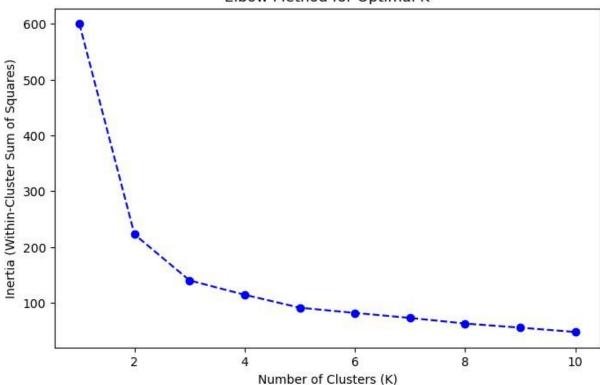
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```
In [2]: #Name: Mane Shivraj Pandurang.
        #Roll No.:37 B.E.A.I & D.S.
        # CL IV (Business Intelligence)
        #Practical No.5
In [3]: import pandas as pd
        import numpy as np
        import seaborn as sns
        import matplotlib.pyplot as plt
        from sklearn.cluster import KMeans
        from sklearn.preprocessing import StandardScaler
        from sklearn.decomposition import PCA
        from sklearn.metrics import silhouette score
        import warnings
        warnings.filterwarnings('ignore')
In [4]: # Load the Iris dataset
        iris = sns.load dataset("iris")
        print(iris.head()) # Display first 5 rows
           sepal_length sepal_width petal_length petal_width species
       0
                   5.1
                                 3.5
                                               1.4
                                                            0.2 setosa
                   4.9
                                 3.0
                                               1.4
                                                            0.2 setosa
       1
       2
                   4.7
                                3.2
                                               1.3
                                                            0.2 setosa
       3
                   4.6
                                 3.1
                                               1.5
                                                            0.2 setosa
       4
                   5.0
                                 3.6
                                               1.4
                                                            0.2 setosa
In [5]: # Drop the categorical 'species' column
        X = iris.drop(columns=['species'])
        # Standardizing the features (scaling)
        scaler = StandardScaler()
        X scaled = scaler.fit transform(X)
In [6]: # Elbow Method to determine the best number of clusters
        inertia = []
        K range = range(1, 11) # Check for K=1 to K=10
        for k in K_range:
            kmeans = KMeans(n clusters=k, random state=42, n init=10)
            kmeans.fit(X scaled)
            inertia.append(kmeans.inertia_)
        # PLot Elbow Curve
        plt.figure(figsize=(8, 5))
        plt.plot(K_range, inertia, marker='o', linestyle='--', color='b')
        plt.xlabel('Number of Clusters (K)')
        plt.ylabel('Inertia (Within-Cluster Sum of Squares)')
        plt.title('Elbow Method for Optimal K')
        plt.show()
```

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Elbow Method for Optimal K



```
In [7]: # Apply K-Means with 3 clusters
        kmeans = KMeans(n_clusters=3, random_state=42, n_init=10)
        clusters = kmeans.fit_predict(X_scaled)
        # Assign clusters to a new column in the dataset
        iris['Cluster'] = clusters
        # Debug check
        print(iris.head()) # Verify the 'Cluster' column is added
          sepal_length
                        sepal_width petal_length petal_width species
                                                                         Cluster
       0
                   5.1
                                 3.5
                                               1.4
                                                            0.2 setosa
                   4.9
                                 3.0
                                               1.4
                                                            0.2 setosa
                                                                                1
       1
       2
                   4.7
                                 3.2
                                               1.3
                                                            0.2 setosa
                                                                                1
       3
                   4.6
                                 3.1
                                               1.5
                                                            0.2 setosa
                                                                                1
```

```
In [8]: silhouette_avg = silhouette_score(X_scaled, clusters)
print(f'Silhouette Score: {silhouette_avg:.2f}')
```

1.4

0.2 setosa

1

Silhouette Score: 0.46

5.0

3.6

```
In [9]: # Apply PCA to reduce dimensions from 4D to 2D
pca = PCA(n_components=2)
X_pca = pca.fit_transform(X_scaled)

# Convert to DataFrame for visualization
pca_df = pd.DataFrame(X_pca, columns=['PC1', 'PC2'])
pca_df['Cluster'] = clusters # Assign clusters
# Scatter plot of PCA components with clusters
plt.figure(figsize=(8, 6))
```

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```
sns.scatterplot(x=pca_df['PC1'], y=pca_df['PC2'], hue=pca_df['Cluster'],
palette='viridis', s=100)
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.title('K-Means Clustering (PCA Reduced)')
plt.legend(title='Cluster')
plt.show()
```

K-Means Clustering (PCA Reduced)

