AI & Robotics



Goals (1/2)



The junior-colleague

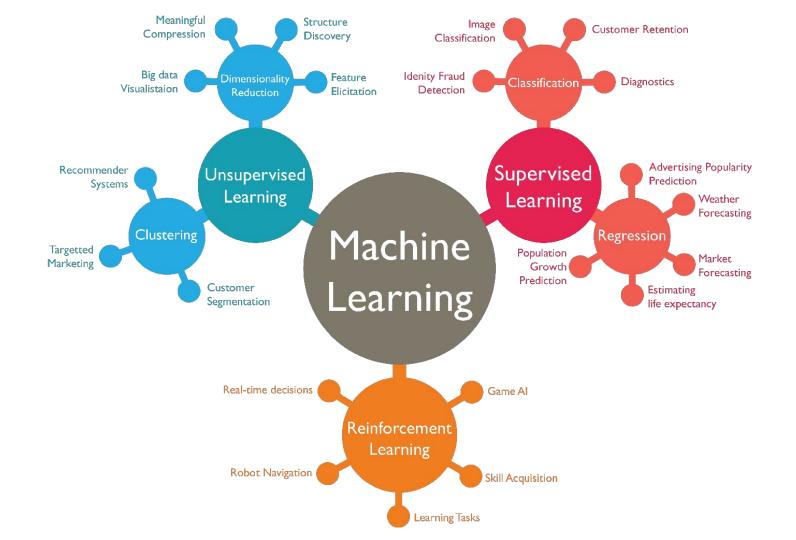
- can explain Unsupervised Learning in their own words
- can describe the general flow of an unsupervised learning pipeline
- can list at least 3 examples of unsupervised learning
- can explain the purpose of clustering
- can explain the difference between clustering and classification
- can explain how to prepare data for clustering
- can explain how the k-means algorithm works in their own words
- can describe the advantages and disadvantages of k-means
- can explain how k-means++ optimizes the centroid choice problem of k-means
- can explain how the Elbow method in the context of k-means works
- can implement k-means and DBSCAN for a given clustering problem

Goals (2/2)



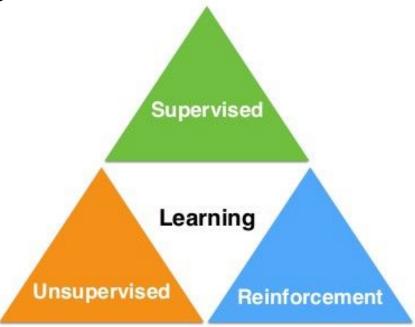
The junior-colleague

- can explain how the DBSCAN clustering algorithm works in their own words
- can describe the advantages and disadvantages of DBSCAN
- can explain how hierarchical clustering works on the basis of a visual representation of a dataset
- can describe the different linkage forms for hierarchical clustering
- can describe the advantages and disadvantages of hierarchical clustering



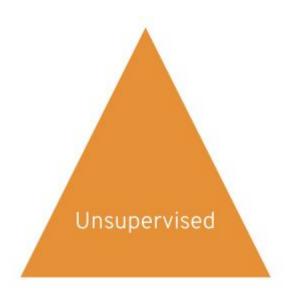
Machine Learning

- Labeled data
- · Direct feedback
- Predict outcome/future

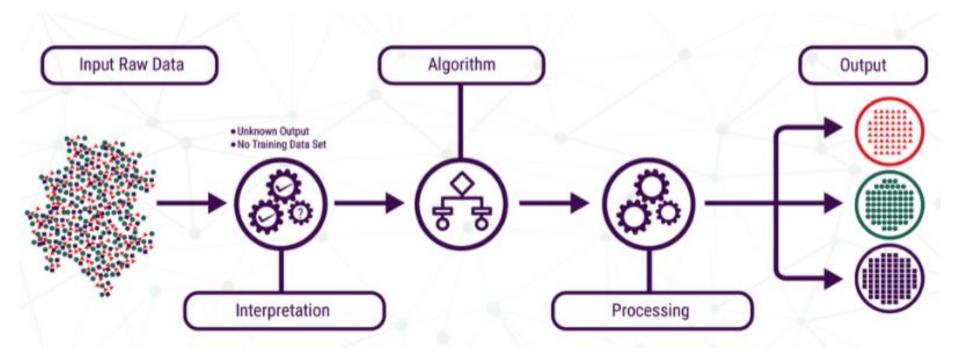


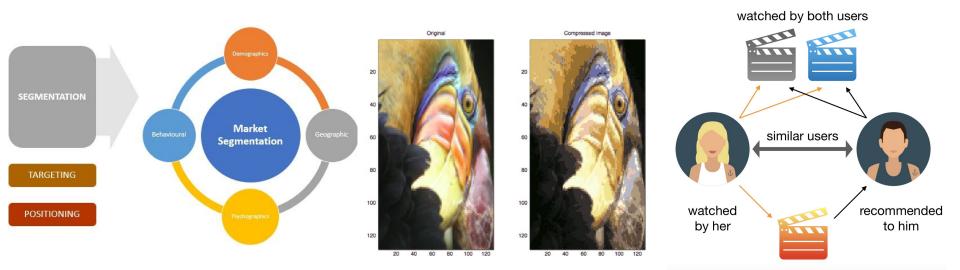
- · No labels
- No feedback
- · "Find hidden structure"

- · Decision process
- · Reward system
- Learn series of actions

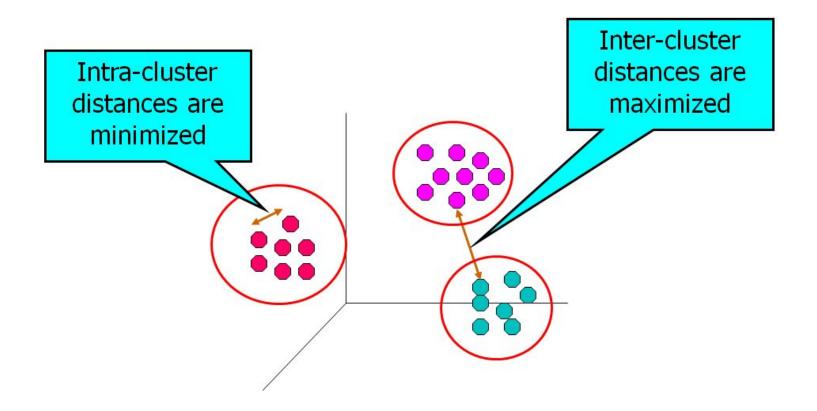


- No labels
- No feedback
- "Find hidden structure"
- Meaningful patterns in unlabeled data
- Usage is on the rise





Clustering



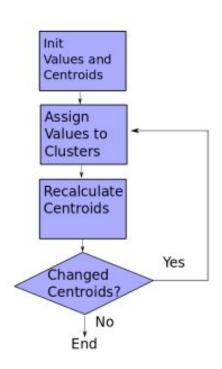
Clustering

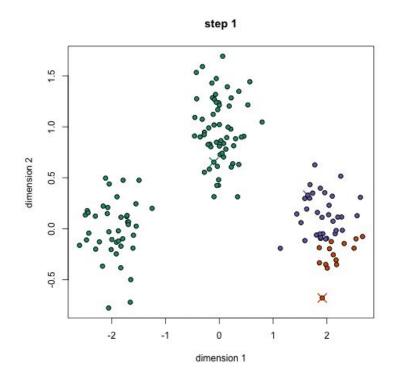
- Centroid-based clustering
 - K-Means
 - Mean-shift
- Density-based Clustering
 - DBSCAN
 - Mean-shift
- Connectivity-based clustering
 - Hierarchical (agglomerative) Clustering

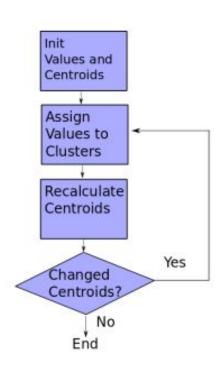
Preparing data

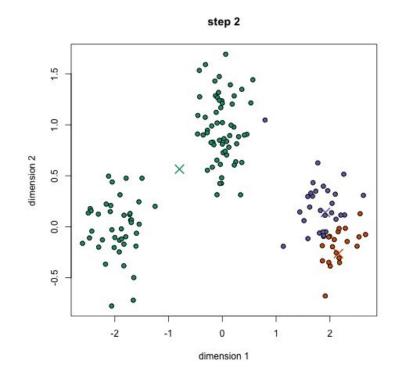
- Categorical variables to numerical data
- Handle missing values
- Clustering uses distance metrics
 - => Scale the data
 - Not necessary for some types of well defined data
 => i.e. clustering geolocation data (longitudes and latitudes)
 - Necessary for data that is of different physical measurements or units

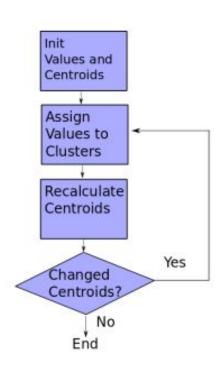
- Very popular clustering algorithm
- Easy to implement
- Naive method
 - => Doesn't know what the number of clusters is
 - => K (# of clusters) is supplied as a parameter
- Finds a local optimum

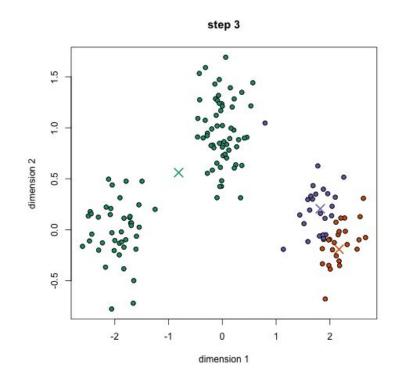


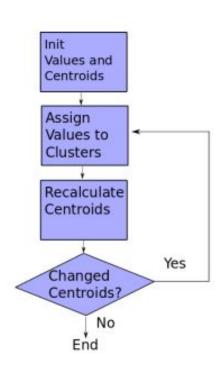


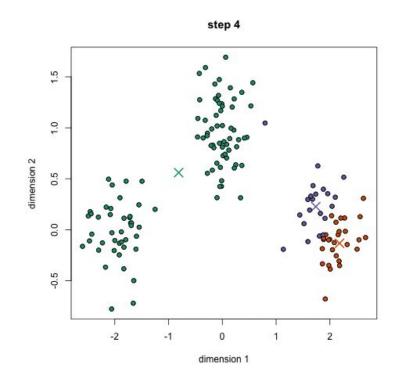


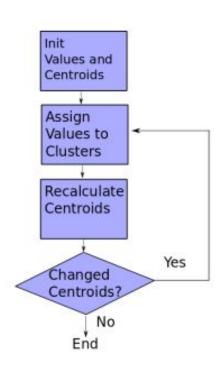


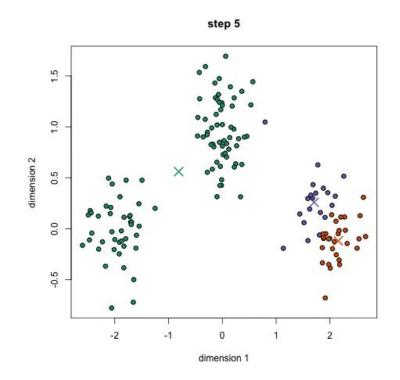


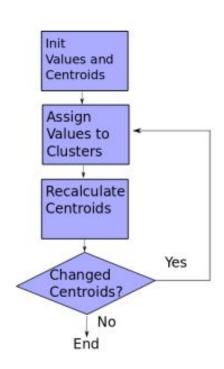


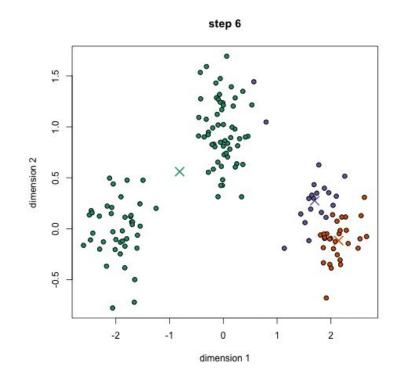


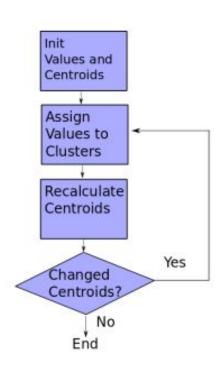


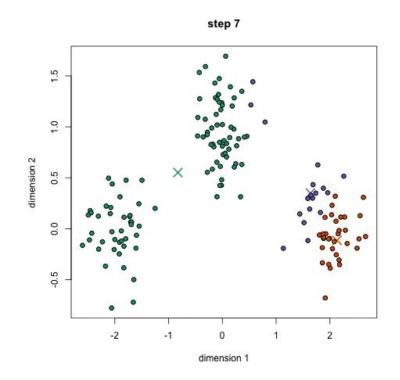


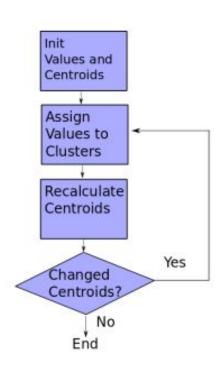


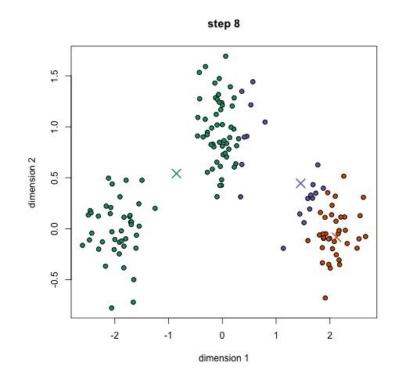


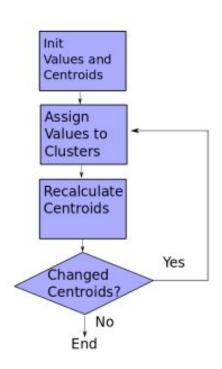


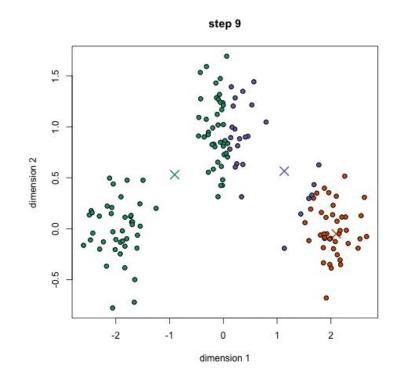


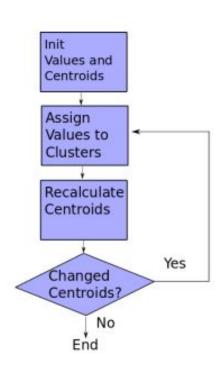


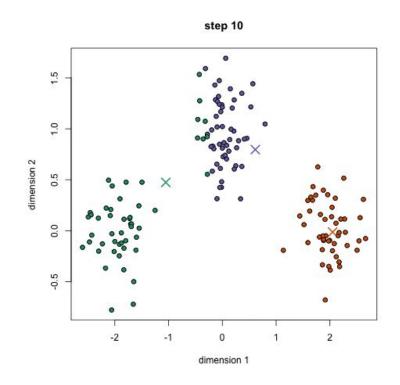


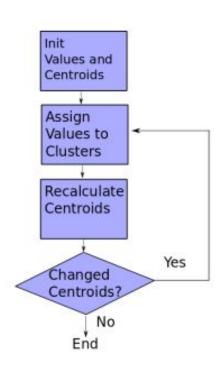


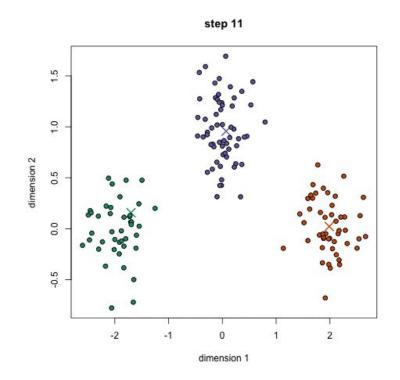


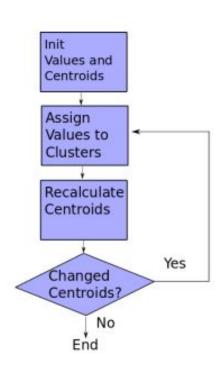


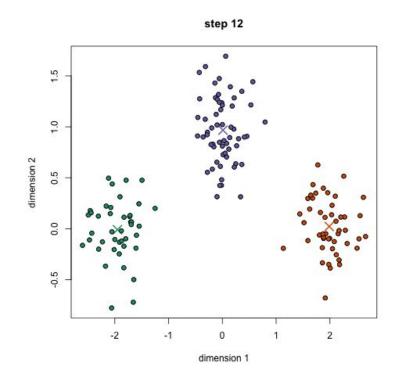








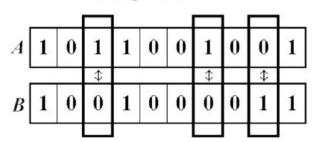


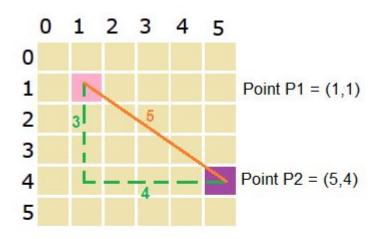


K-Means Clustering: Distance metrics

- Euclidean distance (most used)
- Manhattan distance
- Hamming distance

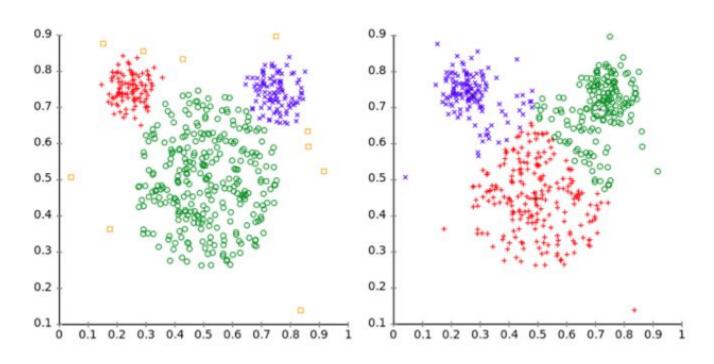




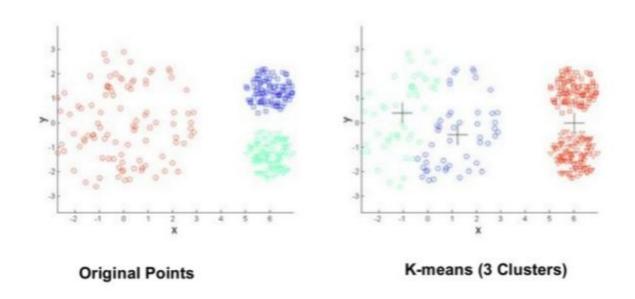


Euclidean distance =
$$\sqrt{(5-1)^2 + (4-1)^2} = 5$$

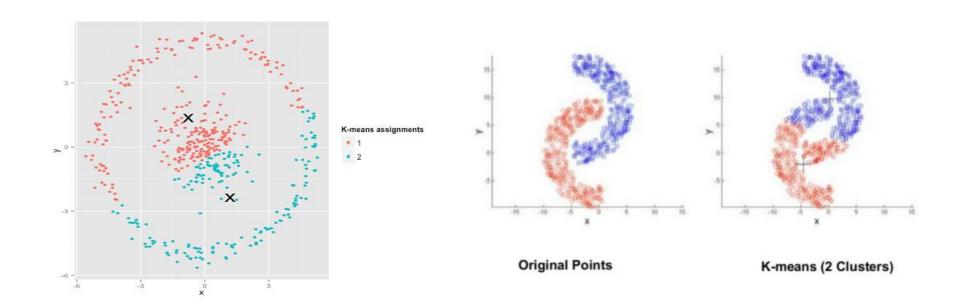
Clusters of different size



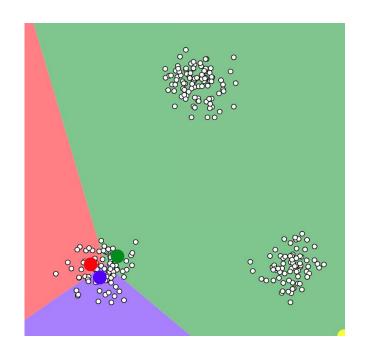
Clusters of different density

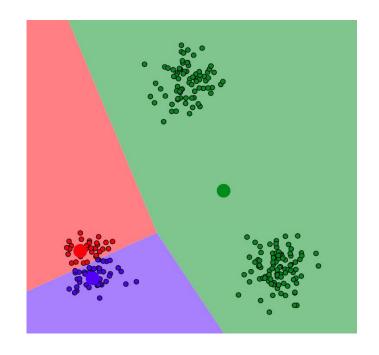


Clusters of non-spherical shape



Choosing the initial centroids





| Advantages | Disadvantages |
|-------------------------------|---|
| Fast and efficient | How to choose k? |
| Simple and easy to understand | How to choose the initial centroids? |
| | Sensitive to outliers |
| | Bad at handling clusters of different size |
| | Bad at handling clusters of different density |
| | Bad at handling clusters of non-spherical shape |

- Choosing centroids
 - K-means++
 - Don't choose randomly
 - Choose the next centroid with a probability proportional to a distance function
 - Default in scikit-learn
- Estimating k
 - Elbow method
 - Silhouette score
 - Calinski-Harabasz index
 - Cluster instability

K-Means Clustering: Elbow Method

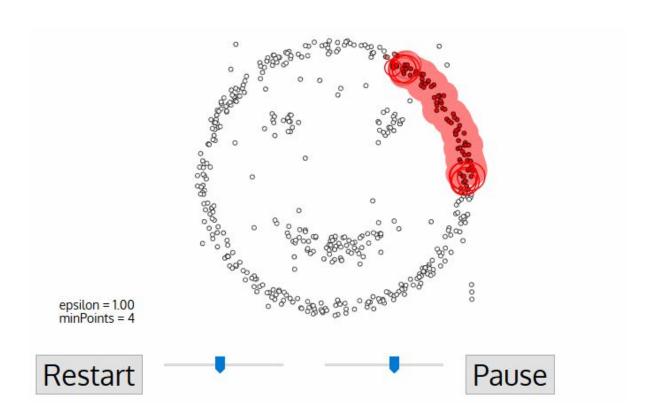


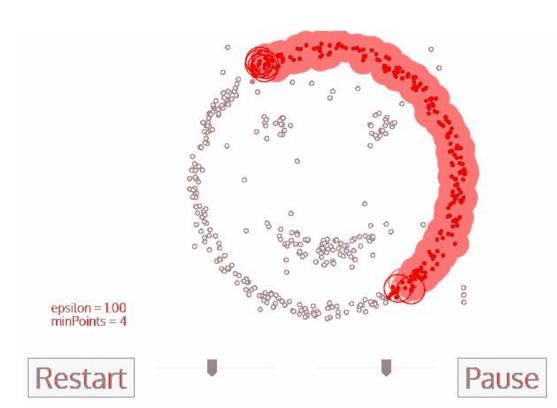
DBSCAN

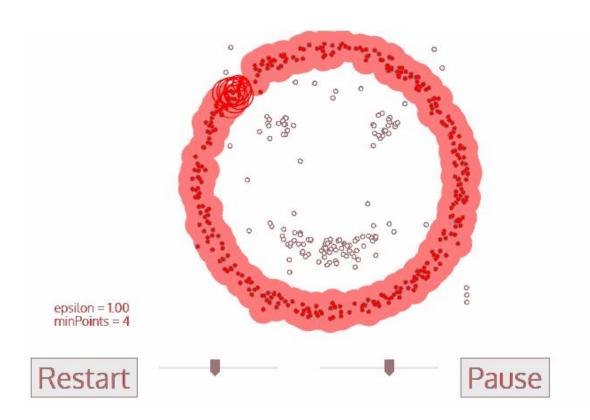
DBSCAN

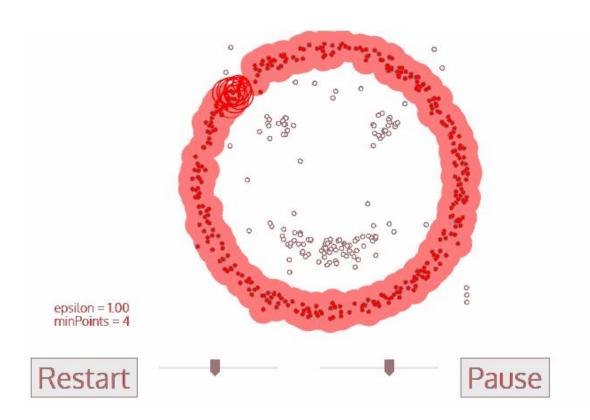
- Density Based Clustering
- A cluster is a high-density area (there are no restrictions on its shape) surrounded by a low-density one
- Better at dealing with non-convex problems
- Doesn't need an initial estimate about the number of expected clusters
- More robust against noise

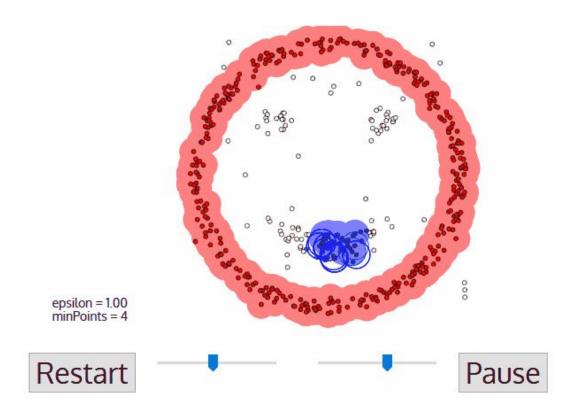
DBSCAN

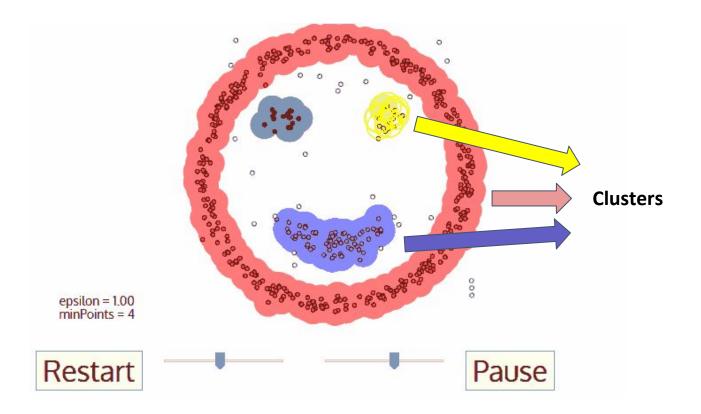


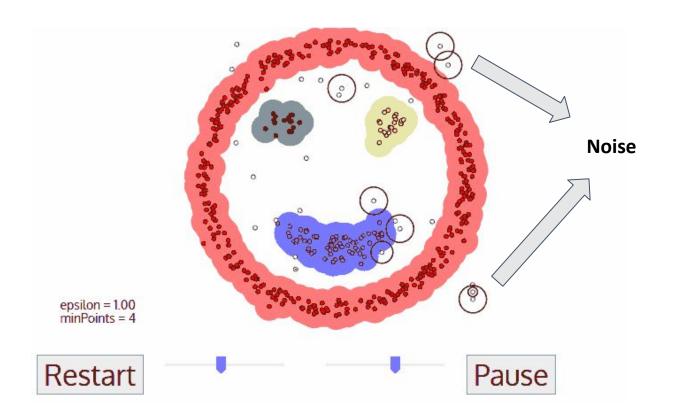












- 1. Start with an unvisited point
- 2. Check neighbourhood (based on parameter epsilon ε)
 - If surrounded by a minimum number of other samples (parameter min_samples)
 - => Part of cluster
 - b. Else:
 - => Noise
 - => In both cases this point is marked as "visited"
- 3. Check neighbours
 - => If they also have a high density, they are merged with the first area
- 4. If point is not in a cluster, continue with a new unvisited data point

Parameters:

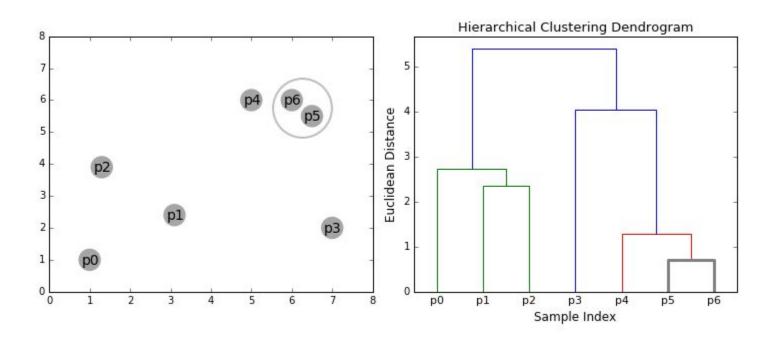
- eps: maximum distance between two neighbors.
 - => Higher values will aggregate more points
 - => Smaller values will create more clusters.
- min_samples : how many surrounding points are necessary to define an area

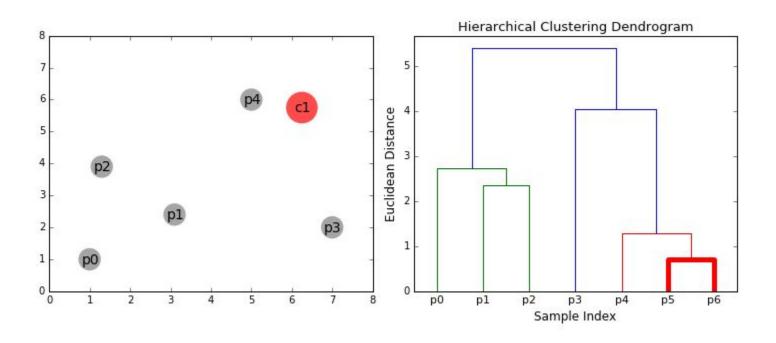
| Advantages | Disadvantages |
|--|---|
| No pre-set number of clusters required | Choosing ε |
| Robust against noise | Choosing minPoints |
| | Bad at handling clusters of different density |

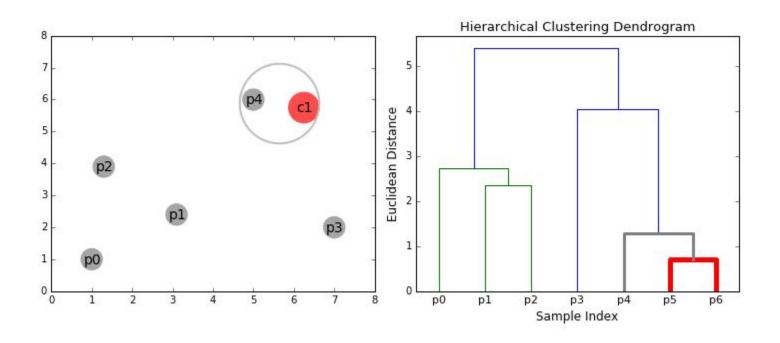
DBSCAN: OPTICS

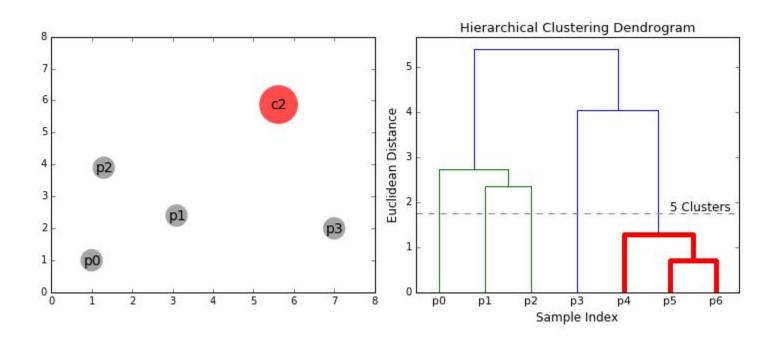
Ordering Points to Identify Cluster Structure

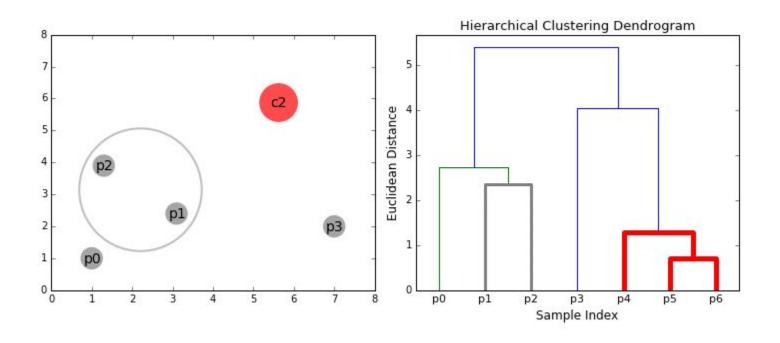
- Generalizes DBSCAN to clusters of different densities
- Epsilon becomes an optional maximum
- => Exact implementation falls outside the extent of this course
- => Important to know there is an optimization possible (check extra material on github)

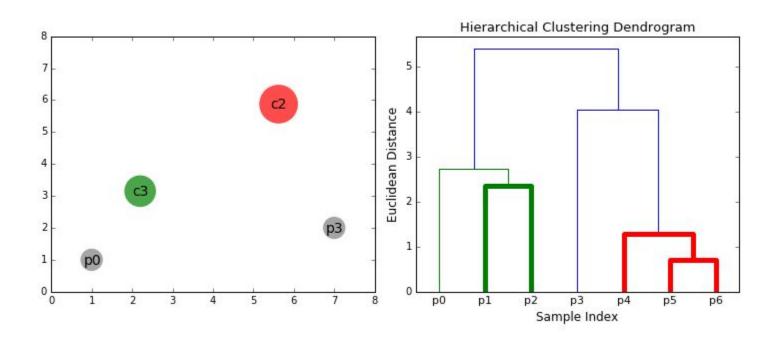


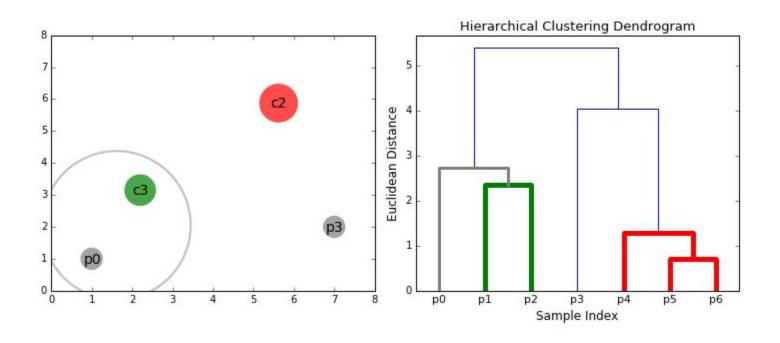


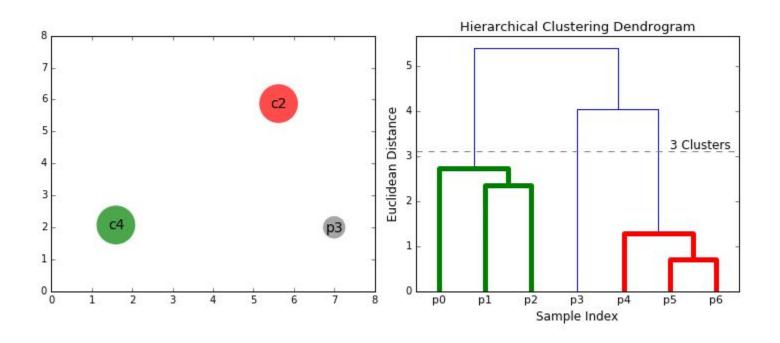










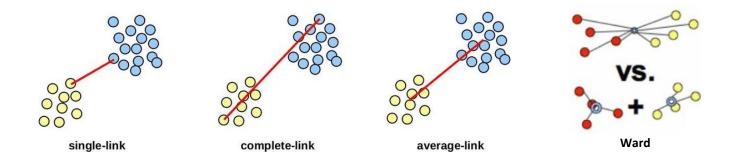


- Start with each data point as a single cluster
 X data points => X clusters
- Each iteration combines two clusters into one=> Choose clusters with smallest average linkage according to distance metric
- 3. Repeat until one cluster contains all data points

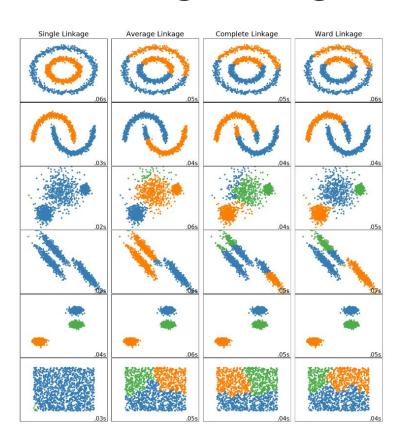
=> Select how many clusters we want in the end

Hierarchical clustering: Linkage

- Single linkage: minimizes the distance between the closest observations of pairs of clusters.
- Maximum or complete linkage: minimizes the maximum distance between observations of pairs of clusters.
- Average linkage: minimizes the average of the distances between all observations of pairs of clusters.
- Ward: minimizes the sum of squared differences within all clusters. It is a variance-minimizing approach (similar to the k-means). If you join 2 clusters, minimize the total distance to the new centroid



Hierarchical clustering: Linkage



| Advantages | Disadvantages |
|---|--|
| Easy to visualize with the dendrogram | Difficult to identify the correct number of clusters from the dendrogram |
| Provides hierarchical relations between clusters | Can be sensitive to noise and outliers based on linkage |
| No a priori information about the number of clusters required | Data points may be incorrectly grouped at an early stage |
| Less influenced by cluster shapes and densities | Slow |

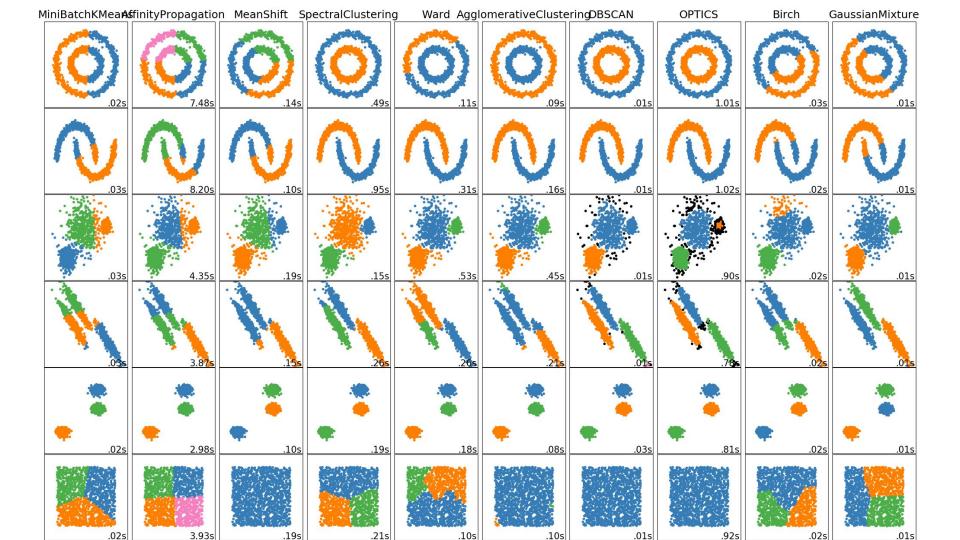
Clustering: Evaluation

- **Distortion/Inertia score:** Computes the sum of squared distances from each point to its assigned center
 - => Assumes convex clusters (only really useful for k-Means)

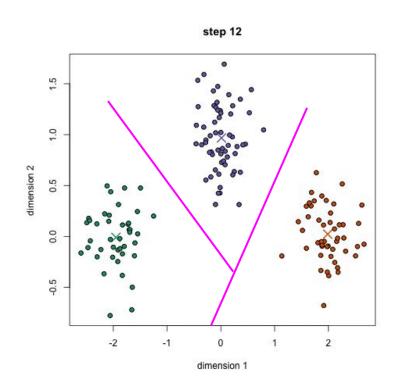
• Silhouette score:

- Measures the distance between each data point, the centroid of the cluster it was assigned to and the closest centroid belonging to another cluster
- A measure of how similar an object is to its own cluster (cohesion) compared to other clusters (separation)
- o [-1, 1]

| RANGE OF SC | INTERPRETATION |
|-------------|---|
| 0.71-1.0 | A strong structure has been found |
| 0.51-0.70 | A reasonable structure has been found |
| 0.26-0.50 | The structure is weak and could be artificial. Try additional methods of data analysis. |
| £ 0.25 | No substantial structure has been found |



From unsupervised to supervised



Once clusters are identified:

- Label the datapoints in the same cluster with the same label
- Apply supervised learning to new datapoints

