

**Aim-:**Implement K-Means clustering/ hierarchical clustering on sales\_data\_sample.csv dataset. Determine the number of clusters using the elbow method.

### INPUT

```
import pandas as pd

import matplotlib.pyplot as plt from sklearn.cluster

import KMeans from sklearn.preprocessing

import StandardScaler from scipy.cluster.hierarchy

import dendrogram, linkage from sklearn.cluster

import AgglomerativeClustering

import numpy as np

import seaborn as sns

# Load the dataset

# Load the dataset with a specified encoding

data = pd.read_csv('sales_data_sample.csv', encoding='ISO-8859-1')

# Preprocess the dataset (choose relevant features)

# Replace 'SALES', 'QUANTITYORDERED', 'PRICEEACH' with actual column names in
your dataset

features = data[['SALES', 'QUANTITYORDERED', 'PRICEEACH']]

# Standardize the features

scaler =StandardScaler()

scaled_features=scaler.fit_transform(features)

# Elbow method to determine the optimal number of clusters for K-Means
```

```

wcss = []

# within-cluster sum of squares
for i in range(1, 11):

    kmeans = KMeans(n_clusters=i, random_state=42)

    kmeans.fit(scaled_features)

    wcss.append(kmeans.inertia_)


# Plot the elbow curve
plt.figure(figsize=(8, 5))

plt.plot(range(1, 11), wcss, marker='o')

plt.title('Elbow Method for Optimal K (K-Means)')

plt.xlabel('Number of Clusters')

plt.ylabel('WCSS (Within-Cluster Sum of Squares)')

plt.show()


# Automatically determine the optimal number of clusters from the elbow #
This function calculates the "elbow" as the point with the maximum curvature
def optimal_k_elbow(wcss):
    x1, y1 = 1, wcss[0]
    x2, y2 = 10, wcss[-1]

    distances = []

    for i in range(10):

        x0 = i + 1

        y0 = wcss[i]

        numerator = abs((y2 - y1) * x0 - (x2 - x1) * y0 + x2 * y1 - y2 * x1)

        denominator = np.sqrt((y2 - y1) ** 2 + (x2 - x1) ** 2)

        distances.append(numerator / denominator)

```

```
return distances.index(max(distances)) + 1
```

```
optimal_k = optimal_k_elbow(wcss)
```

```
# Apply K-Means clustering with the optimal number of clusters
```

```
kmeans = KMeans(n_clusters=optimal_k, random_state=42)
```

```
kmeans_labels = kmeans.fit_predict(scaled_features)
```

```
# Print the number of clusters for K-Means
```

```
print(f'Number of clusters determined by K-Means (Elbow Method): {optimal_k}')
```

```
# Plot K-Means clusters with colored datapoints
```

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

```
sns.scatterplot(x=scaled_features[:, 0], y=scaled_features[:, 1], hue=kmeans_labels,
```

```
palette='Set1', s=100)
```

```
plt.title(f'K-Means Clustering with {optimal_k} Clusters')
```

```
plt.xlabel('Feature 1 (e.g., SALES)')
```

```
plt.ylabel('Feature 2 (e.g., QUANTITYORDERED)')
```

```
plt.show()
```

```
# Perform hierarchical clustering using 'ward' linkage method
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```
linked = linkage(scaled_features, method='ward')
```

```
# Plot the dendrogram showing the first 4 levels (truncate_mode='level', p=4)
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```
plt.figure(figsize=(10, 7))
```

```
dendrogram(linked, orientation='top', distance_sort='descending', show_leaf_counts=True,
```

```
truncate_mode='level', p=4)
```

```

plt.title('Dendrogram for Hierarchical Clustering (Truncated to 4 Levels)')
plt.xlabel('Sample Index')
plt.ylabel('Euclidean Distance')
plt.show()

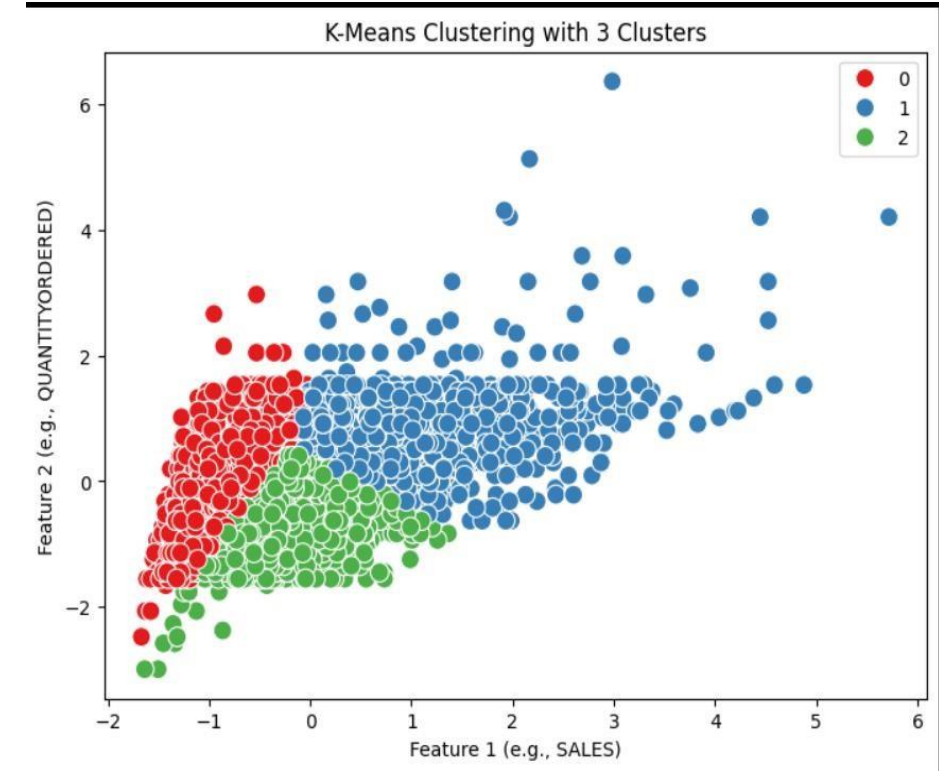
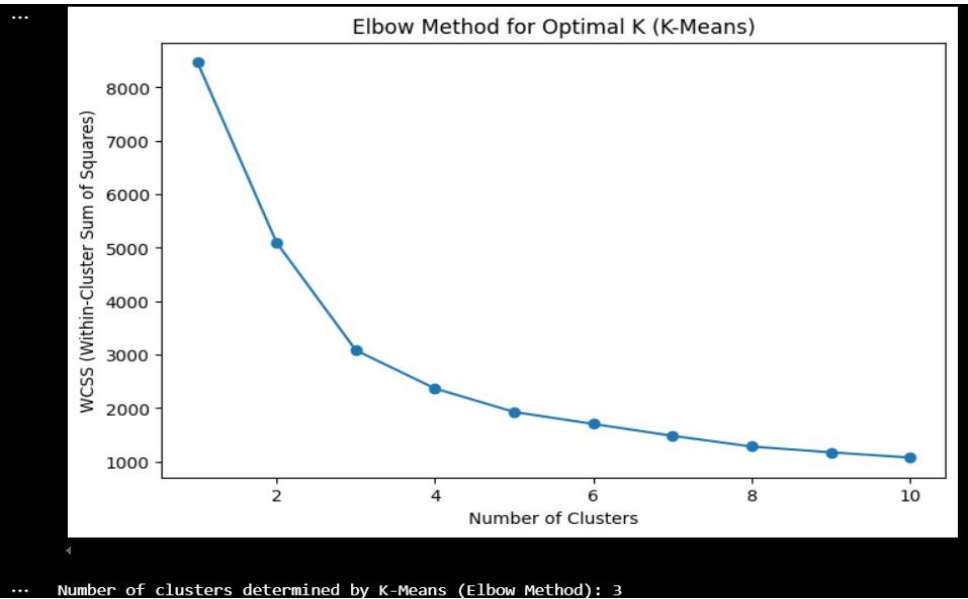
# Automatically determine the number of clusters from the dendrogram
# You can determine this by visually inspecting where the largest gap in the dendrogram
# appears
def get_optimal_clusters_from_dendrogram(linked, threshold=0.7):
    dendrogram_data = dendrogram(linked, no_plot=True)
    distances = np.diff(dendrogram_data['dcoord'], axis=1).ravel()
    threshold_distance = np.percentile(distances, threshold * 100)
    num_clusters = np.sum(distances > threshold_distance) + 1
    return num_clusters

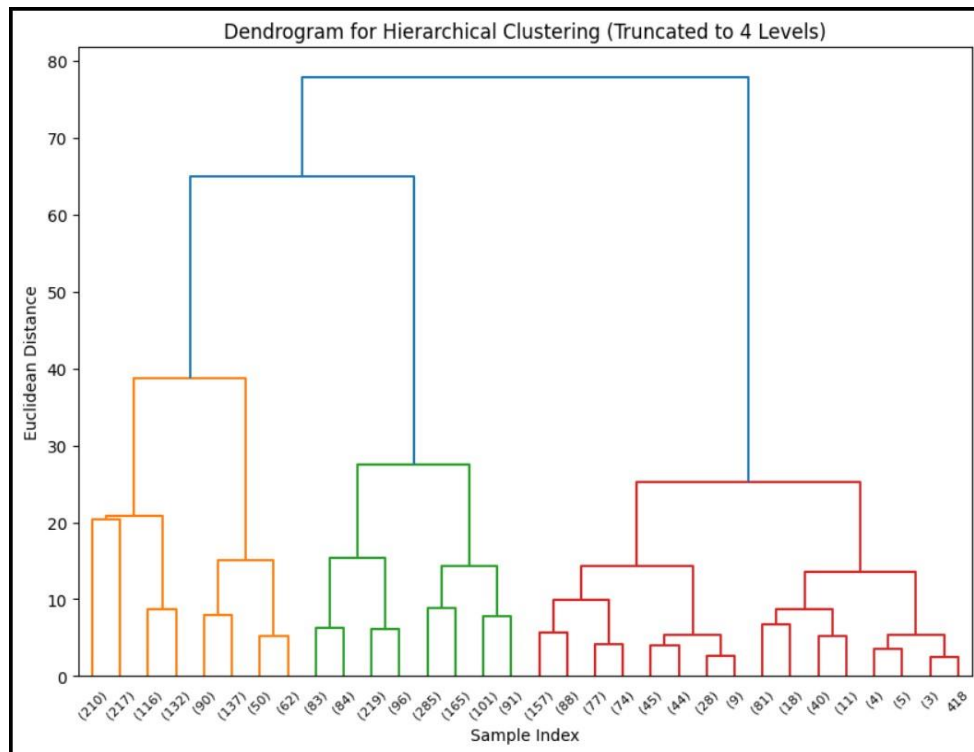
optimal_hc = get_optimal_clusters_from_dendrogram(linked)

# Apply hierarchical clustering with the chosen number of clusters
hc = AgglomerativeClustering(n_clusters=optimal_hc, metric='euclidean', linkage='ward')
hierarchical_labels = hc.fit_predict(scaled_features)

# Print the number of clusters for hierarchical clustering
print(f'Number of clusters determined by Hierarchical Clustering: {optimal_hc}')

```





Number of clusters determined by Hierarchical Clustering: 2541