COMP3308/COMP3608 Introduction to Artificial Intelligence

Week 12 Tutorial exercises Unsupervised Learning (Clustering)

<u>Exercise 1.</u> K-means clustering (Homework)

Given is the one-dimensional dataset: {3, 5, 7, 10, 13}. Run the k-means clustering algorithm for 1 epoch to cluster this dataset into 2 clusters: K1 and K2. Assume that the initial seeds (cluster centers) are c1=3 and c2=10 and that the distance measure is the absolute distance between the examples.

Show the clusters at the end of the epoch and the new cluster centers:

K1 = K2 = c1_new = c2_new =

Solution:

epoch1 – start:

3: seed of cluster $K1 \Rightarrow 3 \in K1$

5: d(c1=3, 5)=2, $d(c2=10, 5)=5 => 5 \in K1$

7: d(c1=3, 7)=4, $d(c2=10, 7)=3 => 7 \in K2$

10: seed of cluster $K2 \Rightarrow 10 \in K2$

13: d(c1=3, 13)=10, $d(c2=10, 13)=3 => 13 \in K2$

The smaller distance for each example is in bold.

=> At the end of the first epoch:

clusters: $K1=\{3, 5\}$ and $K2=\{7, 10, 13\}$

new centroids: c1_new=(3+5)/2=4 and c2_new=(7+10+13)/3=10

Exercise 2. Nearest neighbor clustering

Use the Nearest Neighbor clustering algorithm to cluster examples A, B, C and D described by the following distance matrix. Suppose that the threshold *t* is 3.

	A	В	C	D
Α	0	1	4	5
В		0	2	6
С			0	4
D				0

Solution:

- A is placed in a cluster by itself, so we have $K1=\{A\}$.
- We then look at B if it should be added to K1 or be placed in a new cluster. As d(A,B)=1 < t, B is added to K1, i.e. $K1=\{A,B\}$

- C: added to K1 or placed in a new cluster? d(C,A)=4, d(C,B)=2; the smallest one is d(C,B) and it is < t > C is added to K1, i.e. $K1=\{A,B,C\}$
- D: added to K1 or placed in a new cluster? d(D,A)=5, d(D,B)=6, d(D,C)=4; the smallest is d(D,C) and it is >t => D is placed in a new cluster $K2=\{D\}$
- Final clusters: $K1=\{A,B,C\}$, $K2=\{D\}$

Exercise 3. Hierarchical clustering – single link agglomerative algorithm

Use the **single link** agglomerative clustering to group the data described by the following distance matrix. Draw the dendrogram.

	A	В	C	D
A	0	1	4	5
В		0	2	6
C			0	3
D				0

Solution:

Level 0:

$$(0, 4, \{A\}, \{B\}, \{C\}, \{D\})$$

Level 1: we can merge A and B as $d(A,B) \le 1$ (1, 3, {A,B}, {C}, {D})

The updated matrix is:

	AB	С	D
AB	0	2	5
C		0	3
D			0

Note: the distance between $\{A,B\}$ and C using the single link is $\min(d(A,C), d(B,C))=\min(4,2)=2$. Similarly, the distance between $\{A,B\}$ and D is 5.

Level 2: we can merge $\{A,B\}$ and C as the distance between them<=2 $(2, 2, \{A,B,C\}, \{D\})$

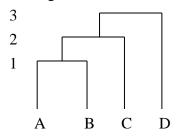
The updated matrix is:

	ABC	D
ABC	0	3
D		0

Level 3: we can merge $\{A,B,C\}$ with D as the distance between them is <=3 $(3,1,\{A,B,C,D\})$

Stop: all items are in 1 cluster.

Dendrogram:



Exercise 4. Hierarchical clustering – complete link agglomerative algorithm

The same task as in the previous exercise but using the **complete link** distance measure.

Solution:

Level 0:

$$(0, 4, \{A\}, \{B\}, \{C\}, \{D\})$$

Level 1: we can merge A and B as the distance between them is ≤ 1 $(1, 3, \{A,B\}, \{C\}, \{D\})$ as $d(A,B) \leq 1$

The updated matrix is:

	AB	C	D
AB	0	4	6
С		0	3
D			0

Note: the distance between $\{A,B\}$ and C using the complete link is $\max(d(A,C), d(B,C))=\max(4,2)=4$. Similarly, the distance between $\{A,B\}$ and D is 6.

Level 2: we can't merge any clusters as all distances are >3 (2, 3, {A,B}, {C}, {D})

Level 3: we can merge C and D as the distance between them is \leq 3 (3, 2, {A,B}, {C,D}

The updated matrix is:

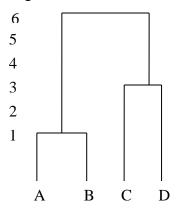
	AB	CD
AB	0	6
CD		0

Level 4: no merging Level 5: no merging

Level 6: we can merge the 2 clusters

Stop: all items are in 1 cluster

Dendrogram:



Exercise 5. Clustering using Weka

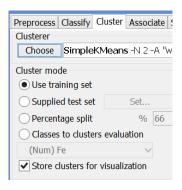
Load the glass.arff data. It describes different types of glass based on their chemical components. The identification of different types of glass is important for criminological investigations – it can be used as evidence.

There are 9 attributes: 1) RI: refractive index and 2)-9) measurement of the following chemical elements: Na (Sodium), Mg (Magnesium), Al (Aluminum), Si (Silicon), K (Potassium), Ca (Calcium), Ba (Barium) and Fe (Iron).

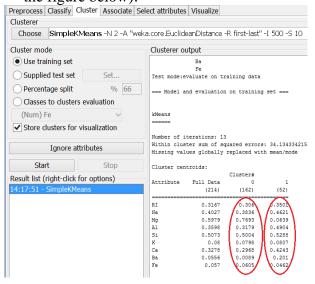
There are 7 classes (types of glass) but for one of them there are no examples in the dataset, so there are 6 classes:

Type of glass: (class attribute)

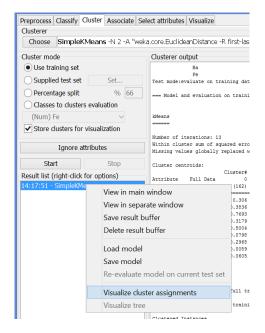
- 1) building_windows_float_processed
- 2) building_windows_non_float_processed
- 3) vehicle_windows_float_processed
- 4) vehicle_windows_non_float_processed (none in this database)
- 5) containers
- 6) tableware
- 7) headlamps
- 1. From the **Preprocess** tab:
 - a) Select the class attribute "type" and remove it. Clustering is an unsupervised method and doesn't use the class attribute.
 - b) Normalise the data using **unsupervised->attribute->Normalize** filter. This is important as the clustering algorithms we will be applying (k-means and hierarchical) are distance-based.
- 2. To perform clustering using the k-means algorithm, click the **Cluster** tab and select the **SimpleKMean**s algorithm. By default it uses k=2 clusters and Euclidean distance. You can see and change the parameters of the algorithm by right clicking on the name, then **Show Properties**.
- 3. Evaluation method: check that **Use training set** is selected and that **Store clusters for visualization** is also selected (both should be the default options).



4. Run the k-means algorithm and analyse the output. It shows the within cluster sum of squared errors (the smaller the better, i.e. high cohesion) and also the centroids for each of the two clusters (circled in the figure below):



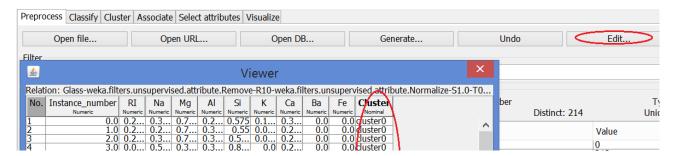
You can visualize the data by using Visualize cluster assignments:



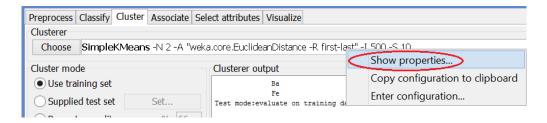
Try different attributes for Y to see which attributes are more discriminative for the 2 clusters. E.g. Mg separates the 2 clusters relatively well:



You can also save the clustering results by clicking **Save** on the Visualization panel. The results will be saved in a .arff file. For each example, at the end of the line, Weka will add the cluster of the example (cluster 0 or cluster 1). You can open and view the saved file with Weka:



Experiment with different number of clusters: right click on **SimpleKMeans** -> **Show properties**, then change k:

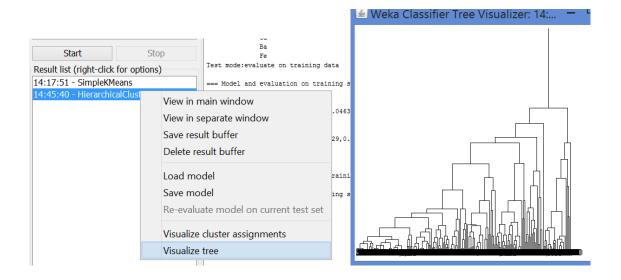


5. Weka also includes an implementation of the hierarchical agglomerative algorithm; it is called **HierarchicalClusterer**:



It includes different ways to measure the distance between the cluster – single link, complete link, etc. Explore them from **Show properties** -> **More.** Select one of them, e.g. the complete link and run the algorithm.

You can visualise the results pairwise as in the k-means algorithm and can also plot the hierarchical tree:



Additional exercises to be done at your own time

Exercise 6. K-means clustering

Use the k-means algorithm and Euclidean distance to group the following 8 examples into 3 clusters: A1=(2,10), A2=(2,5), A3=(8,4), A4=(5,8), A5=(7,5), A6=(6,4), A7=(1,2), A8=(4,9). The distance matrix based on the Euclidean distance is given below:

	A1	A2	A3	A4	A5	A6	A7	A8
A1	0	$\sqrt{25}$	$\sqrt{72}$	$\sqrt{13}$	$\sqrt{50}$	$\sqrt{52}$	$\sqrt{65}$	$\sqrt{5}$
A2		0	$\sqrt{37}$	$\sqrt{18}$	$\sqrt{25}$	$\sqrt{17}$	$\sqrt{10}$	$\sqrt{20}$
A3			0	$\sqrt{25}$	$\sqrt{2}$	$\sqrt{2}$	$\sqrt{53}$	$\sqrt{41}$
A4				0	$\sqrt{13}$	$\sqrt{17}$	$\sqrt{52}$	$\sqrt{2}$
A5					0	$\sqrt{2}$	$\sqrt{45}$	$\sqrt{25}$
A6						0	$\sqrt{29}$	$\sqrt{29}$
A7							0	$\sqrt{58}$
A8								0

Suppose that the initial seeds (centroids) are A1, A4 and A7. Run the k-means algorithm for 1 epoch only. At the end of this epoch show:

- a) the new clusters (i.e. the examples belonging to each cluster)
- b) the centroids of the new clusters

Solution:

a) d(x,y) denotes the Euclidean distance between x and y K1, K2 and k3 are the three clusters seed1=A1=(2,10), seed2=A4=(5,8), seed3=A7=(1,2) epoch1 – start:

A1:

d(A1, seed1)=0 as A1 is seed1 d(A1, seed2)>0

d(A1, seed3) > 0

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=> A1 \in K1
A2:
d(A2, seed1)=d((2,5), (2,10))=sqrt((2-2)^2+(5-10)^2)=sqrt(25)
d(A2, seed2)=d((2,5), (5,8))=sqrt((1-5)^2+(5-8)^2)=sqrt(18)
d(A2, seed3)=d((2,5),(1,2))=sqrt((2-5)^2+(5-2)^2)=sqrt(10)
=> A2 \in K3
A3:
d(A3, seed1)=d((8,4), (2,10))=sqrt(72)
d(A3, seed2)=d((8,4), (5,8))=sqrt(25)
d(A3, seed3)=d((8,4), (1,2))=sqrt(53)
=> A2 \in cluster2
A4:
d(A4, seed1)>0
d(A4, seed2)=0 as A4 is seed2
d(A4, seed3)>0
=> A4 \in K2
A5:
d(A5, seed1)=d((7,5), (2,10))=sqrt(50)
d(A5, seed2)=d((7,5), (5,8))=sqrt(13)
d(A5, seed3)=d((7,5), (1,2))=sqrt(45)
=> A5 \in K2
A6:
d(A6, seed1)=d((6,4), (2,10))=sqrt(52)
d(A6, seed2)=d((6,4), (5,8))=sqrt(17)
d(A6, seed3)=d((6,4), (1,2))=sqrt(29)
=> A6 \in K2
A7:
d(A7, seed1)>0
d(A7, seed2)>0
d(A7, seed3)=0 as A7 is seed3
=> A7 \in K3
A8:
d(A8, seed1)=d((4,9), (2,10))=sqrt(5)
d(A8, seed2)=d((4,9), (5,8))=sqrt(2)
d(A8, seed3)=d((4,9), (1,2))=sqrt(58)
=> A8 \in K2
end of epoch1
new clusters: K1 = \{A1\}, K2 = \{A3, A4, A5, A6, A8\}, K3 = \{A2, A7\}
b) centroids of the new clusters:
K1: (2,10)
K2: (8+5+7+6+4)/5, (4+8+5+4+9)/5=(6,6)
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K3: ((2+1)/2, (5+2)/2)=(1.5, 3.5)