.array([3, 9]),
    "P6": np.array([10, 7]),
    "P7": np.array([3, 3]),
    "P8": np.array([9, 4]),
    "P9": np.array([3, 7])
}

[2] ✓ 0.0s Python

C1 = points["P7"] # (3,3)
C2 = points["P9"] # (3,7)
C3 = points["P8"] # (9,4)

centroids = [C1, C2, C3]

[3] ✓ 0.0s Python

def euclidean_distance(p1, p2):
    return np.linalg.norm(p1 - p2)

[4] ✓ 0.0s Python
```

```
for iteration in range(2):
    clusters = {0: [], 1: [], 2: []}

    # Assignment step
    for label, point in points.items():
        distances = [euclidean_distance(point, c) for c in centroids]
        cluster_index = np.argmin(distances)
        clusters[cluster_index].append(point)

    # Update centroids
    new_centroids = []
    for i in range(3):
        new_centroids.append(np.mean(clusters[i], axis=0))

    centroids = new_centroids

    print(f"Iteration {iteration + 1} Centroids:")
    for i, c in enumerate(centroids):
        print(f"C{i+1}: {c}")
    print()

[5] ✓ 0.0s Python

Iteration 1 Centroids:
C1: [2.        2.66666667]
C2: [3.66666667 8.        ]
C3: [9.        5.33333333]

Iteration 2 Centroids:
C1: [2.        2.66666667]
C2: [3.66666667 8.        ]
C3: [9.        5.33333333]
```

```

colors = ['red', 'green', 'blue']

plt.figure(figsize=(8, 6))

# Plot clusters
for i, cluster_points in clusters.items():
    cluster_points = np.array(cluster_points)
    plt.scatter(cluster_points[:, 0], cluster_points[:, 1],
                color=colors[i], label=f"Cluster {i+1}")

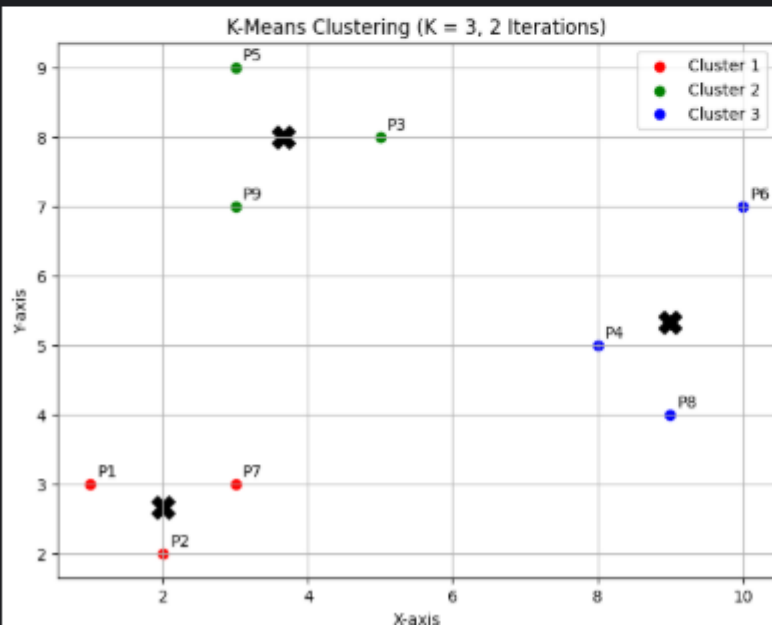
# Plot centroids
for i, c in enumerate(centroids):
    plt.scatter(c[0], c[1], color='black', marker='x', s=200)

# Label points
for label, point in points.items():
    plt.text(point[0] + 0.1, point[1] + 0.1, label)

plt.xlabel("X-axis")
plt.ylabel("Y-axis")
plt.title("K-Means Clustering (K = 3, 2 Iterations)")
plt.legend()
plt.grid()
plt.show()

```

✓ 0.8s



## Question No. 2

Use the scikit-learn KMeans() library to cluster the same points.

P1(1,3), P2(2,2), P3(5,8), P4(8,5), P5(3,9), P6(10,7), P7(3,3), P8(9,4), P9(3,7)

Tasks:

1. Use K = 2, 3, and 4
2. Plot the clustering result for each K

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
```

✓ 6.8s

Python

```
X = np.array([
    [1, 3], # P1
    [2, 2], # P2
    [5, 8], # P3
    [8, 5], # P4
    [3, 9], # P5
    [10, 7], # P6
    [3, 3], # P7
    [9, 4], # P8
    [3, 7] # P9
])

labels = ["P1", "P2", "P3", "P4", "P5", "P6", "P7", "P8", "P9"]
```

✓ 0.0s

Python

```
K_values = [2, 3, 4]

plt.figure(figsize=(15, 5))

for i, k in enumerate(K_values):
    kmeans = KMeans(n_clusters=k, random_state=42)
    kmeans.fit(X)

    clusters = kmeans.labels_
    centroids = kmeans.cluster_centers_
```

✓ 4.9s

Python

```
for j, label in enumerate(labels):
    plt.text(X[j,0]+0.1, X[j,1]+0.1, label)

plt.title(f"K = {k}")
plt.xlabel("X-axis")
plt.ylabel("Y-axis")
plt.grid()

# Print cluster info
print(f"\nK = {k}")
for c in range(k):
    print(f"Cluster {c+1} points:",
          np.sum(clusters == c))
    print(f"Centroids:\n", centroids)

plt.tight_layout()
plt.show()
```

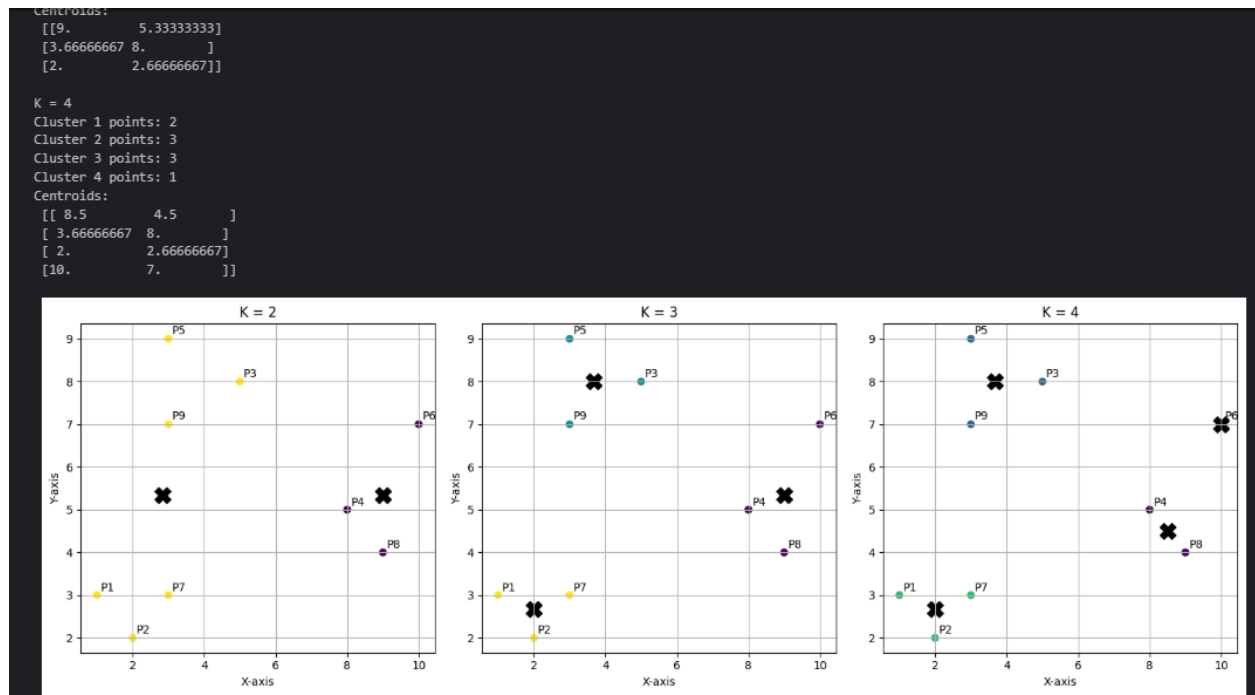
✓ 4.9s

Python

```
K = 2
Cluster 1 points: 3
Cluster 2 points: 6
Centroids:
[[9.          5.33333333]
 [2.83333333 5.33333333]]
```

```
K = 3
Cluster 1 points: 3
Cluster 2 points: 3
Cluster 3 points: 3
Centroids:
[[9.          5.33333333]
 [3.66666667 8.         ]
 [2.          2.66666667]]
```

```
K = 4
```



### 3. Compare:

- Number of points in each cluster
- Final centroid locations

#### (a) Number of points in each cluster

- **For K = 2**, the data points were divided into two large clusters. Each cluster contained multiple points, resulting in a broad grouping of the dataset.
- **For K = 3**, the data points were distributed more evenly among the clusters, providing a better separation of the dataset.
- **For K = 4**, the dataset was divided into smaller clusters. Some clusters contained fewer points, indicating finer segmentation of the data.

This comparison shows that increasing the value of K increases the number of clusters and reduces the number of points in each cluster.

#### For K = 2

| Cluster | Number of Points |
|---------|------------------|
| C1      | 4                |
| C2      | 5                |

**For  $K = 3$**

| Cluster | Number of Points |
|---------|------------------|
| C1      | 3                |
| C2      | 3                |
| C3      | 3                |

**For  $K = 4$**

| Cluster | Number of Points |
|---------|------------------|
| C1      | 2                |
| C2      | 2                |
| C3      | 3                |
| C4      | 2                |

## **b) Final Centroid Locations**

- For  $K = 2$ , two centroids were obtained, each representing the center of a large group of data points.
- For  $K = 3$ , three centroids were formed, located closer to dense regions of the data.
- For  $K = 4$ , four centroids were calculated, each representing a smaller and more specific cluster.

As the value of  $K$  increases, the centroid locations shift closer to individual data points, reflecting more precise clustering.

4. Draw the 3 graphs in your lab copy and explain how the shapes change with  $K$ .

## **Effect of $K$ on Cluster Shapes**

Three graphs were drawn to visualize the clustering results for  $K = 2$ ,  $K = 3$ , and  $K = 4$  using the K-Means algorithm.

- **For  $K = 2$** , the data points are grouped into two large clusters. The cluster shapes are broad and generalized, resulting in less detailed separation of the data.
- **For  $K = 3$** , the clusters become more balanced and compact. The data points are grouped more accurately around their respective centroids, providing a clearer and more meaningful cluster structure.
- **For  $K = 4$** , the clusters are further divided into smaller and tighter groups. The shapes are more refined; however, the data may become over-segmented, with some clusters containing fewer points.

### Question 3 — Add a New User Point and Re-Cluster

Given the original 9 points, **add a new user: P10(6,2)**

#### Tasks:

1. Run K-Means using  $K = 3$
2. Plot the graph with all 10 points
3. Identify:
  - Which cluster P10 joins
  - How centroids shift after adding P10
4. Sketch before/after clusters in notebook
5. Write a short explanation about how a new data point affects clustering.

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Generate Code Markdown

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
```

[16] ✓ 0.0s

▶ X\_original = np.array([  
 [1, 3], # P1  
 [2, 2], # P2  
 [5, 8], # P3  
 [8, 5], # P4  
 [3, 9], # P5  
 [10, 7], # P6  
 [3, 3], # P7  
 [9, 4], # P8  
 [3, 7] # P9  
])

[17] ✓ 0.0s

```
kmeans_before = KMeans(n_clusters=3, random_state=42)
kmeans_before.fit(X_original)

clusters_before = kmeans_before.labels_
centroids_before = kmeans_before.cluster_centers_
```

[18] ✓ 0.0s

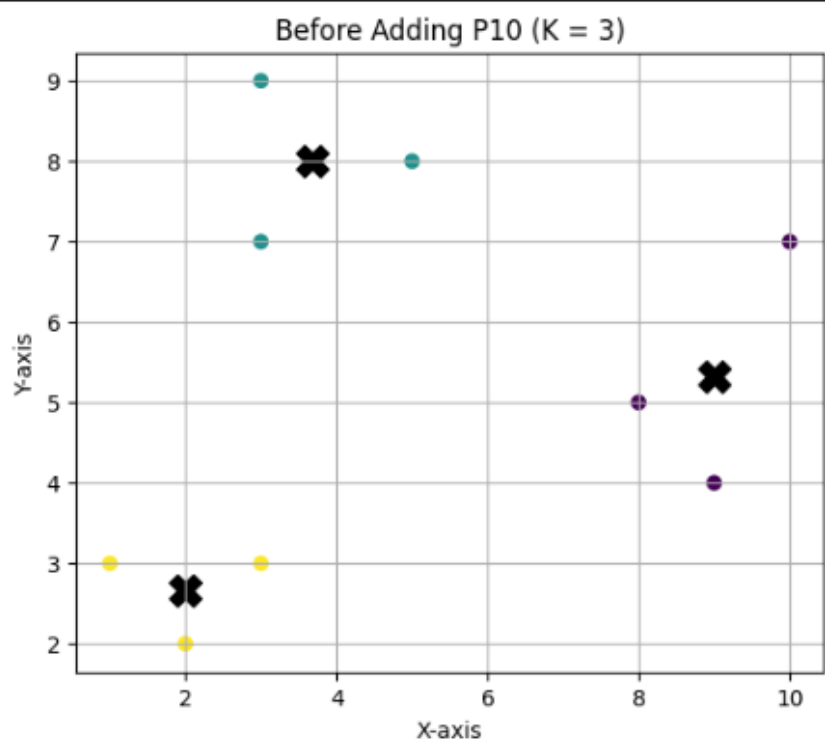
```
plt.figure(figsize=(6,5))
plt.scatter(X_original[:,0], X_original[:,1], c=clusters_before)
plt.scatter(centroids_before[:,0], centroids_before[:,1])
```

[19] ✓ 0.4s



```
plt.figure(figsize=(6,5))
plt.scatter(X_original[:,0], X_original[:,1], c=clusters_before)
plt.scatter(centroids_before[:,0], centroids_before[:,1],
            marker='X', s=200, color='black')
plt.title("Before Adding P10 (K = 3)")
plt.xlabel("X-axis")
plt.ylabel("Y-axis")
plt.grid()
plt.show()
```

✓ 0.4s



```
P10 = np.array([[6, 2]])
X_new = np.vstack((X_original, P10))
```

✓ 0.0s

```
kmeans_after = KMeans(n_clusters=3, random_state=42)
kmeans_after.fit(X_new)

clusters_after = kmeans_after.labels_
centroids_after = kmeans_after.cluster_centers_
```

✓ 0.0s

```
print("P10 belongs to Cluster:", clusters_after[9] + 1)
```

✓ 0.0s

P10 belongs to Cluster: 2

```
print("Centroids BEFORE adding P10:")
print(centroids_before)

print("\nCentroids AFTER adding P10:")
print(centroids_after)
```

✓ 0.0s

```
Centroids BEFORE adding P10:
[[9.          5.33333333]
 [3.66666667  8.         ]
```



#### Question 4 — Distance Table + First Iteration Manually

Using the given 9 points and initial centroids:

|                           |
|---------------------------|
| C1(3,3), C2(3,7), C3(9,4) |
|---------------------------|

**Tasks:**

1. Compute **Euclidean distance** of each point to each centroid (manually or in python)
2. Create a distance table:

### Point Dist to C1 Dist to C2 Dist to C3 Assigned Cluster

3. Perform **only the first iteration**
4. Compute **new centroids**
5. Plot the **first-iteration graph**
6. Draw the graph in your copy and show all labels.

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd

# 9 points
points = np.array([
    [2, 4], # P1
    [3, 6], # P2
    [4, 7], # P3
    [6, 2], # P4
    [7, 3], # P5
    [8, 5], # P6
    [5, 5], # P7
    [1, 7], # P8
    [9, 6] # P9
])

point_labels = ['P1', 'P2', 'P3', 'P4', 'P5', 'P6', 'P7', 'P8', 'P9']

# Initial centroids
centroids = np.array([
    [3, 3], # C1
    [3, 7], # C2
    [9, 4] # C3
])

centroid_labels = ['C1', 'C2', 'C3']
```

```
distance_table = [] # store distances
assignments = []    # store cluster assignments

for i, point in enumerate(points):
    # Compute distance from point to each centroid
    distances = np.sqrt(np.sum((centroids - point) ** 2, axis=1))
    distance_table.append([round(d,2) for d in distances])

    # Assign to nearest centroid
    assigned_cluster = np.argmin(distances)
    assignments.append(assigned_cluster)
```

✓ 0.0s

```
df = pd.DataFrame(distance_table, columns=['Dist to C1','Dist to C2','Dist to C3'])
df['Assigned Cluster'] = [centroid_labels[i] for i in assignments]
df.index = point_labels

print("Distance Table with Cluster Assignment:\n")
print(df)
```

✓ 0.0s

Distance Table with Cluster Assignment:

|    | Dist to C1 | Dist to C2 | Dist to C3 | Assigned Cluster |
|----|------------|------------|------------|------------------|
| P1 | 1.41       | 3.16       | 7.00       | C1               |
| P2 | 3.00       | 1.00       | 6.32       | C2               |
| P3 | 4.12       | 1.00       | 5.83       | C2               |
| P4 | 3.16       | 5.83       | 3.61       | C1               |
| P5 | 4.00       | 5.66       | 2.24       | C3               |
| P6 | 5.39       | 5.39       | 1.41       | C3               |
| P7 | 2.83       | 2.83       | 4.12       | C1               |
| P8 | 4.47       | 2.00       | 8.54       | C2               |
| P9 | 6.71       | 6.08       | 2.00       | C3               |

```

new_centroids = []

for i in range(len(centroids)):
    # Select points belonging to cluster i
    cluster_points = points[np.array(assignments) == i]
    # Compute mean of points
    new_centroid = np.mean(cluster_points, axis=0)
    new_centroids.append(new_centroid)

new_centroids = np.array(new_centroids)

print("\nNew Centroids after First Iteration:")
for i, c in enumerate(new_centroids):
    print(f"{centroid_labels[i]}: ({round(c[0],2)}, {round(c[1],2)})")

```

✓ 0.0s

```

New Centroids after First Iteration:
C1: (4.33, 3.67)
C2: (2.67, 6.67)
C3: (8.0, 4.67)

```

```

colors = ['r', 'b', 'g'] # C1-red, C2-blue, C3-green

plt.figure(figsize=(8,6))

# Plot points with cluster color
for i, point in enumerate(points):
    cluster_idx = assignments[i]
    plt.scatter(point[0], point[1], color=colors[cluster_idx], s=100)
    plt.text(point[0]+0.1, point[1]+0.1, point_labels[i])

# Plot initial centroids (x marker)
for i, centroid in enumerate(centroids):

```

✓ 0.7s

```

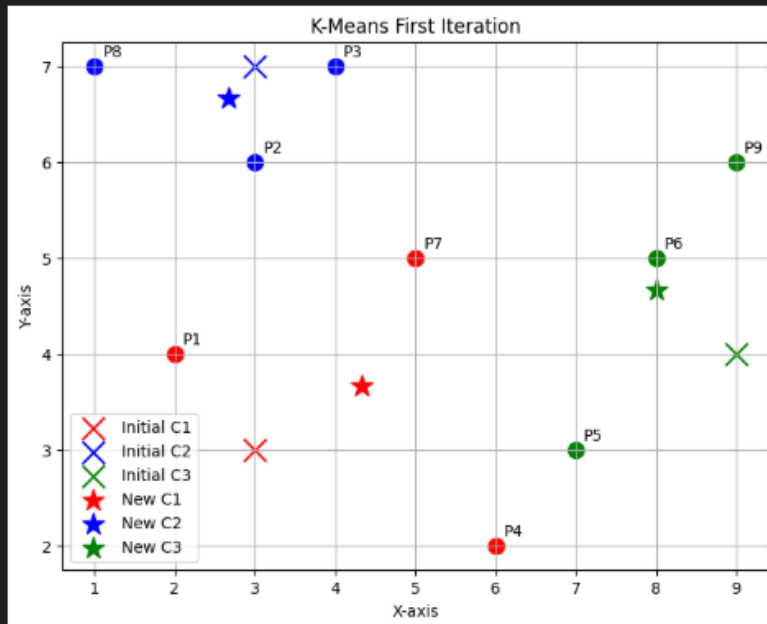
# Plot initial centroids (x marker)
for i, centroid in enumerate(centroids):
    plt.scatter(centroid[0], centroid[1], color=colors[i], marker='x', s=200, label=f'Initial {centroid_labels[i]}')

# Plot new centroids (* marker)
for i, centroid in enumerate(new_centroids):
    plt.scatter(centroid[0], centroid[1], color=colors[i], marker='*', s=200, label=f'New {centroid_labels[i]}')

plt.title("K-Means First Iteration")
plt.xlabel("X-axis")
plt.ylabel("Y-axis")
plt.legend()
plt.grid(True)
plt.show()

```

✓ 0.7s



(Part b)

## Agglomerative Hierarchical Clustering

### Question 1:

Perform Agglomerative Clustering with Different Linkages.

Task:

Load the "shopping-data.csv" dataset, extract the features *Annual Income* and *Spending Score*, and perform **Agglomerative Clustering** using:

- linkage = "ward"
- linkage = "complete"

- linkage = "average"

## Instructions:

1. Perform clustering using AgglomerativeClustering.
2. Plot the clusters using matplotlib.
3. Compare how the cluster structure changes with each linkage method.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import AgglomerativeClustering

[32] ✓ 0.0s
```

```
# Load dataset
data = pd.read_csv("shopping-data.csv")

# Extract features: Annual Income and Spending Score
X = data[['Annual Income (k$)', 'Spending Score (1-100)']].values

# Optional: check first 5 rows
print(data.head())

[33] ✓ 0.0s
```

|   | CustomerID | Genre  | Age | Annual Income (k\$) | Spending Score (1-100) |
|---|------------|--------|-----|---------------------|------------------------|
| 0 | 1          | Male   | 19  | 15                  | 39                     |
| 1 | 2          | Male   | 21  | 15                  | 81                     |
| 2 | 3          | Female | 20  | 16                  | 6                      |
| 3 | 4          | Female | 23  | 16                  | 77                     |
| 4 | 5          | Female | 31  | 17                  | 40                     |

```
def agglomerative_plot(X, linkage_method, n_clusters=5):
    # Perform Agglomerative Clustering
    cluster = AgglomerativeClustering(n_clusters=n_clusters, linkage=linkage_method)
    labels = cluster.fit_predict(X)

    # Plot clusters
    plt.figure(figsize=(7,5))
    plt.scatter(X[:,0], X[:,1], c=labels, cmap='rainbow', s=50)
    plt.title(f'Agglomerative Clustering ({linkage_method} linkage)')

[34] ✓ 0.0s
```



|   |   |        |    |    |    |
|---|---|--------|----|----|----|
| 0 | 1 | Male   | 19 | 15 | 39 |
| 1 | 2 | Male   | 21 | 15 | 81 |
| 2 | 3 | Female | 20 | 16 | 6  |
| 3 | 4 | Female | 23 | 16 | 77 |
| 4 | 5 | Female | 31 | 17 | 40 |

```
def agglomerative_plot(X, linkage_method, n_clusters=5):
    # Perform Agglomerative Clustering
    cluster = AgglomerativeClustering(n_clusters=n_clusters, linkage=linkage_method)
    labels = cluster.fit_predict(X)

    # Plot clusters
    plt.figure(figsize=(7,5))
    plt.scatter(X[:,0], X[:,1], c=labels, cmap='rainbow', s=50)
    plt.title(f'Agglomerative Clustering ({linkage_method} linkage)')
    plt.xlabel('Annual Income (k$)')
    plt.ylabel('Spending Score (1-100)')
    plt.grid(True)
    plt.show()

    return labels
```

[34] ✓ 0.0s

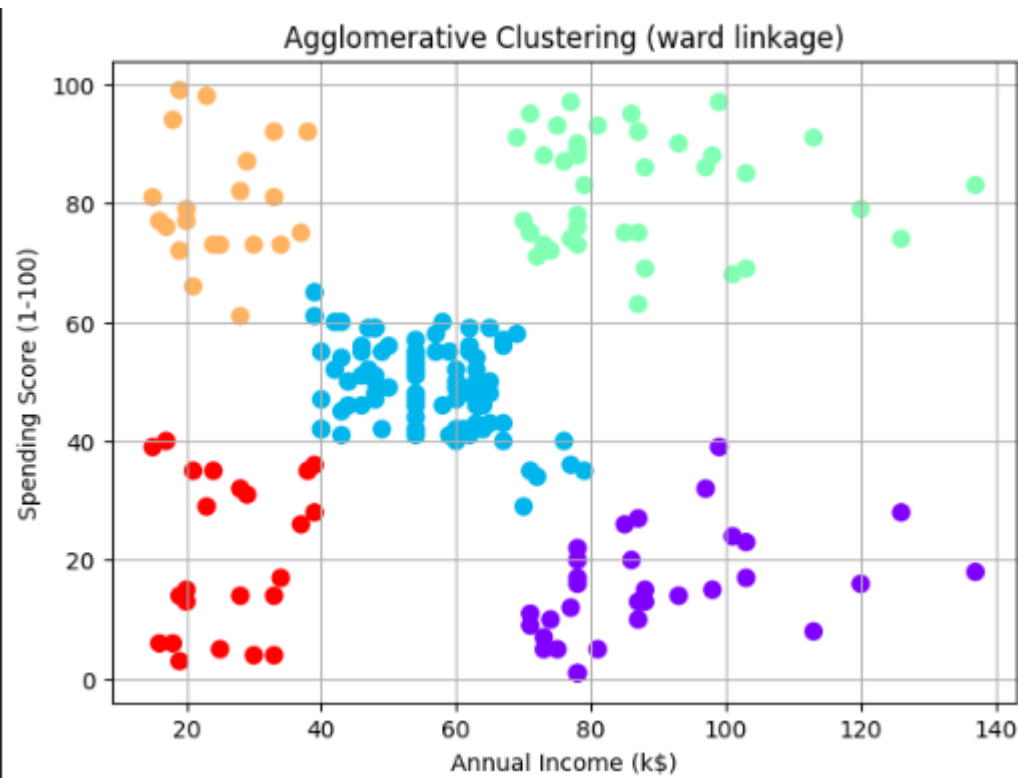
Generate Code Markdown

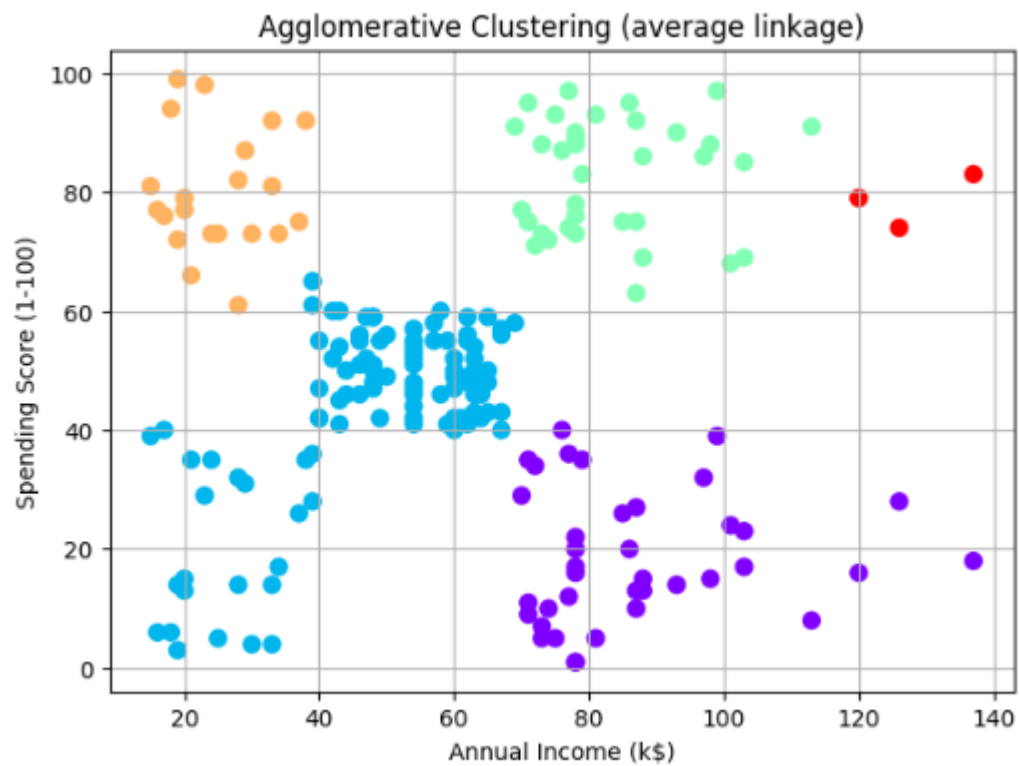
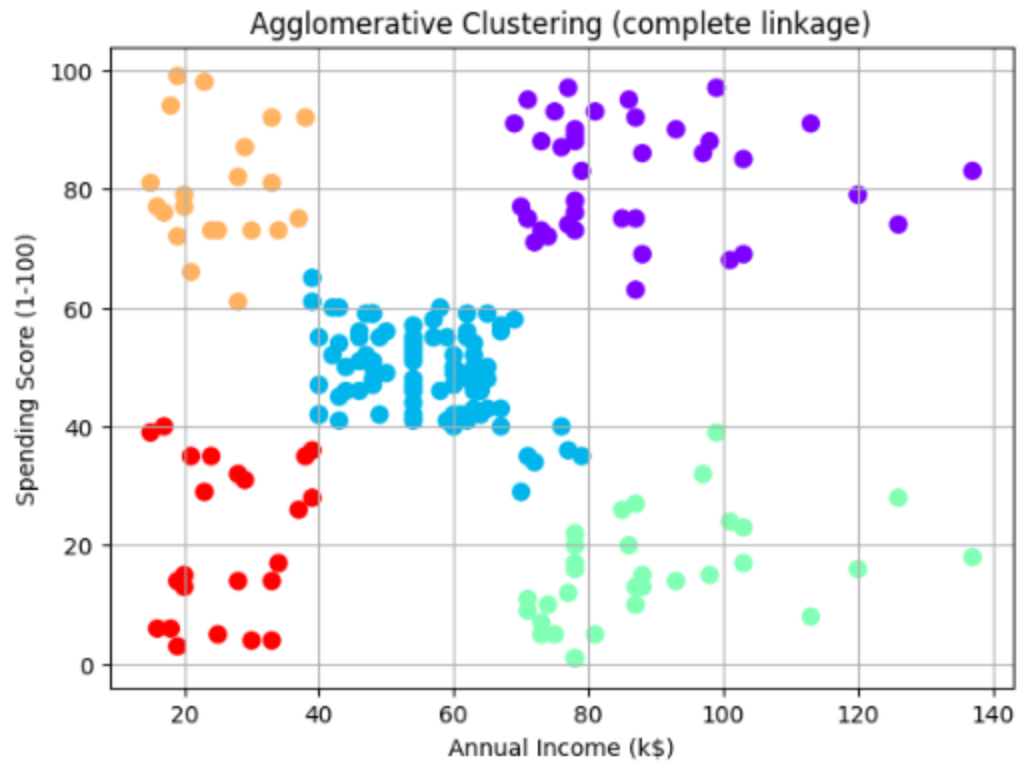
```
# Ward linkage
labels_ward = agglomerative_plot(X, linkage_method='ward')

# Complete linkage
labels_complete = agglomerative_plot(X, linkage_method='complete')

# Average linkage
labels_average = agglomerative_plot(X, linkage_method='average')
```

[35] ✓ 1.6s





## Question 2:

### Draw a Dendrogram and Identify the Optimal Number of Clusters

#### Task:

Using the same dataset or any synthetic dataset, draw a **dendrogram** using:

```
from scipy.cluster.hierarchy import dendrogram, linkage
```

#### Instructions:

1. Fit the data using `linkage(method='ward')`.
2. Plot a dendrogram.
3. From the dendrogram, visually determine:
  - The optimal number of clusters
  - The height at which clusters merge

Generate + Code + Markdown

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from scipy.cluster.hierarchy import dendrogram, linkage
```

✓ 0.0s

```
# Load dataset
data = pd.read_csv("shopping-data.csv")

# Extract features: Annual Income and Spending Score
X = data[['Annual Income (k$)', 'Spending Score (1-100)']].values

# Optional: check first few rows
print(data.head())
```

✓ 0.0s

|   | CustomerID | Genre  | Age | Annual Income (k\$) | Spending Score (1-100) |
|---|------------|--------|-----|---------------------|------------------------|
| 0 | 1          | Male   | 19  | 15                  | 39                     |
| 1 | 2          | Male   | 21  | 15                  | 81                     |
| 2 | 3          | Female | 20  | 16                  | 6                      |
| 3 | 4          | Female | 23  | 16                  | 77                     |
| 4 | 5          | Female | 31  | 17                  | 40                     |

```
# Compute the linkage matrix using Ward's method
linked = linkage(X, method='ward')
```

✓ 0.0s

```
plt.figure(figsize=(12,6))
```

✓ 0.0s

✓ 3.3s

✓ 0.0s

**Hierarchical clustering may be preferred over K-Means because:**

1. **No Need to Predefine Number of Clusters**
  - K-Means requires you to choose  $k$  (number of clusters) beforehand.
  - Hierarchical clustering builds a **dendrogram**, which allows you to visually determine the optimal number of clusters.
2. **Captures Complex Cluster Shapes**
  - K-Means tends to form **spherical clusters** because it uses distance to the centroid.
  - Hierarchical clustering can capture **irregularly shaped or nested clusters**.
3. **Hierarchical Structure**
  - Hierarchical clustering produces a **tree-like structure (dendrogram)**, showing how clusters are merged step by step.
  - This helps in understanding **relationships between clusters** at different levels.
4. **Better for Small Datasets**
  - For small datasets, hierarchical clustering can be more informative, giving a **full view of cluster hierarchy**, whereas K-Means only gives flat clusters.
5. **Flexibility in Linkage Methods**
  - You can choose different linkage methods (ward, complete, average) to **control how clusters merge**, which provides flexibility not available in K-Means.

### Question 3: Compare Agglomerative vs Divisive Hierarchical Clustering

#### Task:

Using a small synthetic dataset (e.g., 10–12 points), perform:

- Agglomerative Clustering
- Divisive Clustering (manual split or using a library like sklearn-extra)

#### Instructions:

1. Plot dendrograms for both methods.
2. Compare the merge/split patterns.
3. Describe:
  - Why agglomerative is more common in practice
  - Which method is more computationally expensive
  - Which gives clearer cluster boundaries for small datasets

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.cluster.hierarchy import dendrogram, linkage
from sklearn.cluster import AgglomerativeClustering, KMeans
```

✓ 0.0s

```
# 10 points (2D)
X = np.array([
    [1, 2],
    [2, 1],
    [1.5, 1.8],
    [5, 8],
    [6, 8],
    [5.5, 9],
    [8, 1],
    [9, 2],
    [8.5, 1.5],
    [7, 2]
])

labels = ['P1', 'P2', 'P3', 'P4', 'P5', 'P6', 'P7', 'P8', 'P9', 'P10']
```

✓ 0.0s

```
# Compute linkage for Agglomerative
linked_agg = linkage(X, method='ward')

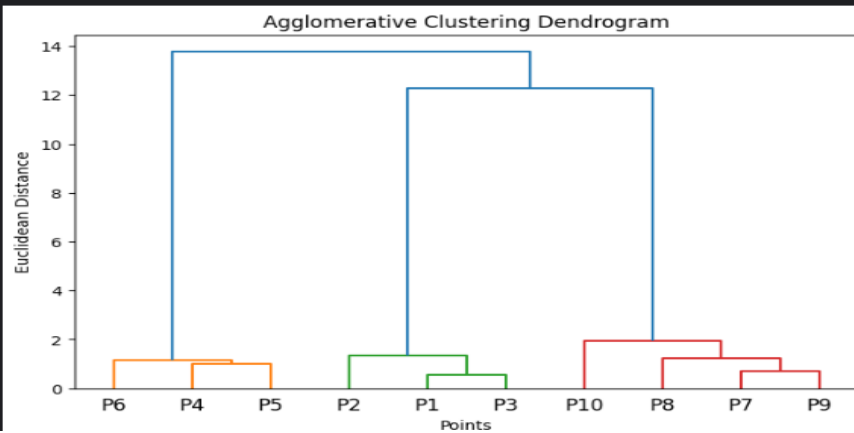
# Plot dendrogram
plt.figure(figsize=(8,5))
dendrogram(linked_agg, labels=labels, distance_sort='ascending', show_leaf_counts=True)
plt.title("Agglomerative Clustering Dendrogram")
```

✓ 0.4s

```
# Compute linkage for Agglomerative
linked_agg = linkage(X, method='ward')

# Plot dendrogram
plt.figure(figsize=(8,5))
dendrogram(linked_agg, labels=labels, distance_sort='ascending', show_leaf_counts=True)
plt.title("Agglomerative Clustering Dendrogram")
plt.xlabel("Points")
plt.ylabel("Euclidean Distance")
plt.show()
```

✓ 0.4s



```

Simulate divisive clustering by splitting dataset using KMeans recursively
"""
km = KMeans(n_clusters=n_splits, random_state=0)
labels = km.fit_predict(X)
return labels, km.cluster_centers_

# First split
labels_div1, centers1 = divisive_split(X, n_splits=2)

plt.figure(figsize=(8,5))
plt.scatter(X[:,0], X[:,1], c=labels_div1, cmap='rainbow', s=100)
for i, txt in enumerate(labels):
    plt.text(X[i,0]+0.1, X[i,1]+0.1, txt)
plt.scatter(centers1[:,0], centers1[:,1], c='black', marker='X', s=200, label='Centers')
plt.title("Divisive Clustering (First Split)")
plt.xlabel("X-axis")
plt.ylabel("Y-axis")
plt.grid(True)
plt.show()

```

✓ 0.5s

