

## **EXPERIMENT 5 : Unsupervised Learning Models: Clustering with K-Means and Dimensionality Reduction with PCA**

### **AIM:**

To demonstrate the application of Unsupervised Learning models, specifically K-Means clustering for grouping data points and Principal Component Analysis (PCA) for dimensionality reduction and visualization, using a suitable dataset.

### **SOURCE CODE:**

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import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.datasets import make_blobs
from sklearn.preprocessing import StandardScaler
from sklearn.cluster import KMeans
from sklearn.decomposition import PCA
from sklearn.metrics import silhouette_score
print("--- Part 1: K-Means Clustering ---")
X, y = make_blobs(n_samples=300, centers=3, cluster_std=0.60, random_state=42)
df_kmeans = pd.DataFrame(X, columns=['Feature_1', 'Feature_2'])
print("\nOriginal K-Means Dataset Head:")
print(df_kmeans.head())
wcss = []
for i in range(1, 11):
    kmeans = KMeans(n_clusters=i, init='k-means++', max_iter=300, n_init=10, random_state=42)
    kmeans.fit(X)
    wcss.append(kmeans.inertia_)

plt.figure(figsize=(10, 6))
plt.plot(range(1, 11), wcss, marker='o', linestyle='--')
plt.title('Elbow Method for Optimal K (K-Means)')
plt.xlabel('Number of Clusters (K)')
plt.ylabel('WCSS')
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plt.grid(True)
plt.show()

optimal_k = 3

kmeans = KMeans(n_clusters=optimal_k, init='k-means++', max_iter=300, n_init=10,
random_state=42)

clusters = kmeans.fit_predict(X)

df_kmeans['Cluster'] = clusters

plt.figure(figsize=(10, 8))

sns.scatterplot(x='Feature_1', y='Feature_2', hue='Cluster', data=df_kmeans,
palette='viridis', s=100, alpha=0.8, legend='full')

plt.scatter(kmeans.cluster_centers_[0], kmeans.cluster_centers_[1],
s=300, c='red', marker='X', label='Centroids')

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

plt.xlabel('Feature 1')

plt.ylabel('Feature 2')

plt.legend()

plt.grid(True)

plt.show()

silhouette_avg = silhouette_score(X, clusters)

print(f"\nSilhouette Score for K-Means (K={optimal_k}): {silhouette_avg:.3f}")

print("\n--- Part 2: Dimensionality Reduction with PCA ---")

X_pca, y_pca = make_blobs(n_samples=500, n_features=4, centers=4, cluster_std=1.0,
random_state=25)

df_pca_original = pd.DataFrame(X_pca, columns=[f'Feature_{i+1}' for i in range(X_pca.shape[1])])

df_pca_original['True_Cluster'] = y_pca

print("\nOriginal PCA Dataset Head (first 5 rows and all columns):")

print(df_pca_original.head())

print(f"Original PCA Dataset Shape: {df_pca_original.shape}")

scaler = StandardScaler()

X_pca_scaled = scaler.fit_transform(X_pca)

df_pca_scaled = pd.DataFrame(X_pca_scaled, columns=[f'Feature_{i+1}' for i in

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range(X_pca_scaled.shape[1]))
print("\nScaled PCA Dataset Head:")
print(df_pca_scaled.head())
pca = PCA(n_components=2)
principal_components = pca.fit_transform(X_pca_scaled)
df_principal_components = pd.DataFrame(data=principal_components,
columns=['Principal_Component_1', 'Principal_Component_2'])
df_principal_components['True_Cluster'] = y_pca
print("\nPrincipal Components DataFrame Head:")
print(df_principal_components.head())
explained_variance = pca.explained_variance_ratio_
print(f"\nExplained Variance Ratio for each Principal Component: {explained_variance}")
print(f"Total Explained Variance by 2 PCs: {explained_variance.sum():.3f}")
plt.figure(figsize=(10, 8))
sns.scatterplot(x='Principal_Component_1', y='Principal_Component_2', hue='True_Cluster',
data=df_principal_components, palette='Paired', s=100, alpha=0.8, legend='full')
plt.title('PCA - Dimensionality Reduction to 2 Principal Components')
plt.xlabel(f'Principal Component 1 ({explained_variance[0]*100:.2f}% variance)')
plt.ylabel(f'Principal Component 2 ({explained_variance[1]*100:.2f}% variance)')
plt.grid(True)
plt.show()
kmeans_pca = KMeans(n_clusters=4, init='k-means++', max_iter=300, n_init=10, random_state=42)
clusters_pca = kmeans_pca.fit_predict(principal_components)
df_principal_components['KMeans_Cluster_on_PCA'] = clusters_pca
plt.figure(figsize=(10, 8))
sns.scatterplot(x='Principal_Component_1', y='Principal_Component_2',
hue='KMeans_Cluster_on_PCA',
data=df_principal_components, palette='viridis', s=100, alpha=0.8, legend='full')
plt.scatter(kmeans_pca.cluster_centers_[0], kmeans_pca.cluster_centers_[1],
s=300, c='red', marker='X', label='Centroids')
plt.title('K-Means Clustering on PCA-Reduced Data')
plt.xlabel('Principal Component 1')

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plt.ylabel('Principal Component 2')
plt.legend()
plt.grid(True)
plt.show()

silhouette_avg_pca = silhouette_score(principal_components, clusters_pca)
print(f'\nSilhouette Score for K-Means on PCA-Reduced Data (K=4): {silhouette_avg_pca:.3f}')
```

## OUTPUT:

