Unsupervised Classification

A partioning-based clustering algorithm: K-means

Data Analysis – juliette.chevallier@insa-toulouse.fr INSA Toulouse, Applied Mathematics, 4th year

1. K-means type methods

- 1.1 General principle
- 1.2 Choice of hyper-parameters

2. DBSCAN: Density-Based Spatial Clustering of Applications with Noise

- 2.1 Principle of DBSCAN methods
- 2.2 Choice of hyper-parameters

K-means type methods

1.1 General principle

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Introduction

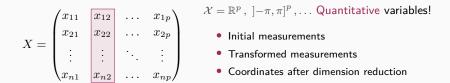
We observe n individuals described by p variables: $x_i = (x_{i1}, x_{i2}, \ldots, x_{ip}) \in \mathcal{X}$

 $X = \begin{pmatrix} x_{11} & x_{12} & \dots & x_{1p} \\ x_{21} & x_{22} & \dots & x_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ x_{21} & x_{22} & \dots & x_{2p} \end{pmatrix}$ • Initial measurements
• Transformed measurements
• Coordinates after dimension reduction

 $\mathcal{X} = \mathbb{R}^p$, $[-\pi, \pi]^p$,... Quantitative variables!

Introduction

We observe n individuals described by p variables: $x_i = (x_{i1}, x_{i2}, \dots, x_{ip}) \in \mathcal{X}$



- Let d be the Euclidean distance: $d(x,y) = \|x-y\|^2$
- **Goal** of *k*-means algorithm: Find a partition of the individuals that minimizes the intra-class inertia, *i.e.* the within-cluster sum of squares (WCSS)

$$\widehat{\mathcal{P}}_{K}^{k-means} \in \operatorname{argmin}_{\mathcal{P}_{K}} \sum_{k=1}^{K} \sum_{i \in \mathcal{C}_{k}} d(\mu_{k}, x_{i})^{2} \quad \text{where} \quad \mu_{k} = \frac{1}{|\mathcal{C}_{k}|} \sum_{i \in \mathcal{C}_{k}} x_{i}$$

$$\mathcal{P}_K = \{\mathcal{C}_1, \dots, \mathcal{C}_K\}$$
 partition of $\llbracket 1, n \rrbracket$

Initialization:

- Choice of the number of classes *K*
- Choice of K initial centroids $\mu_1^{(0)}, \, \dots, \, \mu_K^{(0)}$

Iteration *t*: Repeat:

Allocation update: Point *i* allocated to the nearest centroid

$$i \in \mathcal{C}_k^{(t)}$$
 such that $dig(x_i,\,\mu_k^{(t-1)}ig) = \min_{\ell \in \llbracket 1,K
rbrace} dig(x_i,\,\mu_\ell^{(t-1)}ig)$

$$\mu_{k}^{(t)} = \frac{1}{|\mathcal{C}_{k}^{(t)}|} \sum_{i \in \mathcal{C}_{k}^{(t)}} x_{i}$$

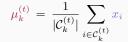
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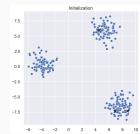
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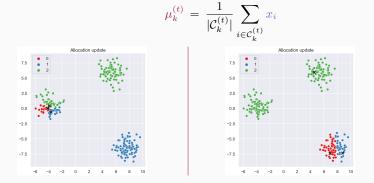
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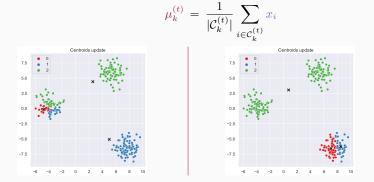
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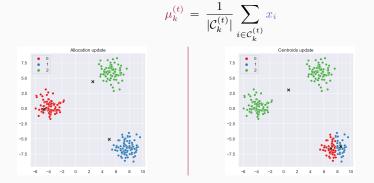
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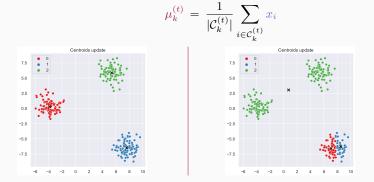
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Assymptotic Behavior of the *k*-means Algorithm

Proposition

The intra-class inertia $I_{Intra} \left(\mathcal{P}_{K}^{(t)}
ight)$ decreases with each step.

 \sim Convergence of the k-means algorithm towards a local minimum of the intra-class inertia.

Sketch of the proof: (Demonstration will be covered in tutorials)

Two key arguments:

1. If point i goes from $\mathcal{C}_k^{(t-1)}$ to $\mathcal{C}_\ell^{(t)}$, then:

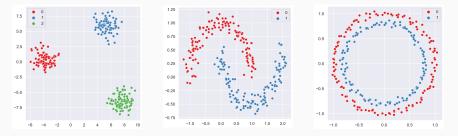
$$dig(x_i,\mu_\ell^{(t)}ig)^2\leqslant dig(x_i,\mu_k^{(t-1)}ig)^2$$

2. $\mu_k^{(t)}$ being the gravity center of $\mathcal{C}_k^{(t)}$,

$$\sum_{i \in \mathcal{C}_k^{(t)}} dig(x_i, \mu_k^{(t)}ig)^2 \leqslant \sum_{i \in \mathcal{C}_k^{(t)}} dig(x_i, \mu_k^{(t-1)}ig)^2$$

Pros: • Relatively efficient (fast),

- Tends to reduce intra-class inertia at each iteration,
- Forms compact and well-separated classes.

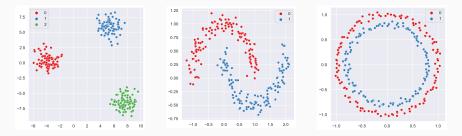


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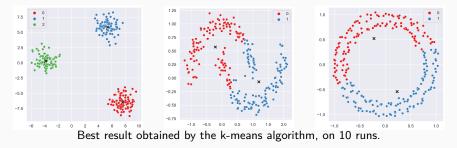
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- Requires the notion of center of gravity,
- Influence of outliers (due to averaging),
- Not suitable for non-convex classes.



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Variants of the *k*-means

k-medoids: More efficient on small dataset, more robust in the presence of noise or outliers.

Idea: Use medoids instead of centroids, *i.e.* points from X.

$$\nu_k \in \operatorname*{argmin}_{y \in X} \sum_{i \in \mathcal{C}_k} d(y, x_i)^2$$

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k-modes: For qualitative data.

(i) Modify the dissimilarity measure to handle qualitative data.

$$d(x_a, x_b) = \sum_{j=1}^{p} \frac{n_{aj} + n_{bj}}{n_{aj} \times n_{bj}} \mathbb{1}_{\{x_{aj} \neq x_{bj}\}}$$

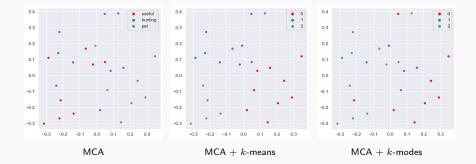
where $n_{aj} = \#\{i \in [\![1, n]\!] \mid x_{aj} = x_{ij}\}.$

(ii) Use of modes instead of centers of gravity.

Remark: For **qualitative** data, we can also use *k*-means algorithm on multiple correspondence analysis (MCA).

Dogs Breeds: *k*-modes vs *k*-means+MCA

Analysis of 27 dog breeds based on 6 descriptive qualitative: size (3), weight (3), velocity (3), intelligence (3), affection (2) and aggressiveness (2).



K-means type methods

1.1 General principle

1.2 Choice of hyper-parameters

Choice of the Number of Classes K



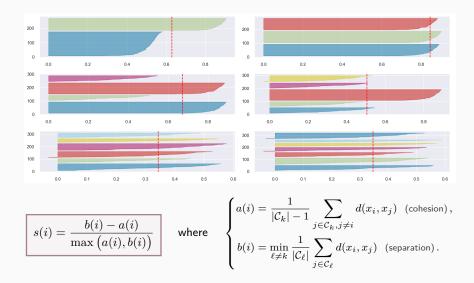
- Elbow method for the intra-class inertia IIntra
 - For each value of $K \in \{2, \ldots, K_{max}\}$, we obtain a classification \mathcal{P}_K ,
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 - We select the one where we observe a significant jump in intra-class inertia.
- Other criteria based on inertia
 - **R-Square**: $RSQ(K) = \frac{I_{Inter}(\mathcal{P}_K)}{I_{Tot}} = 1 \frac{I_{Intra}(\mathcal{P}_K)}{I_{Tot}}$ K: Elbow on the RSQ curve.
 - Semi-Partial R-Square: $SPRSQ(K) = \frac{I_{Inter}(\mathcal{P}_K) I_{Inter}(\mathcal{P}_{K-1})}{I_{Tot}}$ K: largest reduction of the QSPRS.
 - Calinski-Harabasz (CH): $PseudoF(K) = \frac{I_{Inter}(\mathcal{P}_K)}{I_{Intra}(\mathcal{P}_K)} \times \frac{n-K}{K-1}$ K: Peak on the CH curve.

Silhouette Score



- The better the classification, the closer the silhouette score is to 1
- Negative score in case of bad classification

Choice of Initial Centriods



- A judicious choice can favour the convergence towards a global minimum!
- Selection based on additional knowledge, or on a preliminary study of the data: histograms, *etc.*
- Repeat the method N times, and select the partition \mathcal{P}_K with the lowest intra-class inertia.

Choice of Initial Centriods [Arthur and Vassilvitskii, 2006]

 Forgy Initialization
 Random Partition Method
 k-means++

Any K points from the data, at random.

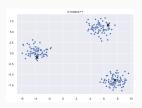
- Random assignment of a cluster ID to each data point,
- Average by ID of the points.

- Choose a random point,
- Next centroid so that it lies at a large distance from the first one, with high probability: Sample a point from a probability distrib. proportional to the distance to the first centroid,
- Remaining pts generated by a probability disctrib. proportional to the distance of each point from its nearest

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 Image: state stat



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Not a good choice for *k*-means!

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DBSCAN: Density-Based Spatial Clustering of Applications with Noise

2.1 Principle of DBSCAN methods2.2 Choice of hyper-parameters

• Two key parameters:

ε: The distance that specifies the neighborhoods.
 Two points are considered to be neighbors if the distance between them are less than or equal to ε.

MinPts: Minimum number of data points to define a cluster.

- Algorithmic steps for DBSCAN clustering:
 - 1. Arbitrarily picking up a point in the dataset (until all points have been visited).
 - 2. If there are at least MinPts points within a radius of ε to the point then we consider all these points to be part of the same cluster.
 - 3. The clusters are then expanded by recursively repeating the neighborhood calculation for each neighboring point

See aaronscotthq.com/2020-05-28-scott_dbscan for an animation.

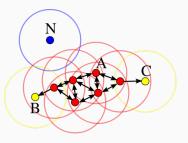
Points classified as core point, border point, or outlier:

Core point: There are at least MinPts number of points (including itself) in its ε -neighborhood:

```
\#\{i\in \llbracket 1,n\rrbracket\mid d(x_a,x_i)<\varepsilon\}\geqslant MinPts\,.
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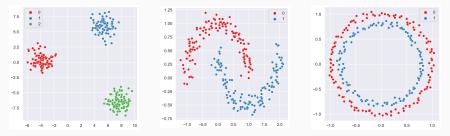
Border point: Belongs to the ε-neighborhood of a core point, but is not a central point (not enough dense neighborhood).

> Outlier: Neither a central point nor a border point (In particular, not classified).



Pros: • Does not require to specify number of clusters beforehand,

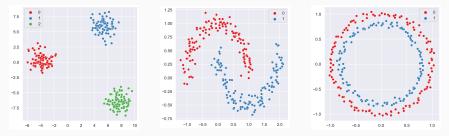
- Performs well with clusters of arbitrary shapes,
- Robust to outliers and able to detect them.



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- **Cons:** In some cases, determining an appropriate neighborhood distance ε is not easy and requires domain knowledge,
 - Not well suited if the clusters are very different from each other in terms of intra-cluster densities.

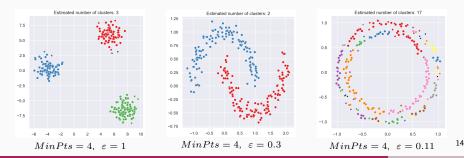
Characteristics of the clusters defined by the combination $\varepsilon - MinPts$, and we pass only one couple $\varepsilon - MinPts$ to the algorithm.



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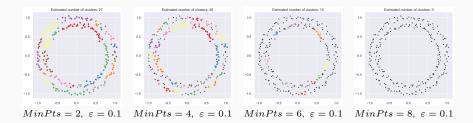


DBSCAN: Density-Based Spatial Clustering of Applications with Noise

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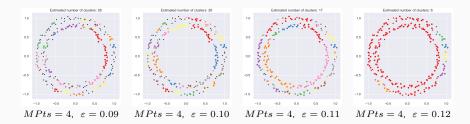
Determining Minimum Samples MinPts



MinPts: to be defined using domain knowledge and familiarity with the data set.

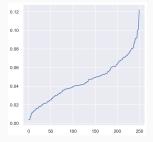
- $MinPts \ge p+1$,
- The larger the data set, the larger *MinPts*,
- The noisier the data set, the larger *MinPts*,
- For 2d data, use DBSCAN's default value of MinPts = 4 [Ester et al., 1996]
- For more than 2d data, choose MinPts = 2p [Sander et al., 1998], or MinPts = ln(n).

Determining the distance ε



 ε : Based on the average distance between each point and its MinPts nearest neighbors (MinPts - NN distance)

- For *each* point of the dataset, compute its *MinPts* - *NN* distance,
- Plot this distances in ascending,
- We choose ε as the value of the MinPts NN distance where an "elbow" is observed (maximum curvature).



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