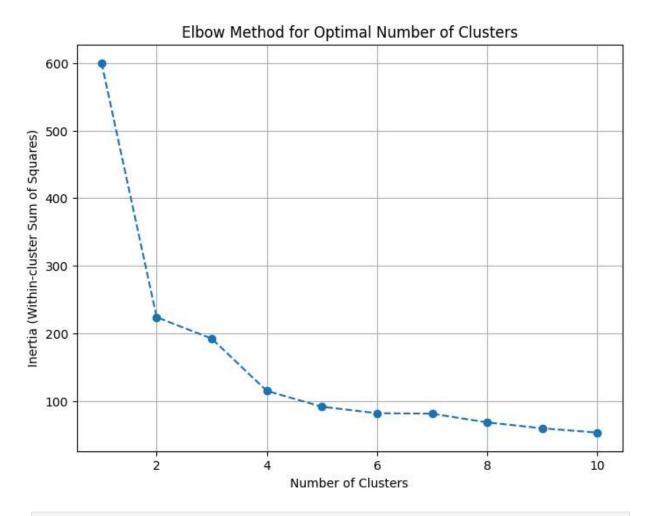
```
In [2]: import pandas as pd
         import numpy as np
         import matplotlib.pyplot as plt
         from sklearn.cluster import KMeans
         from sklearn.preprocessing import StandardScaler
         from sklearn.decomposition import PCA
In [4]: # Load the Iris dataset
         df = pd.read_csv("Iris.csv")
In [6]: # Select features (attributes) for clustering (e.g., sepal length, sepal width, pet
         X = df.iloc[:, 1:-1] # Exclude the first column (id) and the last column (species)
In [8]: # Standardize the feature matrix (important for K-Means)
         scaler = StandardScaler()
         X_scaled = scaler.fit_transform(X)
In [10]: # Determine the optimal number of clusters using the elbow method
         inertia = []
         for k in range(1, 11):
             kmeans = KMeans(n_clusters=k, random_state=42)
             kmeans.fit(X scaled)
             inertia.append(kmeans.inertia_)
In [11]: print(inertia)
        [600.0, 223.73200573676345, 192.03717409190028, 114.68221609937967, 91.295444740669
        8, 81.76026132860625, 80.98238131032988, 68.08623905064636, 59.38528882045367, 52.98
        999721015859]
In [14]: # Plot the elbow curve
         plt.figure(figsize=(8, 6))
         plt.plot(range(1, 11), inertia, marker='o', linestyle='--')
         plt.xlabel('Number of Clusters')
         plt.ylabel('Inertia (Within-cluster Sum of Squares)')
         plt.title('Elbow Method for Optimal Number of Clusters')
         plt.grid()
         plt.show()
```



In []:	:	
In []:	*	