

11813184

Data Mining

Assignment Five

10 points

Due: Thursday, November 16 @ 11:50 PM

Question 1 (3 points). Use the similarity matrix in the table below to perform complete link hierarchical clustering *by hand (do not use Python)*. Show your results by drawing a dendrogram. The dendrogram should clearly show the order in which the points are merged.

	p1	p2	р3	p4	p5
p1	1.00	0.10	0.41	0.55	0.35
p2	0.10	1.00	0.64	0.47	0.98
р3	0.41	0.64	1.00	0.44	0.85
p4	0.55	0.47	0.44	1.00	0.76
p5	0.35	0.98	0.85	0.76	1.00

Question 2 (4 points). In this problem, you will perform K-means clustering by hand (do not use Python), with K = 2, on a small example with 6 observations and 2 features. The observations are as follows:

OBSERVATION	X1	X2
1	1	4
2	1	3
3	0	4
4	5	1
5	6	2
6	4	0

- (a) Plot the observations by hand
- (b) Arbitrarily partition the observations into 2 groups. Report the cluster label for each observation.
- (c) Compute the centroid for each cluster.
- (d) Assign each observation to the centroid to which it is closest, in terms of Euclidean distance. Report the cluster label for each observation.
- (e) Repeat (c) and (d) until the answers obtained stop changing.
- (f) In your plot from (a), color the observations according to the cluster labels obtained.

Question 3 (3 points). Apply K-means clustering on the Iris dataset using Python and visualize cluster assignments from K = 2 to K = 5 clusters. What value of K seems the best? You can include the following statements to load the iris data:

from sklearn import datasets
iris = datasets.load_iris()

Good luck!

Question :

To perform complete link hierarchical clustering, we should consider each point in the similarity matrix as the initial centroids. Each stage where there are similar clusters (the closest), we merged them. We form the K clusters by assigning all the points to the closest centroid, after which, we would recalculate the centroid of each cluster, and keep doing this until the centroids do not change. We would do this iteratively until all points in the matrix are accounted for fall into a single cluster.

1) For this question, the initial step would be to look at the proximity matrix and use the distance formula d=1-s, where d is the distance and s is the similarity measure. We would merge them as cluster "p2 union p5", which can just be written as "25", getting p25 as the smallest value.

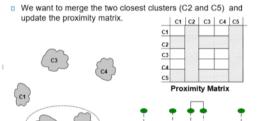
	p1	p2	р3	p4	p5
p1	d(1) = 0				
p2	d(0.10) = 0.90	d(1) = 0			
р3	d(0.41) = 0.59	d(0.64) = 0.36	d(1) = 0		
p4	d(0.55) = 0.45	d(0.47) = 0.53	d(0.44) = 0.56	d(1) = 0	
p5	d(0.35) = 0.65	d(0.98) = 0.02	d(0.85) = 0.15	d(0.76) = 0.24	d(1) = 0

2) After we do that, we calculate the similarity of each point respective to that "25" cluster. It depends on what technique we want to use for inter-cluster similarity, but for this example, we use MAX for complete linkage, which calculates the maximum distance between any pair of data points, one from each of the two clusters.

These two steps would be repeated. We find the similarity of a point with respect to the cluster we created. See below.

	p25	p1	р3	p4
p25	d(1) = 0			

- Select K points as the initial centroids.
- 2: repeat
- Form K clusters by assigning all points to the closest centroid.
- Recompute the centroid of each cluster.
- 5: until The centroids don't change



DETROIT MERCY

p1	max(d(0.10), d(0.35)) = 0.90	d(1) = 0		
р3	max(d(0.64), d(0.85)) = 0.36	d(0.41) = 0.59	d(1) = 0	
p4	max(d(0.47), d(0.76)) = 0.53	d(0.55) = 0.45	d(0.44) = 0.56	d(1) = 0

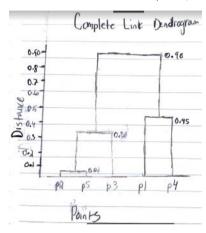
The smallest value we have in this matrix is 0.36. From that, we can say that p3 is clustered with original cluster "25" to now be "235". We repeat the process again with "235".

	p235	p1	p4
p235	d(1) = 0		
p1	max(0.90, 0.59) = 0.90	d(1) = 0	
p4	max(0.53, 0.56) = 0.56	d(0.55) = 0.45	d(1) = 0

The smallest value we have in this matrix is 0.45 from p1 x p4. From that, we can say that they are clustered as "14".

	p235	p14
p235	d(1) = 0	
p14	max(0.90, 0.56) = 0.90	d(1) = 0

The next smallest value and since it is the only one left, we have 0.90 from p235 x p14. See below for the complete link dendrogram.

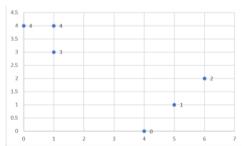


Question 2

Consider K = 2 initial clusters, see below.

Part A:

Observation X1		X2
1	1	4
2	1	3
3	0	4
4	5	1
5	6	2
6	1	0



Used a scatterplot in Excel to create this.

Part B: Let's arbitrarily assign Cluster 1: {1, 2, 3} and Cluster 2: {4, 5, 6}.

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Cluster	Centroid	Centroid		
	x1	x2		
1	((1+1+0)/3) = 0.6667	((4+3+4)/3) = 3.6667		
2 ((5+6+4)/3) = 5		((1+2+0)/3) = 1		

Cluster 1 centroid: (x1, x2) Cluster 2 centroid: (x1, x2). Cluster 1 centroid: (0.67, 3.67) Cluster 2 centroid: (5, 1).

Euclidean formula can be seen below.

$$d = \sqrt{(x_2 - x_1)^2 + (y_2 - y_1)^2}$$

Observation	Cluster 1	Cluster 2	
1	SQRT((3.67-4)^2+(0.67-1)^2) = 0.4666	SQRT((1-4)^2+(5-1)^2) = 5	

2	SQRT((3.67-3)^2+(0.67-1)^2) = 0.7468	SQRT((1-3)^2+(5-1)^2) = 4.4721
3	SQRT((3.67-4)^2+(0.67-0)^2) = 0.7468	SQRT((1-4)^2+(5-0)^2) = 5.8309
4	SQRT((3.67-1)^2+(0.67-5)^2) = 5.0870	SQRT((1-1)^2+(5-5)^2) = 0
5	SQRT((3.67-2)^2+(0.67-6)^2) = 5.5854	SQRT((1-2)^2+(5-6)^2) = 1.4142
6	SQRT((3.67-0)^2+(0.67-4)^2) = 4.9555	SQRT((1-0)^2+(5-4)^2) = 1.4142

Observation 1 is closer to Cluster 1.

Observation 2 is closer to Cluster 1.

Observation 3 is closer to Cluster 1.

Observation 4 is closer to Cluster 2.

Observation 5 is closer to Cluster 2.

Observation 6 is closer to Cluster 2.

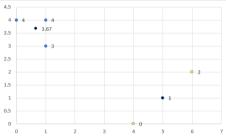
No need to update the clusters.

Part E:

Because each observation is mapped to the appropriate cluster, and that there are no additional observations for any of these clusters to reallocate to, the answers stop changing. Fortunately, my selection of the correct observations was inferred upon through the aid of the visual scatterplot.

Part F:

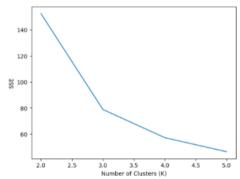
Observation	X1	X2
1	1	4
2	1	3
3	0	4
4	5	1
5	6	2
6	4	0
7	0.67	3.67
Q	5	1



Used a scatterplot in Excel to create this.

* Note: Not able to see Observation 4 due to centroid being right on top of it.

Question 3



The elbow method analysis suggests that the optimal value of K for the given dataset is K=3. The "elbow point" in the sum of squared distances (SSE) plot indicates that adding more clusters beyond K=3 does not significantly reduce the SSE. Therefore, K=3 provides a reasonable balance between capturing the underlying structure in the data and avoiding overcomplexity, making it the best choice in this context.