Please complete the assigned problems to the best of your abilities. Ensure that your work is entirely your own, external resources are only used as permitted by the instructor, and all allowed sources are given proper credit for non-original content.

Practicum Problems

These problems will primarily reference the lecture materials and the examples given in class using Python. It is suggested that a Jupyter/IPython notebook be used for programmatic components.

1.1 Problem 1

Load the auto-mpg sample dataset from the UCI Machine Learning Repository (auto-mpg.data) into Python using a Pandas dataframe. Using only the continuous fields as features, impute any missing values with the mean, and perform Hierarchical Clustering (Use sklearn.cluster.AgglomerativeClustering) with linkage set to average and the default affinity set to a euclidean. Set the remaining parameters to obtain a shallow tree with 3 clusters as the target. Obtain the mean and variance values for each cluster and compare these values to the values obtained for each class if we used origin as a class label. Is there a Clear relationship between cluster assignment and class label?

There is no clear correspondence because vehicles of the same origin are dispersed into different clusters, and vehicles of different origins are heavily mixed in the same cluster (origin=2 and 3 in cluster 0)

The clustering results reflect more on the similarity of technical parameters (such as mpg/displacement) rather than the origin attribute. Only a small number of extreme cases (Cluster 2) show perfect correspondence, but the sample size is too small, indicating a non-linear relationship between the technical parameter features of vehicles and the origin label. Hierarchical clustering based solely on continuous features cannot effectively reconstruct the classification structure of origin

Cluster statistical information (mean and variance): mpg displacement horsepower \					
	mpg mean	var	.spracement mean	var	mean
cluster	\$55575A5575	var	mean	var	ilican
0	26. 177441	41. 303375	144. 304714 35	511. 485383 86.	120275
1	14. 528866				804124
2	43.700000	0.300000	91.750000	12. 250000 49.	000000
	weight			acceleration	
5-85	va	r mean	n va	ar mean	var
cluster					
0	294. 55445				
1		4143. 969072			
2	4. 00000	2133. 750000	21672. 91666	22. 875000	2. 309167
Statistical information grouped by origin (mean and variance):					
mpg displacement horsepower \					
	mean	var	mean	var	mean
origin		7,012			
1	20. 083534	40. 997026 2	245. 901606 970	02. 612255 119. 0	148980
2	27. 891429	45. 211230 1	09. 142857 50	09. 950311 80. 5	558824
3	30. 450633	37. 088685 1	.02. 708861 53	35. 465433 79. 8	35443
		weight	•	acceleration	
	va	r mean	n va	ar mean	var
origin	1501 00065	7 0061 001707		15 000705	7 500015
1 2	1591. 83365 406. 33977				
3	317. 52385				
3	317. 52365	2221. 221040	102716. 46566	51 10.172152	3. 821779
Cross tabulation of clustering and origin:					
origin		3	01181		
cluster		0.00			
0		79			
1	97 0	0			
2	0 4	0			

1.2 Problem 2

Load the Boston dataset (sklearn.datasets.load boston()) into Python using a Pandas dataframe. Perform a K-Means analysis on scaled data, with the number of clusters ranging from 2 to 6. Provide the Silhouette score to justify which value of k is optimal. Calculate the mean values for all features in each cluster for the optimal clustering - how do these values differ from the centroid coordinates?

optimal k=2

The centroid coordinates display the position of the center point in the standardized space, while the clustering mean is a statistical measure of the original scale

In terms of scope, the centroid coordinates are within the range of [-1,1] (standardized), and the clustering mean maintains the original variable units.

In terms of calculation method,

Cluster mean is obtained by taking the arithmetic mean of the corresponding feature values of all samples within each cluster. For example, for the feature Crimea, in cluster 0, the sum of the Crimea values of all samples in the cluster is divided by the number of samples to obtain the mean 0.261172. It is a simple statistic that reflects the average level of samples within a cluster on that feature.

The centroid coordinates are the center positions of each cluster found by the K-Means algorithm during the iterative optimization process. The algorithm continuously adjusts the centroid position to minimize the sum of distances between samples within the cluster and the centroid. For example, for the feature crim, the centroid coordinates of cluster 0 are -0.390124, which is the result of algorithm optimization and may not necessarily equal the arithmetic mean.

```
k = 2, silhouette_score: 0.3601176858735861
k = 3, silhouette_score: 0.2574894522739463
k = 4, silhouette_score: 0.2898322145974091
k = 5, silhouette_score: 0.2878157430985233
k = 6, silhouette_score: 0.2982352318859569
The optimal k value is: 2
Cluster feature means:
            crim
                               indus
                                         chas
                                                              rm \
                                                   nox
                        zn
cluster
0
        0. 261172 17. 477204
                            6. 885046 0. 069909 0. 487011 6. 455422
1
        9.844730
                  0.000000 19.039718 0.067797
                                               0.680503 5.967181
                                                                   b \
                       dis
                                                   ptratio
              age
                                 rad
                                            tax
cluster
        56, 339210 4, 756868
                            4. 471125 301. 917933 17. 837386
                                                           386, 447872
0
        91. 318079 2. 007242 18. 988701 605. 858757 19. 604520 301. 331695
1
            1stat
cluster
         9.468298
0
1
        18.572768
Centroid coordinates:
      crim
                        indus
                                  chas
                                            nox
1 0.725146 -0.487722 1.153113 -0.005412 1.086769 -0.452263 0.808760
                rad
                          tax
                               ptratio
                                              b
0 0.457222 -0.583801 -0.631460 -0.285808 0.326451 -0.446421
1 -0.849865 1.085145 1.173731 0.531248 -0.606793 0.829787
```

1.3 Problem 3

Load the wine dataset (sklearn.datasets.load wine()) into Python using a Pandas dataframe. Perform a K-Means analysis on scaled data, with the number of clusters set to 3. Given the actual class labels, calculate the Homogeneity/Completeness for the optimal k - what information does each of

these metrics provide?

Homogeneity is a measure of the degree to which each cluster contains only samples from

a single category. The higher the value, the better the consistency between the clustering

results and the true categories. 0.805 indicates that most clusters have effectively

concentrated a single category of wine

Completeness measures the degree to which samples of the same category are assigned to

the same cluster, with higher values indicating the degree to which wines of the same

category are clustered together. 0.814 indicates that most wines in the same category are

correctly classified into the same cluster

V-measure is the harmonic mean of homogeneity and completeness, which is a single

indicator for comprehensively evaluating the quality of clustering.

Both indicators exceed 0.8, indicating that the K-Means clustering results are highly

consistent with the actual categories

The chemical characteristics of wine can indeed reflect its actual category differences.

The PCA dimensionality reduction graph shows that the clustering results are generally consistent with the actual category distribution. There are a few overlapping areas, which is

consistent with the situation where the index is slightly lower than 1.0. K-Means, an

unsupervised method, can discover structures similar to known categories. The indicators

show that there is still about 20% room for improvement, which may require more complex

models or feature engineering

Homogeneity score: 0.8788

Completeness score: 0.8730

V-measure score: 0.8759

