

Intelligent Systems

Excercise 7 – Clustering

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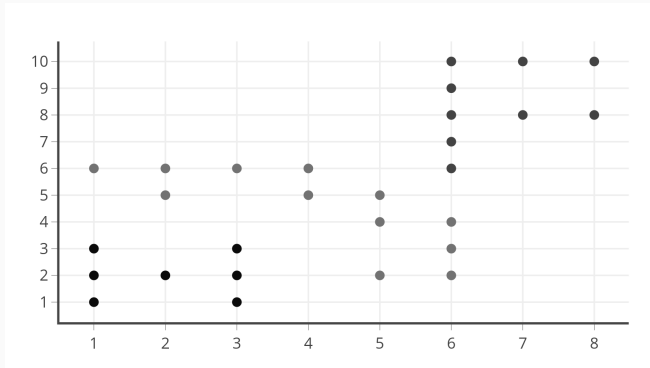
1. Single, Complete und Average Linkage
2. c-Means basics
3. Apply c-Means Clustering

Single, Complete und Average Linkage

- A. How does Single Linkage Clustering work? Visualise the procedure using the data set of the figure with $C = 3$ and with $C = 2$. As distance measure choose the Manhattan distance.**
- B. What are Pros and Cons of Single Linkage in comparison to complete Linkage regarding the treatment of outliers and the tendency of producing chains?
- C. What is the difference between Single Linkage, Complete Linkage, and Average Linkage? How would Complete or Average Linkage cluster the data points in the figure with $C = 2$ and the Manhattan distance?

1. A SINGLE LINKAGE CLUSTERING

How does Single Linkage Clustering work? Visualise the procedure using the data set of the figure with $C = 3$ and with $C = 2$. As distance measure choose the Manhattan distance.



Given a set of samples x_i mit $i = 1, \dots, N$ and additionally a number of cluster c that are required to be found. ($N \geq c$).

1. First, partition the samples such that every sample is assigned to a different cluster. Given N samples results in N clusters:
 - $C_i = \{x_i\}$
2. These N clusters are merged stepwise:
 - 2.1 Determine two clusters with minimum distance according to the criterion nearest neighbor and merge them.
 - 2.2 If the number of clusters is still above c , repeat step 2.1.

A) SINGLE LINKAGE EXAMPLE

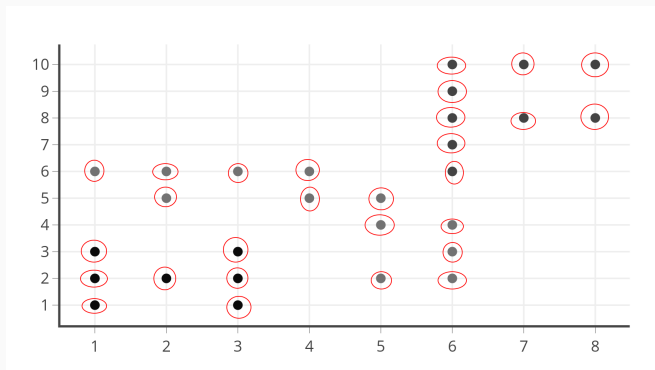


Figure 1: *

Every sample is assigned to a different cluster.

A) SINGLE LINKAGE EXAMPLE $C=3$

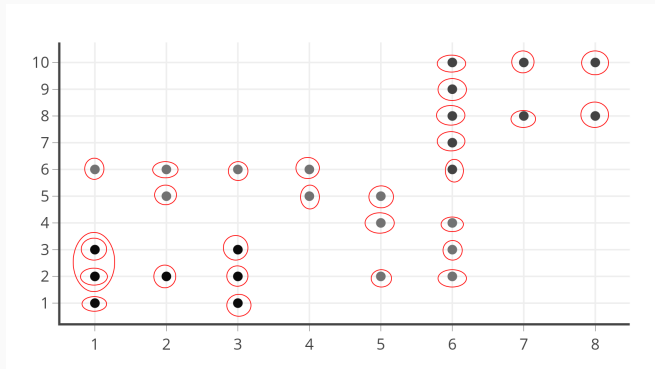


Figure 2: *

Merge two clusters, until the number of required clusters $c = 3$ is reached.

A) SINGLE LINKAGE EXAMPLE $C=3$

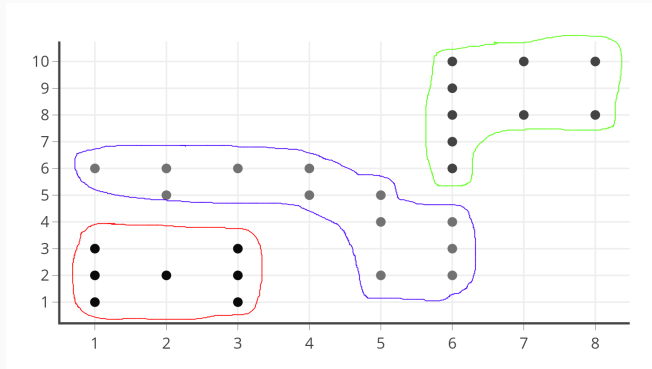


Figure 3: *

Resulting 3 clusters.

A) SINGLE LINKAGE EXAMPLE $C=2$

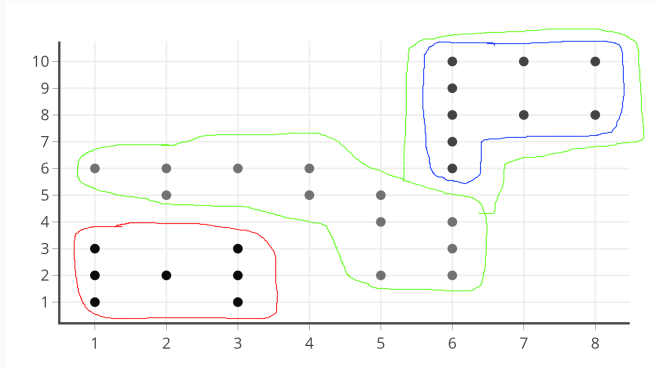


Figure 4: *

Merge two clusters, until the number of required clusters $c = 2$ is reached.

A) SINGLE LINKAGE BEISPIEL C=2

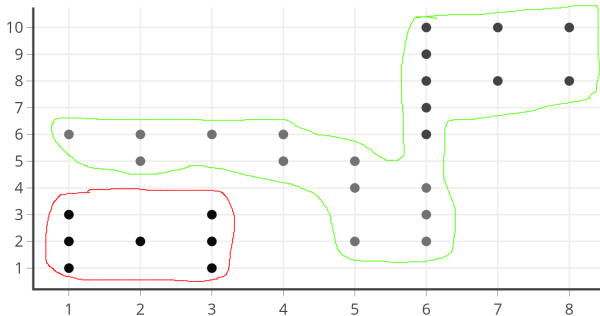


Figure 5: *

The possible result shows a poor clustering. The middle cluster should be merged to the bottom cluster → chaining problem

- A. How does Single Linkage Clustering work? Visualise the procedure using the data set of the figure with $C = 3$ and with $C = 2$. As distance measure choose the Manhattan distance.
- B. What are Pros and Cons of Single Linkage in comparison to complete Linkage regarding the treatment of outliers and the tendency of producing chains?**
- C. What is the difference between Single Linkage, Complete Linkage, and Average Linkage? How would Complete or Average Linkage cluster the data points in the figure with $C = 2$ and the Manhattan distance?

What are Pros and Cons of Single Linkage in comparison to complete Linkage regarding the treatment of outliers and the tendency of producing chains?

Pros:

- Outliers are detected, because they will be added lastly

Cons:

- Only two samples matters for merging two clusters → What if these are noise or outliers?
- Chaining problem: long chains are able to form clusters with a large maximum point distance (Remark: sometimes chaining is required for clustering geographical data, i.e. rivers)

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- C. What is the difference between Single Linkage, Complete Linkage, and Average Linkage? How would Complete or Average Linkage cluster the data points in the figure with $C = 2$ and the Manhattan distance?**

- Similarly as Single Linkage but maximum distant neighbor as distance measure

Pros:

- No chaining as in Single Linkage
For example: In Exercise 1A with $c = 2$ the bottom and middle cluster will be merged definitely.

Cons:

- Outliers will be merged to clusters most likely and won't be detected as outliers

- Similarly as Single Linkage but average distant neighbor as distance measure
- Average Linkage is a trade-off between Single Linkage and Complete Linkage
- In many cases this approach succeeds.
- For example:
The middle cluster in exercise 1A with $c = 2$ will be merged correctly to the bottom cluster.

c-Means basics

- A. Visualise and explain with the help of the figure how the **c-Means algorithm works.**
- B. What are Pros and Cons of the c-Means algorithm?
- C. Which steps can be applied to optimise the clustering results?

Given N d -dimensional samples $X = \{x_i\}_{i=1}^N \subset \mathbb{R}^d$ and the number of required clusters $c \in \mathbb{N}$

1. Distribute c cluster centres $c_j, j = 1, \dots, c$ (arbitrary) in the space \mathbb{R}^d

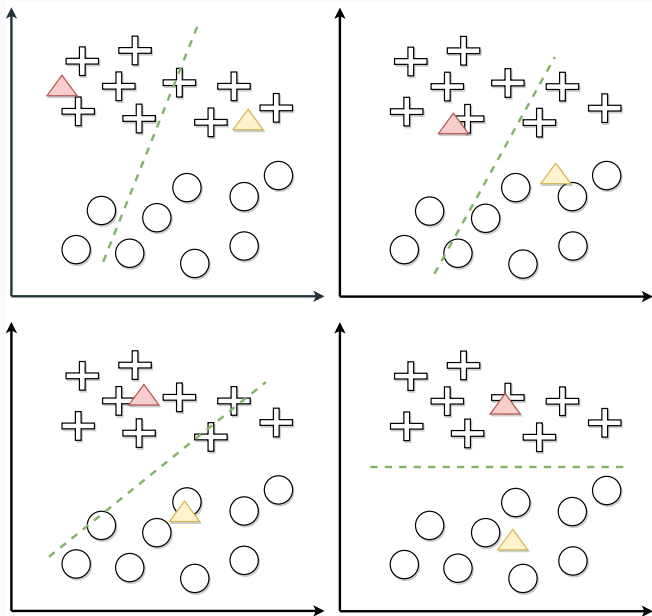
2. Assign every sample to the nearest cluster centre
 $class(x_i) = \underset{j=1, \dots, k}{\operatorname{argmin}} \|x_i - c_j\|_2$

3. Distribute new cluster centres

$$c_j = \frac{1}{|\{x \in X | class(x) = j\}|} \sum_{x \in \{x' \in X | class(x') = j\}} x$$

4. Repeat steps 2)-3) and terminate after k iterations or if no new cluster assignment took place

3. A C-MEANS PROCEDURE II



- A. Visualise and explain with the help of the figure how the *c*-Means algorithm works.
- B. What are Pros and Cons of the *c*-Means algorithm?**
- C. Which steps can be applied to optimise the clustering results?

What are Pros and Cons of the c -Means algorithm?

Pros:

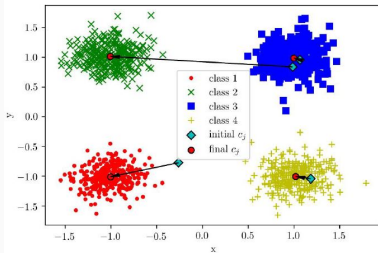
- Easy, understandable
- Every sample is assigned to a cluster
- Algorithm terminates

Cons:

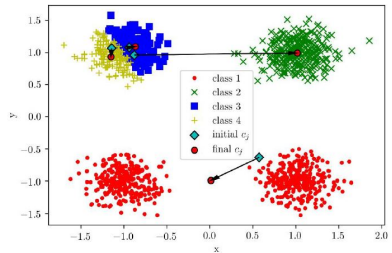
- Greedy procedure \rightarrow convergence at local minimum possible
- Number of clusters has to be known or estimated
- Also outliers will be assigned to clusters
- Sensible according to poor initial cluster centres placement

- A. Visualise and explain with the help of the figure how the *c*-Means algorithm works.
- B. What are Pros and Cons of the *c*-Means algorithm?
- C. Which steps can be applied to optimise the clustering results?**

Which steps can be applied to optimise the clustering results?
In the figure: How can we prevent the clustering in the right image?



a) Successful clustering



b) Failed clustering

Unlucky placement of cluster centres:

- Repeat procedure with different randomised initial cluster centres placements
- Intelligent cluster centres placement

Not suitable number of clusters:

- Begin with small c and evaluate the minimum cluster distance with ascending c
- Decide according to elbow method

3. C ELBOW-CRITERION

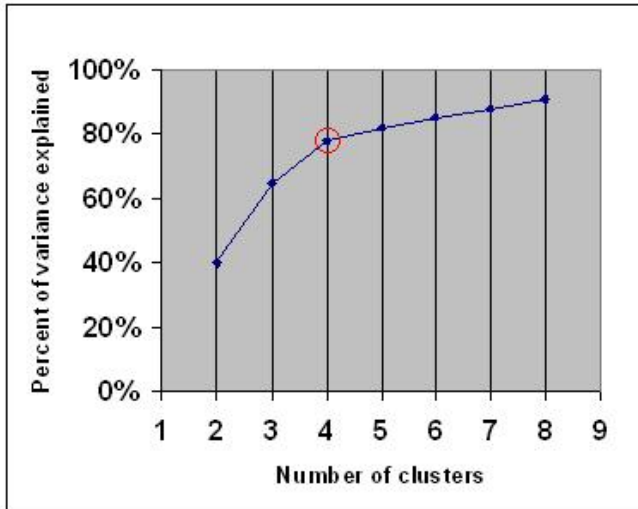


Figure 6: *

- **Solution 1:**

Increase c and merge similar clusters

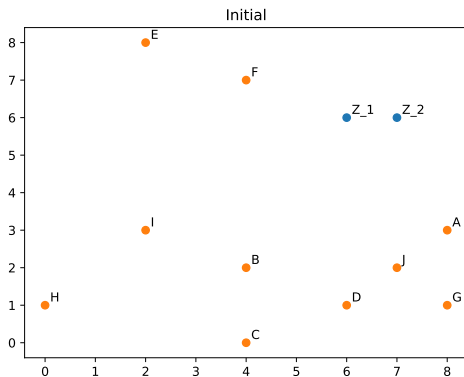
- **Solution2:**

Apply clever cluster centre distribution:

- Approach 1: Random distribution
 - a) Select c samples randomly as cluster centre
 - b) Choose random points within minimum sphere containing all samples.
- Approach 2: Sort samples x_1, \dots, x_N according distance to global centre $c_{global} = \frac{1}{N} \sum_{i=1}^N x_i$ and select nearest $(1 + (j - 1) \cdot \lfloor \frac{N}{c} \rfloor)$ -th sample as cluster centre.

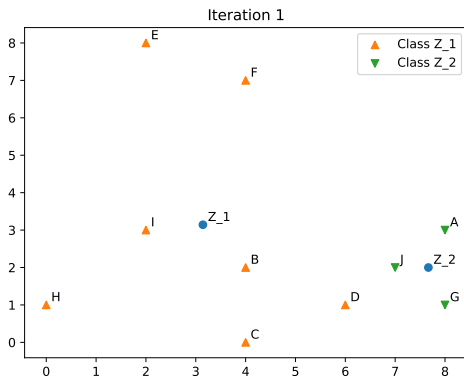
Apply c-Means Clustering

A. Proceed 4 iterations of the c-Means clustering on the points given in the figure. Note the Euclidean distances into the table. Remark: You can use a ruler to measure the Euclidean distances.



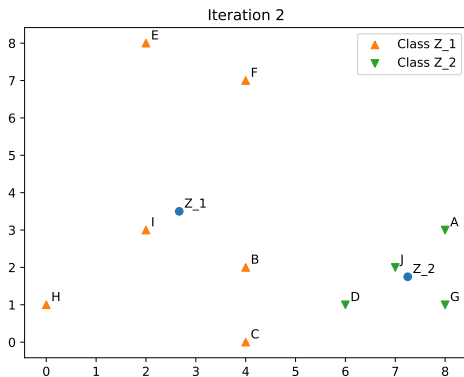
Iteration		<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	<i>F</i>	<i>G</i>	<i>H</i>	<i>I</i>	<i>J</i>
0	$Z_1 = (6.0, 6.0)$	3.6	4.5	6.3	5.0	4.5	2.2	5.4	7.8	5.0	4.1
0	$Z_1 = (7.0, 6.0)$	3.2	5.0	6.7	5.1	5.4	3.2	5.1	8.6	5.8	4.0

2. A ITERATION 1



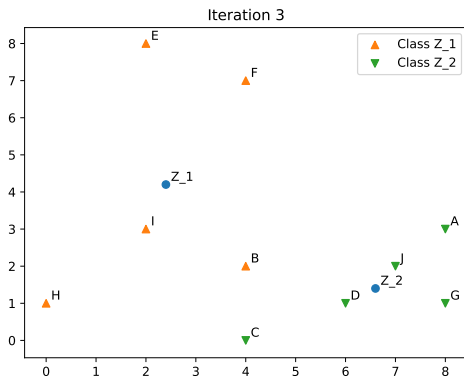
Iteration		<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	<i>F</i>	<i>G</i>	<i>H</i>	<i>I</i>	<i>J</i>
1	$Z_1 = (3.1, 3.1)$	4.9	1.4	3.3	3.6	5.0	4.0	5.3	3.8	1.2	4.0
1	$Z_1 = (7.7, 2.0)$	1.1	3.7	4.2	1.9	8.3	6.2	1.1	7.7	5.8	0.7

2. A ITERATION 2



Iteration		<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	<i>F</i>	<i>G</i>	<i>H</i>	<i>I</i>	<i>J</i>
2	$Z_1 = (2.7, 3.5)$	5.4	2.0	3.7	4.2	4.5	3.7	5.9	3.7	0.8	4.6
2	$Z_1 = (7.2, 1.8)$	1.5	3.3	3.7	1.5	8.2	6.2	1.1	7.3	5.4	0.4

2. A ITERATION 3



Iteration		<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	<i>F</i>	<i>G</i>	<i>H</i>	<i>I</i>	<i>J</i>
3	$Z_1 = (2.4, 4.2)$	5.7	2.7	4.5	4.8	3.8	3.2	6.4	4.0	1.3	5.1
3	$Z_2 = (6.6, 1.4)$	2.1	2.7	3.0	0.7	8.0	6.2	1.5	6.6	4.9	0.7