Data Science SS20



Machine Learning II

Unsupervised Learning: Clustering

Machine Learning II



Outline

- Clustering Algorithms
 - K-Means
 - DBSCAN

Introduction to ML



Basic Types of Machine Learning Algorithms

Supervised Learning

Unsupervised Learning

Reinforcement Learning

- NO Labeled data
- NO Direct and quantitative evaluation
- Explore structure of data

Clustering Algorithms



Definition

Cluster analysis or **clustering** is the task of **grouping a set of objects** in such a way that objects in the same group (called a cluster) are **more similar** (in some sense) to each other than to those in other groups (clusters). [Wikipedia]

Introduction to ML



Unsupervised Learning

Data without "labels" (x_1, x_2, \dots, x_n)

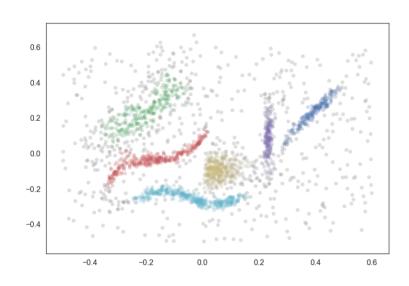
- Clustering
- Outlier Detection (e.g. Defect or Intrusion detection)

Clustering Algorithms



Introduction

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Example 2d data set

Clustering Algorithms



Motivation

- Standard technique for data exploration and anlysis
- Objective: find inherent structures in data
- Just like Multivariate Statistics
 - For high dimensional data
 - Geometric (manifold) and statistical motivations



Definition:

Given a set of observations (x_1, x_2, \dots, x_n)

where each observation is a *d*-dimensional real vector, **K-Means** clustering aims to partition the *n* observations into $k \le n$

sets
$$S = \{S_1, S_2, \dots, S_k\}$$

so as to minimize the within-cluster sum of squares:

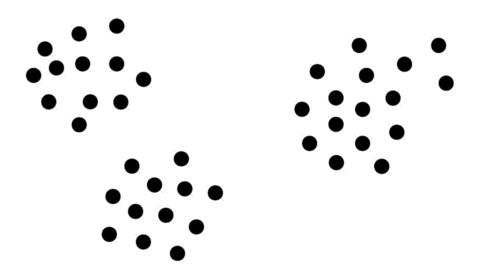
$$\arg \min[S] \sum_{i=1}^{k} \sum_{x \in S_i} ||x - \mu_i||$$



Intuition:

$$\arg \min[S] \sum_{i=1}^{k} \sum_{x \in S_i} ||x - \mu_i||$$

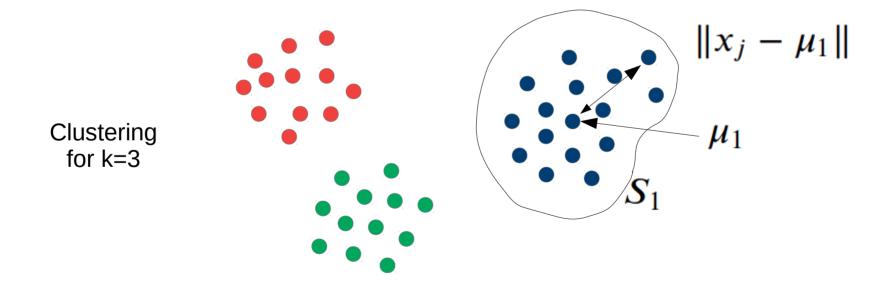
Data:





Intuition:

$$\arg \min[S] \sum_{i=1}^{k} \sum_{x \in S_i} ||x - \mu_i||$$





Basic Algorithm:

Init (t=1): Select *k* random cluster centers

$$\mu_1^{(1)} := x_{r1}, \mu_2^{(1)} := x_{r2}, \dots, \mu_k^{(1)} := x_{rk} \quad for \quad x_{rj} \in X$$

Repeat *n* times:

1. For step t: Assign all samples to "closest" center

$$S_i^{(t)} = \{x_p : \|x_p - \mu_i^{(t)}\|^2 \le \|x_p - \mu_j^{(t)}\|^2 \quad \forall j, 1 \le j \le k\}$$

2. Re-Compute cluster centers

$$\mu_i^{(t+1)} = \frac{1}{|S_i^{(t)}|} \sum_{x_j \in S_i^{(t)}} x_j$$



Intuition:

$$\mu_1^{(1)} := x_{r1}, \mu_2^{(1)} := x_{r2}, \dots, \mu_k^{(1)} := x_{rk} \quad for \quad x_{rj} \in X$$

Random!

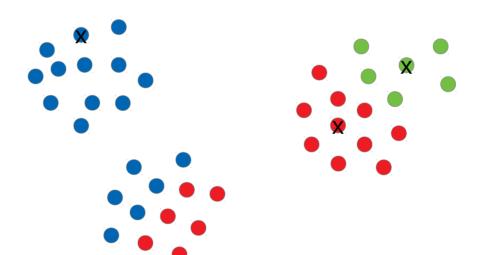
Init:



Intuition:

$$S_i^{(t)} = \{x_p : \|x_p - \mu_i^{(t)}\|^2 \le \|x_p - \mu_i^{(t)}\|^2 \quad \forall j, 1 \le j \le k\}$$

Step 1:

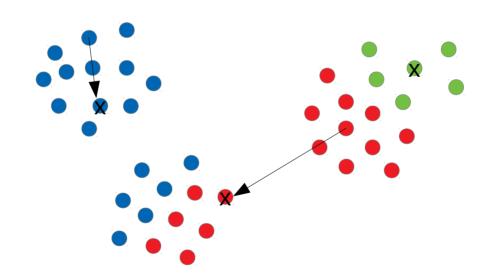




Intuition:

$$\mu_i^{(t+1)} = \frac{1}{|S_i^{(t)}|} \sum_{x_j \in S_i^{(t)}} x_j$$

Step 2:

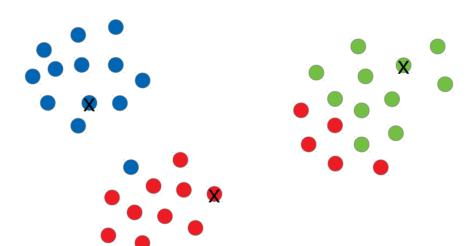




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Step 1:

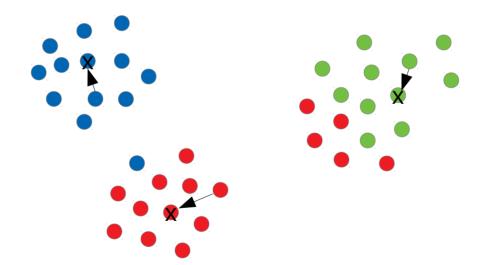




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Step 2:

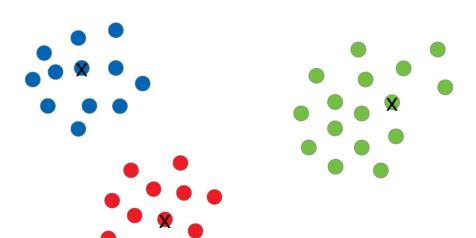




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Step 1:

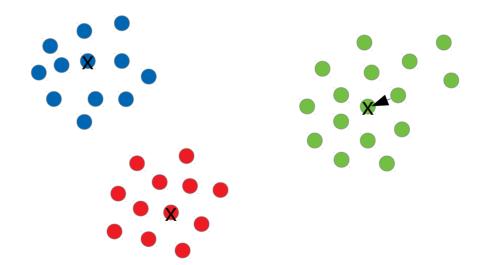




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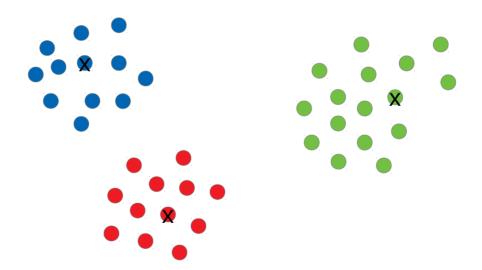
Step 2:





Intuition: Convergence after fix t or fix means

Step 1:



t=5,6,7,...



Evaluation:

- Very simple but effective clustering algorithm
- Advantages:
 - Easy to implement
 - Easy to parallelize
- Disadvantages
 - Need to know k in advance (or search for best k)
 - High complexity: HP-hard (exponential in data dimension)
 - Problem with non-blob shaped (non-convex) clusters



Evaluation:



More practical examples in the Lab session....



Definition

Density-based spatial clustering of applications with noise (DBSCAN) is a density-based clustering **non-parametric** algorithm:

Given a set of points in some space, it groups together points that are closely packed together (points with many nearby neighbors), marking as outliers points that lie alone in low-density regions (whose nearest neighbors are too far away).



Definition

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In contrast to K-Means, the number of clusters is not given



Definitions in DBSCAN

Given a set of observations (x_1, x_2, \dots, x_n) we need:

Some distance function on the data: $d(x_i, x_j)$

Parameter: radius \mathcal{E}

Set of Neighbors: $\mathbb{N}(x_i)$

Parameter: Minimum number of Neighbors n_{ε}



Definitions in DBSCAN

Density at a data sample: number of neighbors in radius

$$\mathbb{d}(x_i) := |\mathbb{N}(x_i)| = |\{x_p : d(x_p, x_i) \le \varepsilon\}|$$

Core samples: all samples with a density higher than a threshold

$$\{x_p: \mathbb{d}(x_p) \geq n_{\varepsilon}\}$$

Reachable samples: all samples with at least one neighbor

$$\{x_p: d(x_p) \ge 1\}$$



Definitions in DBSCAN

Outlier samples: All samples without neighbors

$$\{x_p: \mathbb{d}(x_p)=0\}$$



Basic Algorithm

Init: mark all samples as core, reachable or outlier

Remove outlier

For all core samples: choose next random core sample and recursively merge it's

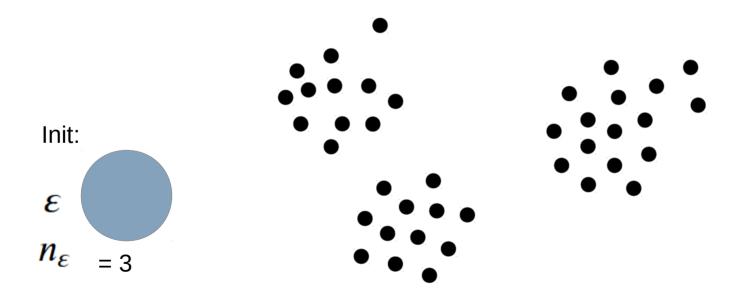
neighborhood with all neighbors that are also core samples.

Increment Cluster ID.

For all reachable samples: assign to closest cluster

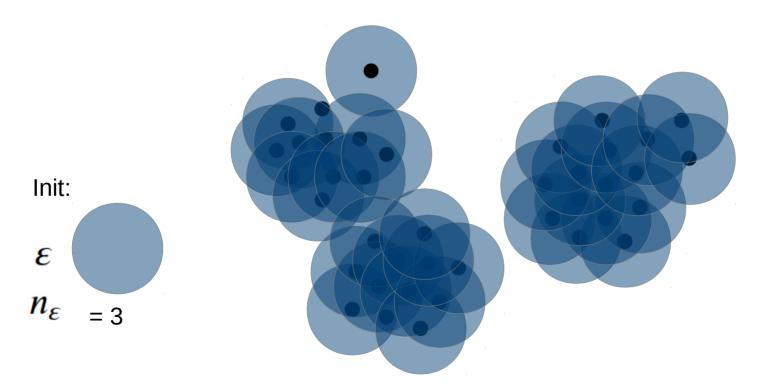


Intuition:



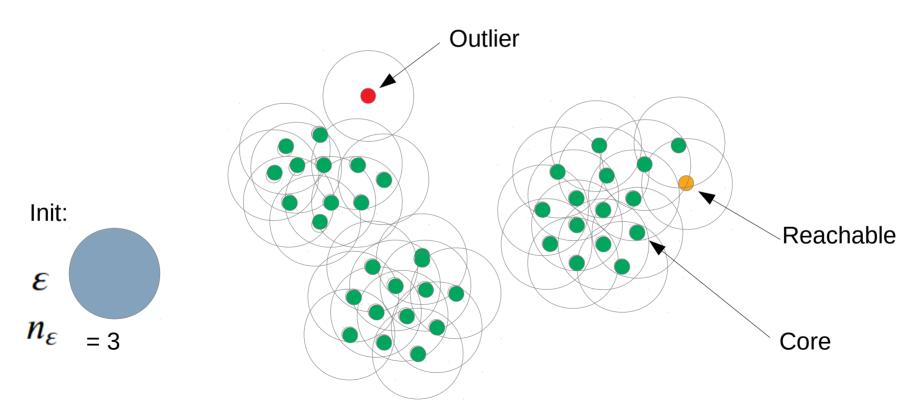


Density:



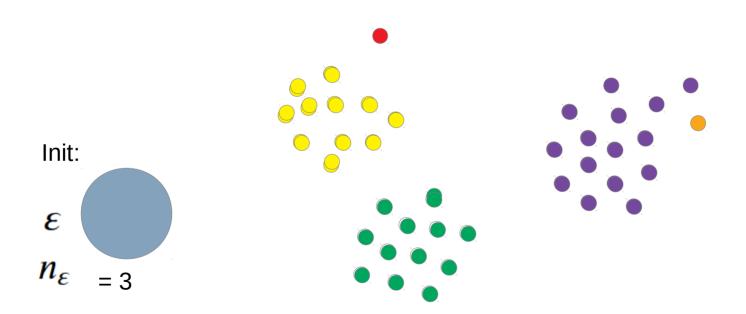


Core - Reachable - Outlier:



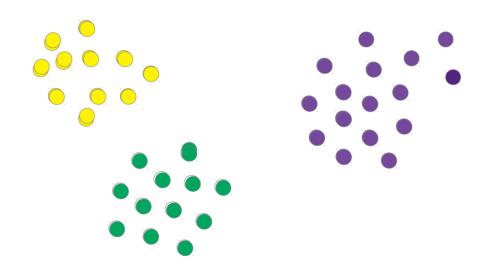


Merge:





Final:





Evaluation:

- Very common Clustering Algorithm
- Advantages:
 - Does not need number of clusters
 - Works well for non convex clusters
 - Fast implementation possible (R-Trees)
- Disadvantages
 - Has two hyper-parameters to optimize
 - Fails on data with high variance in density
 - Not deterministic



Evaluation:



More practical examples in the Lab session....

Clustering Algorithms: Evaluation



How to evaluate clustering:

Visually → use dimension reduction techniques to visualize 2d or 3d

Clustering Algorithms: Evaluation



How to evaluate clustering:

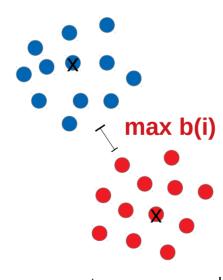
- Visually → use dimension reduction techniques to visualize 2d or 3d
- Quantitative quality measures (what is a good cluster?)
 - Low intra cluster variance

$$a(i) = rac{1}{|C_i|-1} \sum_{j \in C_i, i
eq j} d(i,j)$$

High extra cluster variance

$$b(i) = \min_{i
eq j} rac{1}{|C_j|} \sum_{j \in C_i} d(i,j)$$

For each data point $i \in C_i$ (data point i in the cluster C_i)



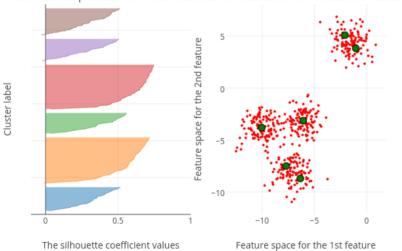
Clustering Algorithms: Evaluation



Silhouette Diagrams: finding the best number of clusters

Silhouette analysis for KMeans clustering on sample data with n_clusters = 6

The silhouette plot for the various clustersThe visualization of the clustered data.



Silhouette analysis for KMeans clustering on sample data with n_clusters = 2

The silhouette plot for the various clustersThe visualization of the clustered data.



[plots: https://plot.ly/scikit-learn/plot-kmeans-silhouette-analysis/]

Discussion



There are many more Clustering algorithms available...

 $\dots \rightarrow$ lab session.