

```
In [1]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
import seaborn as sns
from sklearn.decomposition import PCA

import warnings
warnings.filterwarnings('ignore')
```

```
In [2]: # Load Iris dataset directly from seaborn
df = sns.load_dataset("iris")

# Display first 5 rows
df
```

```
Out[2]:   sepal_length  sepal_width  petal_length  petal_width  species
      0           5.1         3.5          1.4         0.2    setosa
      1           4.9         3.0          1.4         0.2    setosa
      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
     ...
     145          6.7         3.0          5.2         2.3  virginica
     146          6.3         2.5          5.0         1.9  virginica
     147          6.5         3.0          5.2         2.0  virginica
     148          6.2         3.4          5.4         2.3  virginica
     149          5.9         3.0          5.1         1.8  virginica
```

150 rows × 5 columns

```
In [3]: # Select features (attributes) for clustering (e.g., sepal_length, sepal_width, petal_length, petal_width)
X = df.iloc[:, 1:-1] # Exclude the first column (id) and the last column (species)
```

```
In [18]: X
```

```
Out[18]:   sepal_width  petal_length  petal_width
      0           3.5          1.4         0.2
      1           3.0          1.4         0.2
      2           3.2          1.3         0.2
      3           3.1          1.5         0.2
      4           3.6          1.4         0.2
     ...
     145          3.0          5.2         2.3
     146          2.5          5.0         1.9
     147          3.0          5.2         2.0
     148          3.4          5.4         2.3
     149          3.0          5.1         1.8
```

150 rows × 3 columns

```
In [4]: # Standardize the feature matrix (important for K-Means)
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
```

```
In [20]: # 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_)
```

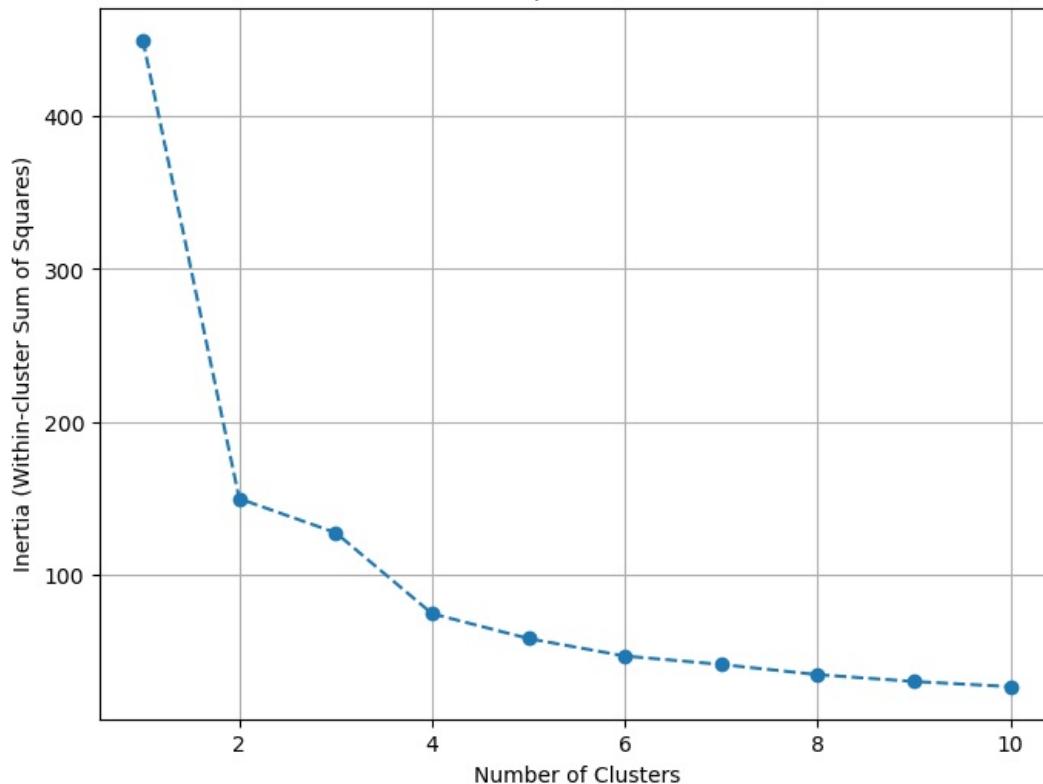
```
In [22]: inertia
```

```
Out[22]: [450.0,  
 149.5644912164707,  
 127.31742412220018,  
 74.24391953311728,  
 57.94932812049257,  
 46.507154272195756,  
 40.97340775686134,  
 34.29606921621072,  
 29.819669802832685,  
 26.442321160268044]
```

```
In [7]: # 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()
```

Elbow Method for Optimal Number of Clusters



```
In [ ]:
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