## Question1.1

#### Here is the code

```
import pandas as pd
import numpy as np
from sklearn.cluster import AgglomerativeClustering
from sklearn.preprocessing import StandardScaler
from sklearn.impute import SimpleImputer
url = "https://archive.ics.uci.edu/ml/machine-learning-databases/auto-mpg/auto-mpg.data"
columns = ['mpg', 'cylinders', 'displacement', 'horsepower', 'weight', 'acceleration', 'model_year', 'origin', 'car_name']
df = pd.read_csv(url, sep='\s+', names=columns) # Use sep='\s+' instead of delim_whitespace
continuous_cols = ['mpg', 'cylinders', 'displacement', 'horsepower', 'weight', 'acceleration', 'model_year']
df['horsepower'] = pd.to_numeric(df['horsepower'], errors='coerce')
X = df[continuous cols].copy()
imputer = SimpleImputer(strategy='mean')
X imputed = imputer.fit transform(X)
X_imputed = pd.DataFrame(X_imputed, columns=continuous_cols)
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X_imputed)
clustering = AgglomerativeClustering(n clusters=3, linkage='average', metric='euclidean') # Use metric instead of affinit
labels = clustering.fit_predict(X_scaled)
df['cluster'] = labels
cluster_stats = df.groupby('cluster')[continuous_cols].agg(['mean', 'var'])
print("Cluster Statistics (Mean and Variance):")
print(cluster_stats)
origin_stats = df.groupby('origin')[continuous_cols].agg(['mean', 'var'])
print("\nOrigin Class Statistics (Mean and Variance):")
print(origin_stats)
crosstab = pd.crosstab(df['cluster'], df['origin'])
print("\nCrosstab of Cluster vs Origin:")
print(crosstab)
print("\nAnalysis:")
if crosstab.max().max() / crosstab.sum().sum() > 0.5:
    print("There is a clear relationship between cluster assignments and origin labels.")
else:
    print("There is no clear relationship between cluster assignments and origin labels.")
```

Here is the output

horsepower weight acceleration \	Cluster	Statistics	(Mean and					
Cluster 0		mpg	3			displacemen	it	
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	2	45366	950					
	201	0 4	3.00					

Analysis:
There is no clear relationship between cluster assignments and origin labels.

From the image, we can see that Cluster 1 and Cluster 2 have a strong association with the origin categories, while Cluster 0 is a mixed category primarily containing American and Japanese cars but with features closer to European cars. The hierarchical clustering did not fully distinguish between origin=1 (American) and origin=3 (Japanese) vehicles.

Therefore, my conclusion is:

There is a partial association between cluster assignments and class labels, but no particularly strong or clear relationship.

# Question1.2

#### Here is my code

```
import pandas as pd
import numpy as np
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import silhouette score
try:
    from sklearn.datasets import load boston
    boston = load_boston()
    X = pd.DataFrame(boston.data, columns=boston.feature names)
    dataset_name = "Boston"
except ImportError:
    from sklearn.datasets import fetch_california_housing
    boston = fetch california housing()
    X = pd.DataFrame(boston.data, columns=boston.feature names)
    dataset name = "California Housing"
scaler = StandardScaler()
X scaled = scaler.fit transform(X)
silhouette scores = []
for k in range(2, 7):
    kmeans = KMeans(n clusters=k, n init=10, random state=42) # Explicitly set n init
    labels = kmeans.fit_predict(X_scaled)
    score = silhouette score(X scaled, labels, sample size=1000, random state=42) # Limit sample size
    silhouette_scores.append((k, score))
optimal_k = max(silhouette_scores, key=lambda x: x[1])[0]
print(f"Optimal number of clusters for {dataset name}: {optimal k}")
print(f"Silhouette scores for k=2 to 6: {silhouette_scores}")
kmeans_optimal = KMeans(n_clusters=optimal_k, n_init=10, random_state=42)
labels optimal = kmeans optimal.fit predict(X scaled)
X['cluster'] = labels_optimal
cluster means = X.groupby('cluster').mean()
print(f"\nMean values for each cluster in {dataset name}:")
print(cluster_means)
centroids = pd.DataFrame(scaler.inverse_transform(kmeans_optimal.cluster_centers_), columns=boston.feature_names)
print(f"\nCentroid coordinates for {dataset name}:")
print(centroids)
print(f"\nDifference between cluster means and centroids in {dataset name}:")
print(cluster means - centroids)
```

### Here is the output

```
Optimal number of clusters for California Housing: 2
Silhouette scores for k=2 to 6: [(2, 0.3368433874602563), (3, 0.3368433874602563), (4, 0.3026463930769773), (5, 0.32344236329026266), (6, 0.3232918404758
Mean values for each cluster in California Housing:
         MedInc HouseAge AveRooms AveBedrms Population AveOccup \
cluster
       3.918104 28.412773 5.225159 1.075685 1532.241745 3.098100
      3.805276 28.952057 5.710036 1.125614 1278.279590 3.032817
        Latitude Longitude
cluster
0
      33.945736 -118.010096
      37.956526 -121.719940
Centroid coordinates for California Housing:
    MedInc HouseAge AveRooms AveBedrms Population AveOccup Latitude \
0 3.918418 28.414681 5.225372 1.075692 1532.198896 3.097956 33.945450
1 3.804868 28.949303 5.709630 1.125593 1278.397166 3.033030 37.955996
   Longitude
0 -118,009704
1 -121.719626
Difference between cluster means and centroids in California Housing:
         MedInc HouseAge AveRooms AveBedrms Population AveOccup \
cluster
0
     -0.000315 -0.001908 -0.000213 -0.000007 0.042849 0.000144
      0.000407 0.002754 0.000406 0.000021 -0.117576 -0.000213
1
       Latitude Longitude
cluster
       0.000286 -0.000392
1
       0.000530 -0.000315
```

Based on the output results, it can be concluded that the optimal k is 2, with the two clusters primarily reflecting differences in housing characteristics between the northern and southern regions of California.

Cluster 0 is concentrated in the southern region,

Cluster 1 is concentrated in the northern region.

The differences between the mean values and centroid coordinates are very minor, indicating that the K-Means clustering results are reliable. The mean values and centroid coordinates are almost numerically idendical.

## Question1.3

### Here is my code

```
import os
import pandas as pd
import numpy as np
from sklearn.datasets import load_wine
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import homogeneity_score, completeness_score
os.environ["OMP_NUM_THREADS"] = "1"
wine = load_wine()
X = pd.DataFrame(wine.data, columns=wine.feature names)
y = wine.target
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
kmeans = KMeans(n clusters=3, n init=10, random state=42) # Explicitly set n init
labels = kmeans.fit_predict(X_scaled)
homogeneity = homogeneity_score(y, labels)
completeness = completeness_score(y, labels)
print("Homogeneity Score:", homogeneity)
print("Completeness Score:", completeness)
Here is the output
     Homogeneity Score: 0.8788432003662366
     Completeness Score: 0.8729636016078731
```

The homogeneity and completeness scores are 0.878 and 0.873, respectively, indicating that the K-Means clustering performs well on the Wine dataset.

Using homogeneity and completeness together provides a comprehensive evaluation of how well the clustering results align with the true labels. The current high scores demonstrate that K-Means effectively captures the class structure of the Wine dataset.