

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
from sklearn.model_selection import train_test_split
from sklearn.cluster import KMeans
from scipy.cluster.hierarchy import dendrogram, linkage
```

```
country_data= pd.read_csv('Country-data.csv')
print(country_data.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 167 entries, 0 to 166
Data columns (total 10 columns):
#   Column      Non-Null Count  Dtype
---  ---
0   country     167 non-null    object
1   child_mort  167 non-null    float64
2   exports     167 non-null    float64
3   health      167 non-null    float64
4   imports     167 non-null    float64
5   income      167 non-null    int64
6   inflation   167 non-null    float64
7   life_expec  167 non-null    float64
8   total_fer   167 non-null    float64
9   gdp         167 non-null    int64
dtypes: float64(7), int64(2), object(1)
memory usage: 13.2+ KB
None
```

```
print(country_data.isnull().sum()) ##Since there are no null alues we dont have any NAN values to remove
```

```
country      0
child_mort   0
exports      0
health       0
imports      0
income       0
inflation    0
life_expec   0
total_fer    0
gdp          0
dtype: int64
```

```
country_data.head()
```

```
country  child_mort  exports  health  imports  income  inflation  life_expec  tc
0  Afghanistan      90.2     10.0    7.58     44.9    1610         9.44        56.2
1    Albania        16.6     28.0    6.55     48.6    9930         4.49        76.3
2    Algeria        27.3     38.4    4.17     31.4   12900        16.10        76.5
3    Angola        119.0     62.3    2.85     42.9    5900        22.40        60.1
Antigua and
```

We will check if there are any duplicates on the basis of country to check if there are any double entries of any country; From the result it is clear that we dont have any duplicate entries

```
duplicates_in_one_column = len(country_data['country']) - len(country_data['country'].drop_duplicates())
print(f"Number of duplicates on the basis of country column: {duplicates_in_one_column}")
```

```
Number of duplicates on the basis of country column: 0
```

Now we will Stadardise the data

```
columns_to_normalize = country_data.select_dtypes(include=['float64', 'int64']).columns
```

```
train_X, test_X=train_test_split(country_data[columns_to_normalize],test_size=0.3, random_state=1)
```

```
scaler=StandardScaler()
```

```
scaler.fit(train_X)
```

```
StandardScaler()
```

```
train_X=scaler.transform(train_X)
```

```
test_X=scaler.transform(test_X)
```

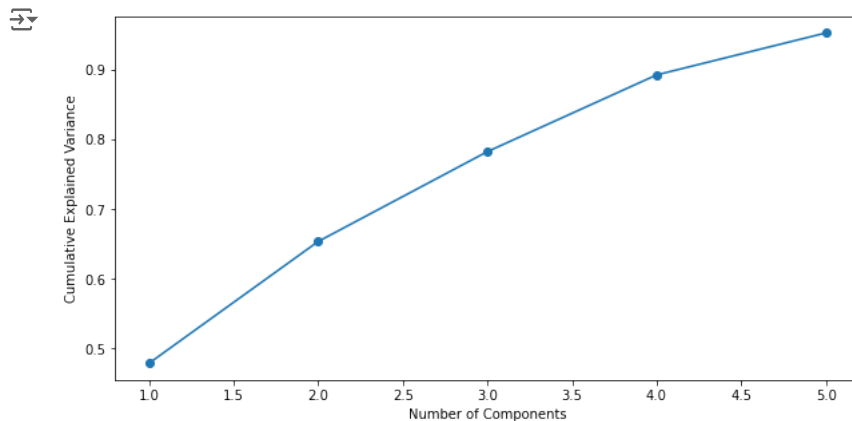
```
train_X = pd.DataFrame(train_X, columns=columns_to_normalize)
test_X= pd.DataFrame(test_X, columns=columns_to_normalize)
```

Now, PCA(Principle Component Analysis Technique) is applied to reduce the dimensionality of the data while retaining 90% of the variance as it will help to simplify the complexity of the data and aided in visualization.

```
# Initialize PCA, retaining 90% of variance
pca = PCA(n_components=0.9)
```

```
# Fit and transform the normalized data
X_train_pca = pca.fit_transform(train_X)
X_test_pca = pca.transform(test_X)
```


```
# Plot the explained variance to understand the components
plt.figure(figsize=(10, 5))
plt.plot(range(1, len(pca.explained_variance_ratio_) + 1), pca.explained_variance_ratio_.cumsum(), marker='o')
plt.xlabel('Number of Components')
plt.ylabel('Cumulative Explained Variance')
plt.show()
```

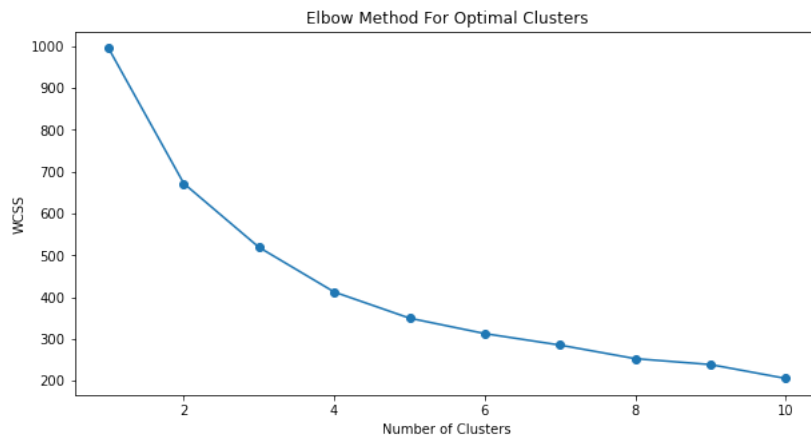


I used K-mean Clustering Elbow method to get most optimal number of clusters, according to which country will be labelled. I assumed the most optimal clustered as 4.

```
wcss = []
for i in range(1, 11):
    kmeans = KMeans(n_clusters=i, random_state=42)
    kmeans.fit(X_train_pca)
    wcss.append(kmeans.inertia_)
```

```
plt.figure(figsize=(10, 5))
plt.plot(range(1, 11), wcss, marker='o')
plt.xlabel('Number of Clusters')
plt.ylabel('WCSS')
plt.title('Elbow Method For Optimal Clusters')
plt.show()
```

 C:\Users\257380\anaconda3\lib\site-packages\sklearn\cluster\\_kmeans.py:1036: UserWarning  
warnings.warn()



# Assuming the elbow method suggests 4 clusters

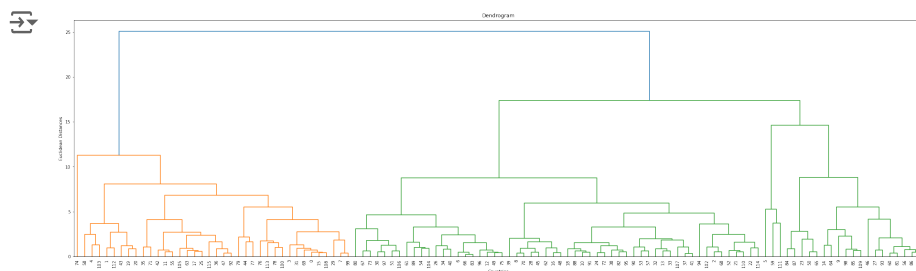
```
kmeans = KMeans(n_clusters=4, random_state=42)
train_clusters = kmeans.fit_predict(X_train_pca)
test_clusters = kmeans.predict(X_test_pca)
```

```
# Add the cluster labels to the original data
country_data.loc[train_X.index, 'Cluster'] = train_clusters
country_data.loc[test_X.index, 'Cluster'] = test_clusters
```

Then, Hierarchical clustering is performed using ward linkage which provided a dendrogram that helped to visualize nested relationships between countries.

```
# Using 'ward' linkage to minimize variance within clusters
Z = linkage(X_train_pca, method='ward')

plt.figure(figsize=(35, 10))
dendrogram(Z, labels=train_X.index, leaf_rotation=90, leaf_font_size=10)
plt.title('Dendrogram')
plt.xlabel('Countries')
plt.ylabel('Euclidean Distances')
plt.show()
```



```
from sklearn.cluster import AgglomerativeClustering

# Fit Agglomerative Clustering
hc = AgglomerativeClustering(n_clusters=4, affinity='euclidean', linkage='ward')
train_hc_clusters = hc.fit_predict(X_train_pca)

# Summarize clusters
country_data.loc[train_hc_clusters, 'HC_Cluster'] = train_hc_clusters

# Summarize clusters
cluster_summary = country_data.groupby('Cluster').mean()
print(cluster_summary)

# Assuming lower GDP per capita, higher infant mortality, and lower life expectancy indicate higher need
priority_clusters = cluster_summary.sort_values(by=['gdp', 'life_expec', 'child_mort'], ascending=[True, False, True])

# Extract countries in the highest priority cluster
priority_cluster_index = priority_clusters.index[0]
priority_countries = country_data[country_data['Cluster'] == priority_cluster_index]

print("Recommended Countries for Aid Allocation according to highest clusters that is Child Mortality(child_mort) ,GDP(gdp) and\n Lower Life Expectancy(life_expec):")
print(priority_countries[['country', 'gdp', 'life_expec', 'child_mort', 'Cluster']])
```

	child_mort	exports	health	imports	income	inflation \
Cluster						
0.0	45.458621	38.016897	5.793448	42.451724	14382.896552	10.979379
1.0	38.708333	42.750483	6.919167	47.247765	18897.050000	7.116533
2.0	40.450000	29.900000	7.660000	36.950000	17680.000000	10.165000
3.0	42.932000	40.175200	7.552800	54.508000	12959.440000	5.381160

	life_expec	total_fer	gdp	HC_Cluster
Cluster				
0.0	70.703448	3.217241	10571.827586	0.862069
1.0	70.435000	2.851500	15242.150000	1.750000
2.0	71.550000	3.220000	17505.000000	3.000000
3.0	68.528000	3.032800	9500.680000	0.760000

Recommended Countries for Aid Allocation according to highest clusters that is Child Mortality(child\_mort) ,GDP(gdp) and Lower Life Expectancy(life\_expec):

	country	gdp	life_expec	child_mort	Cluster
0	Afghanistan	553	56.2	90.2	3.0
5	Argentina	10300	75.8	14.5	3.0
14	Belarus	6030	70.4	5.5	3.0
15	Belgium	44400	80.0	4.5	3.0
21	Botswana	6350	57.1	52.5	3.0
24	Bulgaria	6840	73.9	10.8	3.0
25	Burkina Faso	575	57.9	116.0	3.0
31	Central African Republic	446	47.5	149.0	3.0
33	Chile	12900	79.1	8.7	3.0
35	Colombia	6250	76.4	18.6	3.0
48	El Salvador	2990	74.1	19.2	3.0
56	Gambia	562	65.5	80.3	3.0
60	Greece	26900	80.4	3.9	3.0
64	Guinea-Bissau	547	55.6	114.0	3.0
65	Guyana	3040	65.5	37.6	3.0
67	Hungary	13100	74.5	6.0	3.0
73	Ireland	48700	80.4	4.2	3.0
81	Kiribati	1490	60.7	62.7	3.0
85	Latvia	11300	73.1	7.8	3.0
86	Lebanon	8860	79.8	10.3	3.0
93	Madagascar	413	60.8	62.2	3.0
98	Malta	21100	80.3	6.8	3.0
101	Micronesia, Fed. Sts.	2860	65.4	40.0	3.0
106	Mozambique	419	54.5	101.0	3.0
109	Nepal	592	68.3	47.0	3.0

Start coding or generate with AI.