LECTURER: Nghia Duong-Trung

MACHINE LEARNING

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UNIT 2

CLUSTERING

STUDY GOALS

- Know the definitions and terms used for clustering
- Comprehend common applications of clustering analysis
- Understand different methods for clustering analysis
- Analise the advantages and limitations of the clustering methods
- Implement clustering methods in Python

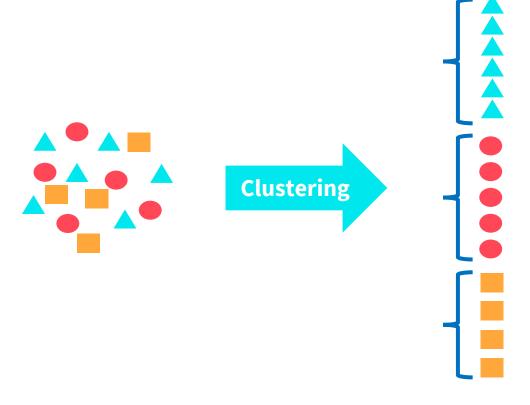




- 1. What is clustering and its application?
- 2. What are the popular methods of clustering, their advantages and limitations?
- 3. How to implement the clustering methods in python?

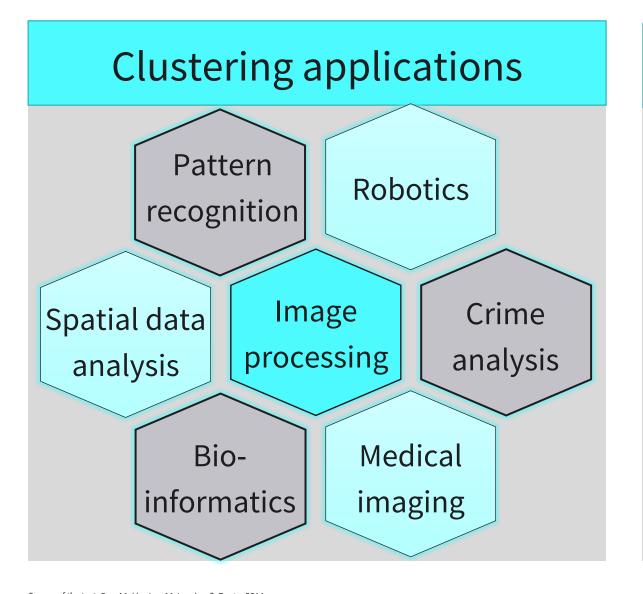
INTRODUCTION

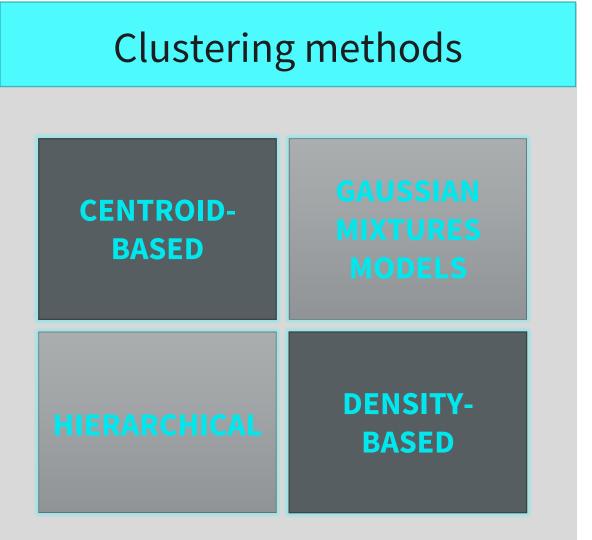
- Clustering is an unsupervised learning technique to:
 - reveal meaningful partitions,hierarchies
 - find association rules of data
 - gather data points into groups and extract useful information



An example of clustering

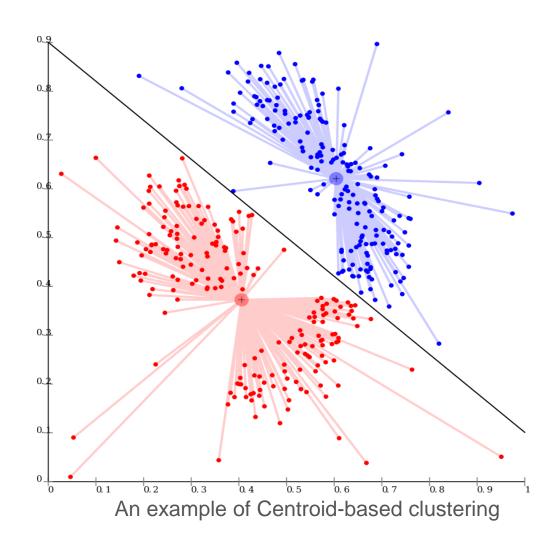
INTRODUCTION





2.1 CENTROID-BASED CLUSTERING

- Centroid: the arithmetic average position of all the points
- Centroid-based clustering
 searches for a pre-determined
 number of clusters
- Each data point is assigned to the cluster achieving the minimum distance from the centroid



2.1 CENTROID-BASED CLUSTERING

- K-Means Clustering partitions
 data points into K clusters
- Three main steps: initialization, assignment, update
- Advantages: simple to employ,
 efficient for large datasets
- Limitation: K must be determined,
 initialization-dependent, problems with
 high-dimensional data and outliers

Step 1 Step 2 Result Step 3

Algorithm of K-Means clustering

Source of the text: Data Science Team, 2020; Pragathi, Jayanthi, & Malathi, 2018; Zhu, 2022. Source of the image: File:K Means Example Step 1.svg (2020).

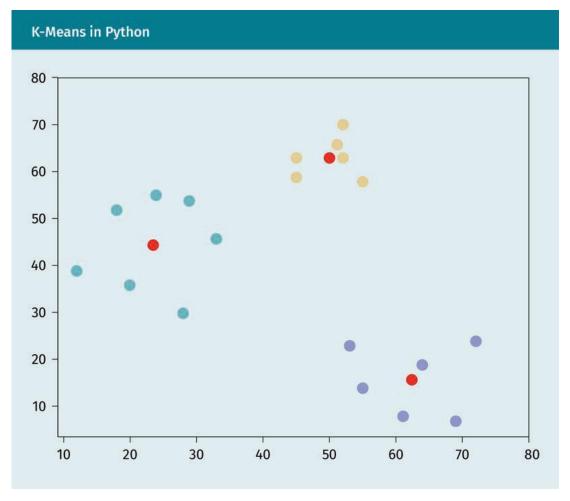
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File: K Means Example Step 3.svg (2020).

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2.1 CENTROID-BASED CLUSTERING

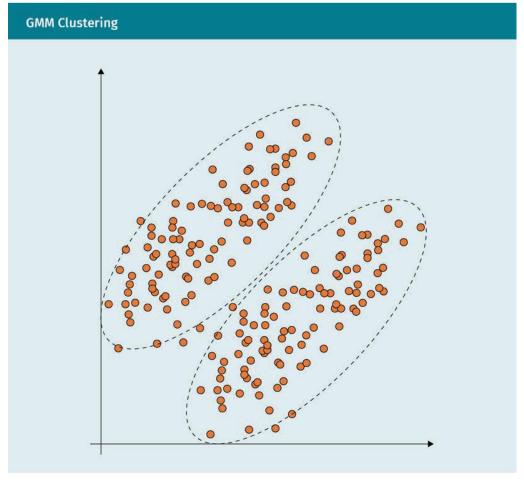
```
#K-Means clustering with Python
     >>> import pandas as pd # For reading datasets
     >>> import numpy as np # For computations
     >>> import matplotlib.pyplot as plt # For visualization
     >>> from pandas import DataFrame # For creating data frame
    >>> from sklearn.cluster import KMeans
     >>> Data={
    'x': [12, 20, 28, 18, 29, 33, 24, 45, 45, 52, 51, 52, 55, 53, 55, 61, 64, 69, 72],
     'y': [39, 36, 30, 52, 54, 46, 55, 59, 63, 70, 66, 63, 58, 23, 14, 8, 19, 7, 24]
10
     >>> df = DataFrame(Data,columns=['x','y'])
    # Create and fit the KMeans model
     >>> kmeans = KMeans(n clusters=3).fit(df)
    # Find the centroids of the clusters
    >>> centroids = kmeans.cluster centers
    # Get the associated cluster for each data record
    >>> kmeans.labels
     # Display the clusters contents and their centroids
     >>> plt.scatter(df['x'], df['y'], c= kmeans.labels_.astype(float), s=50,
     alpha=0.5)
     >>> plt.scatter(centroids[:, 0], centroids[:, 1], c='red', s=50)
    >>> plt.show()
```



Scatter Plot of Dataset With Clusters Identified Using K-means Clustering

2.2 GAUSSIAN MIXTURE MODELS CLUSTERING

- GMM: probabilistic generative models
- Each data point is assigned to the cluster returning the highest probability



An example of GMM clustering

- The optimum means (μ) and standard deviations (σ) are found by applying the Expectation maximization algorithm
- Advantages: efficient for datasets
 with complex data distribution
- Limitation: slow, heavy computation,
 and can get stuck in local-maximum

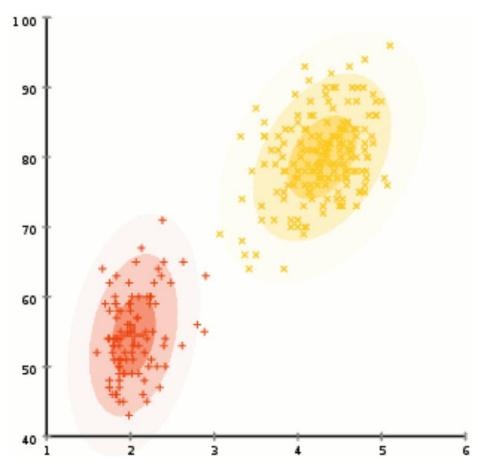
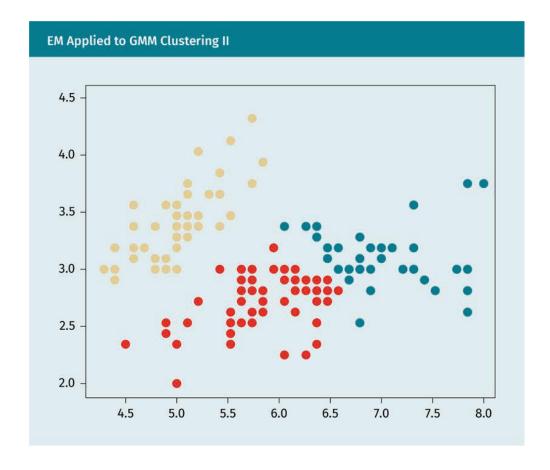


Illustration of expectation-maximization algorithm

2.2 GAUSSIAN MIXTURE MODELS CLUSTERING

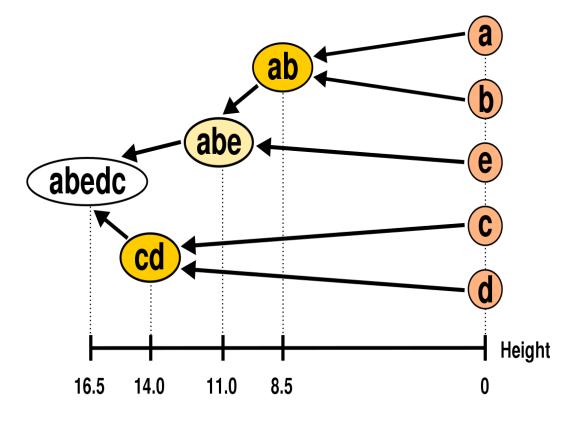
```
# GMM clustering with Python
>>> import numpy as np
>>> import pandas as pd
>>> import matplotlib.pyplot as plt
>>> from pandas import DataFrame
>>> from sklearn import datasets
>>> from sklearn.mixture import GaussianMixture
>>> iris = datasets.load_iris() # load the iris dataset
>>> X = iris.data[:,:2] # select first two columns
>>> d = pd.DataFrame(X) # turn it into a dataframe
>>> plt.scatter(d[0], d[1])
>>> plt.show() # plot the data
>>> gmm = GaussianMixture(n_components = 3)
>>> gmm.fit(d) # fit the data as a mixture of 3 Gaussians
>>> labels = gmm.predict(d) # predict the cluster of each data record
>>> print('Converged:',gmm.converged') # Check if the model has converged
>>> means = gmm.means_ # get the final "means" for each cluster
>>> covariances = gmm.covariances # get the final standard deviations
>>> d['labels']= labels
>>> d0 = d[d['labels']== 0]
>>> d1 = d[d['labels']== 1]
>>> d2 = d[d['labels']== 2]
>>> plt.scatter(d0[0], d0[1], c ='r')
>>> plt.scatter(d1[0], d1[1], c ='yellow')
>>> plt.scatter(d2[0], d2[1], c ='g')
>>> plt.show() # plot the data records in each clusters in different color
```



Scatter Plot of Dataset With Clusters Identified Using GMM Clustering

2.3 HIERARCHICAL CLUSTERING

- The clusters are built in a hierarchy as a **tree**
- Three levels of clusters:
 - Universe: root
 - Intermediate
 - Single-points. leaves

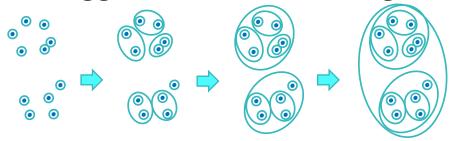


Tree-view (dendrogram) of hierarchical clustering

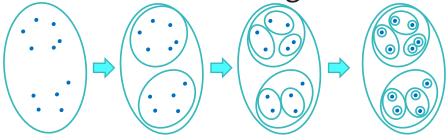
2.3 HIERARCHICAL CLUSTERING

- Two types of clustering:
 - Agglomerative: bottom-up
 - Divisive: top-down
- Data points are merged/split
 based on their distance
- Advantages: easy to implement, intuitive, no assumption
- Limitation: not suitable for large datasets, sensitive to outliers

Agglomerative clustering



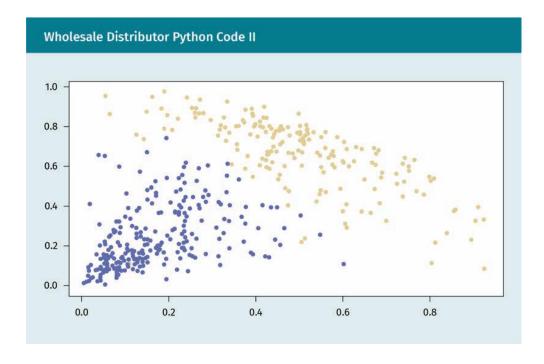
Divisive clustering



An example of hierarchical clustering

2.3 HIERARCHICAL CLUSTERING

```
# Agglomerative Clustering with Python
>>> import pandas as pd
>>> import numpy as np
>>> import matplotlib.pyplot as plt
>>> data = pd.read csv('Wholesale customers data.csv')
# Normalize the dataset to get all the features at the same scale
>>> from sklearn.preprocessing import normalize
>>> data scaled = normalize(data)
>>> data scaled = pd.DataFrame(data scaled, columns=data.columns)
# Draw the dendrogram to find the optimum number of clusters
>>> import scipy.cluster.hierarchy as shc
>>> plt.figure(figsize=(10, 7))
>>> plt.title("Dendrograms")
>>> dend = shc.dendrogram(shc.linkage(data_scaled, method='ward'))
>>> plt.show()
# From the dendrogram, we decide the optimum number of clusters is 2
# apply hierarchical clustering foe two clusters only
>>> from sklearn.cluster import AgglomerativeClustering
>>> cluster = AgglomerativeClustering(n clusters=2, affinity='euclidean', link
age='ward')
>>> cluster.fit predict(data scaled)
# visualize the two clusters
>>> plt.figure(figsize=(10, 7))
>>> plt.scatter(data_scaled['Milk'], data_scaled['Grocery'], c=cluster.labels )
>>> plt.show()
```

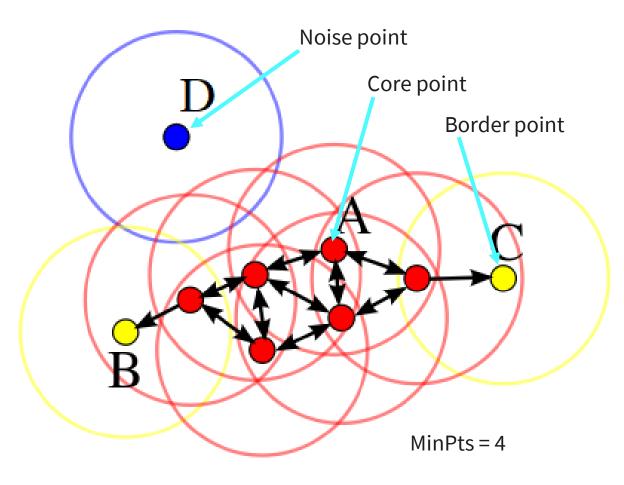


Scatter Plot of Dataset With Clusters Identified Using Agglomerative Clustering

2.4 DENSITY-BASED CLUSTERING

- Clusters are identified by grouping "dense" data points together, which permits:
 - the representation of arbitrarily shaped clusters,
 - the learning of outliers
- ε-Neighborhood of a point p in the dataset d:

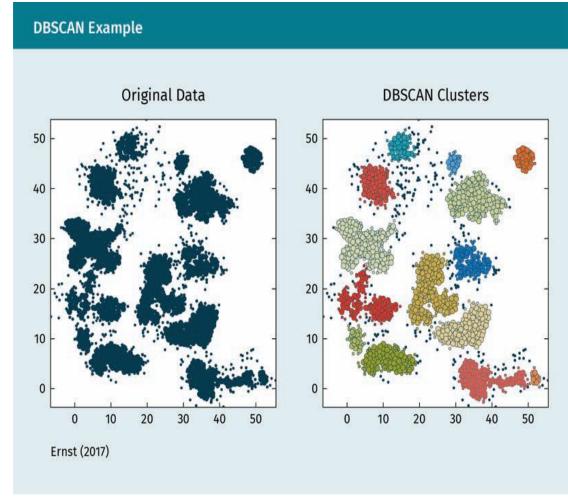
$$N(p) = \{ q \in d | dist(p, q) \le \varepsilon \}$$



Algorithm of Density-based Clustering

2.4 DENSITY-BASED CLUSTERING

- Density-based spatial clustering for data with noise:
 - 1. Initialize MinPts and ε
 - 2. Start at a random data point p
 - 3. Check if p is a core point
 - 4. Density-connected clustering
 - 5. Start at new point, repeat 3 & 4
- Advantages: estimation of cluster number is not required
- Limitation: ε and MinPts setting up could be tricky



An example of DBSCAN Clustering

2.4 DENSITY-BASED CLUSTERING

```
# DBSCAN clustering with Python
>>> import numpy as np
>>> from sklearn.cluster import DBSCAN
>>> from sklearn import metrics
>>> from sklearn.datasets.samples generator import make blobs
>>> from sklearn.preprocessing import StandardScaler
>>> from pylab import *
>>> import matplotlib.pyplot as plt
>>> # Generate sample data
>>> centers = [[1, 1], [-1, -1], [1, -1]]
>>> X, labels true = make blobs(n samples=750, centers=centers,
cluster std=0.4, random state=0)
>>> X = StandardScaler().fit transform(X) # standardize the dataset
>>> xx, yy = zip(*X)
>>> db = DBSCAN(eps=0.3, min_samples=10).fit(X) # Set up parameters
>>> core samples = db.core sample indices
>>> core_samples_mask = np.zeros_like(db.labels_, dtype=bool)
>>> core samples mask[db.core sample indices ] = True
>>> n clusters = len(set(labels))-(1 if -1 in labels else 0) # Nr of clusters
>>> labels = db.labels
>>> outliers = X[labels == -1] # find the outliers
>>> cluster1 = X[labels == 0] # Get the contents of each cluster
>>> cluster2 = X[labels == 1]
>>> cluster3 = X[labels == 2]
>>> unique labels = set(labels)
>>> colors = ['y', 'b', 'g', 'r']
```

```
>>> for k, col in zip(unique_labels, colors): if k == -1: col = 'k'
class_member_mask = (labels == k)

xy = X[class_member_mask & core_samples_mask]

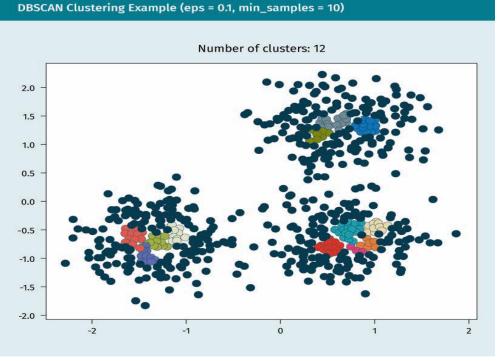
plt.plot(xy[:, 0], xy[:, 1], 'o', markerfacecolor=col,
markeredgecolor='k',markersize=6)

xy = X[class_member_mask & ~core_samples_mask]

plt.plot(xy[:, 0], xy[:, 1], 'o', markerfacecolor=col,
markeredgecolor='k', markersize=6)

>>> plt.title('number of clusters: %d' %n_clusters_)

>>> plt.show()
```



Using DBSCAN Clustering

REVIEW STUDY GOALS



- Know the definitions and terms used for clustering
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SESSION 2

TRANSFER TASK

Create the dataset using the following commands:

>>> from sklearn.datasets import make_blobs

>>> dataset,_ = make_blobs(n_samples=100, random_state=12)

Partition the data points into 3 clusters by using:

- 1. K-Means clustering
- 2. GMM clustering
- 3. Agglomerative clustering
- 4. DBSCAN with $\varepsilon=1$ and MinPts=10

TRANSFER TASK PRESENTATION OF THE RESULTS

Please present your results.

The results will be discussed in plenary.





- 1. Which of the following clustering algorithms suffers from the problem of convergence to local optima?
 - a) Agglomerative clustering
 - b) Only Expectation-Maximization clustering
 - C) K- Means clustering and expectation-maximization clustering
 - d) DBSCAN



2. For clustering to be applied, what is the minimum number of variables/ features required?

- a) 0
- b) 1
- c) 2
- d) 3



- 3. Is it possible that assignment of observations to clusters does not change between successive iterations in K-Means?
 - a) Yes
 - b) No
 - c) Cannot say
 - d) It depend on the number of data points

LEARNING CONTROL QUESTIONS



- 4. What could be the possible reason(s) for producing two different dendrograms using the agglomerative clustering algorithm for the same dataset?
 - a) Proximity function used
 - b) Number of data points used
 - c) Number of variables used
 - d) All these options are possible

LEARNING CONTROL QUESTIONS



- 5. K-Means clustering with Euclidean distance measure produces which kind of cluster shapes?
 - a) Spherical
 - b) Ellipsoid
 - c) Arbitrary
 - d) Cubic

LIST OF SOURCES

Text:

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File:DBSCAN-Illustration.svg. (2020, October 7). In Wikimedia Commons, the free media repository. Retrieved, January 25, 2023, from https://commons.wikimedia.org/w/index.php?title=File:DBSCAN-Illustration.svg. (2020, October 7). In Wikimedia Commons, the free media repository. Retrieved, January 25, 2023, from https://commons.wikimedia.org/w/index.php?title=File:DBSCAN-Illustration.svg. (2020, October 7). In Wikimedia Commons, the free media repository. Retrieved , January 25, 2023, from https://commons.wikimedia.org/w/index.php?title=File:DBSCAN-Illustration.svg. (2020, October 7). In Wikimedia Commons, the free media repository. (2020, October 7). In Wikimedia Commons wikimedia.org/w/index.php?title=File:DBSCAN-Illustration.svg.

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