

MLRF Lecture 02

J. Chazalon, LRDE/EPITA, 2019

Clustering

Lecture 02 part 03

Clustering: finding groups in data (1/2)

Many techniques:

Useful for Practice session 2

- Connectivity models: hierarchical clustering, ...
cluster = set of neighbors
- Centroid models: k-means, ...
cluster = centroid point
- Distribution model: Gaussian mixtures models est. w. Expectation Maxim., ...
cluster = statistical distribution (center + spread in each direction)
- Density models: DBSCAN, ...
cluster = dense region
- Graph-based models: HCS (Highly Connected Subgraphs) algorithms, ...
cluster = clique in the graph
- ...

Useful for Practice session 4
(eventually)

Clustering: finding groups in data (2/2)

Always the same goal:

- Minimise the differences between elements within the same cluster
- Maximise the differences between elements within different clusters

Number of clusters:

- Many methods require to choose it beforehand
- Several techniques to adjust the number of clusters automatically

Outliers rejection:

- Some techniques do not assign lonely points to any cluster

⇒ Focus on HAC and K-Means today

Hierarchical Agglomerative Clustering (HAC)

Hierarchical Agglomerative Clustering

Algorithm

Initialization:

- Create a cluster from each element

While more than 1 cluster:

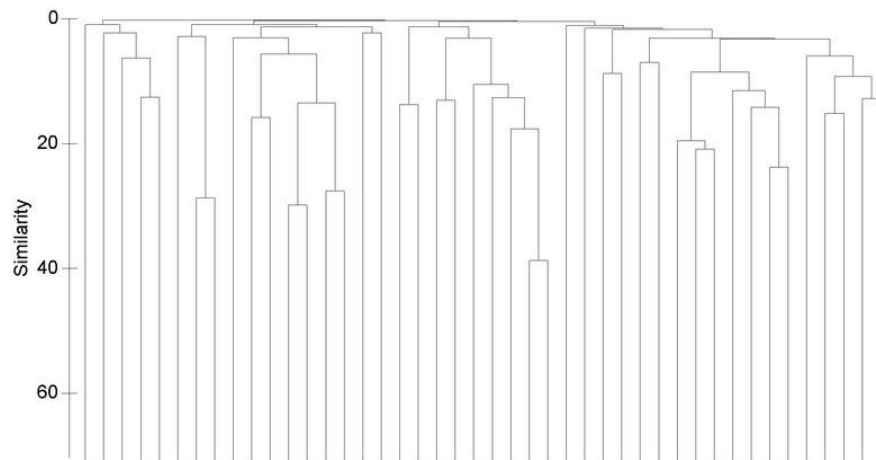
- Merge the two closest clusters

Challenge: distance between clusters

- New center = mean of points
- or distance = maximal distance
- or...

Not suitable for large datasets

Output: Dendrogram



K-Means

K-Mean clustering (again) (1/3)

The K-means algorithm aims to choose centroids that minimise the inertia, or within-cluster sum-of-squares criterion:

$$\sum_{i=0}^n \min_{\mu_j \in C} (||x_i - \mu_j||^2)$$

- ie it does not maximizes inter-cluster distance
- ie it puts centers so as to get the best coverage (may not be on a density peak!)

K-Mean clustering (again) (2/3)

Algorithm

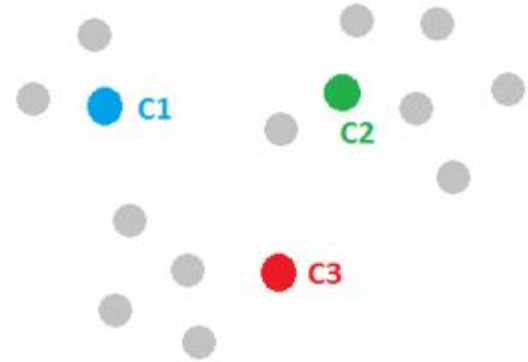


K-Mean clustering (again) (2/3)

Algorithm

Initialization:

- (randomly) select cluster centers

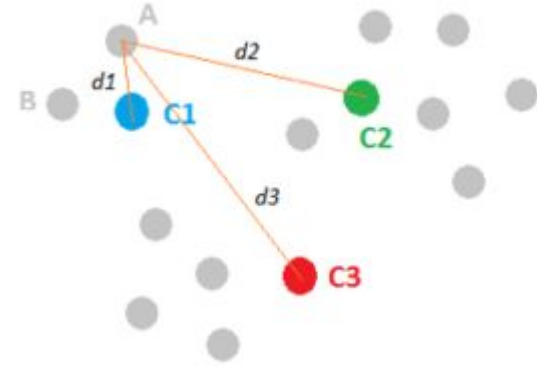


K-Mean clustering (again) (2/3)

Algorithm

Initialization:

- (randomly) select cluster centers
- Calculate distance points \leftrightarrow centers

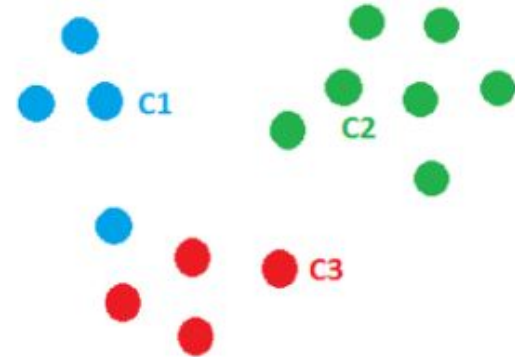


K-Mean clustering (again) (2/3)

Algorithm

Initialization:

- (randomly) select cluster centers
- Calculate distance points \leftrightarrow centers
- Assign each point to closest center

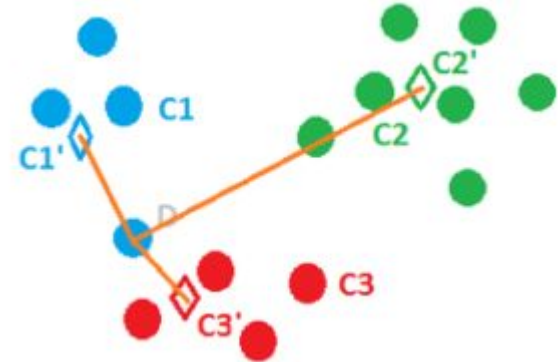


K-Mean clustering (again) (2/3)

Algorithm

Initialization:

- (randomly) select cluster centers
- Calculate distance points \leftrightarrow centers
- Assign each point to closest center
- Update cluster centers: avg of points



K-Mean clustering (again) (2/3)

Algorithm

Initialization:

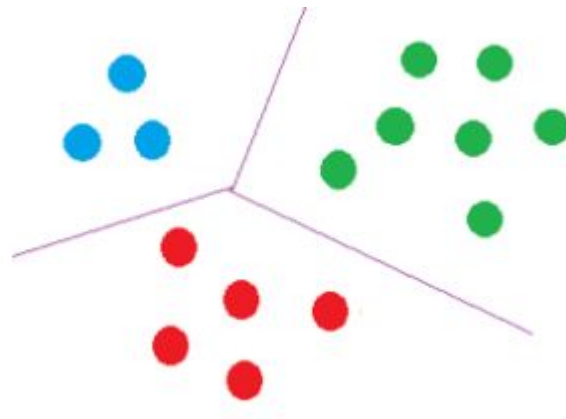
- (randomly) select cluster centers

Loop until converged:

- Calculate distance points \leftrightarrow centers
- Assign each point to closest center
- Update cluster centers: avg of points

Result: centroid centers

- local maximas
- tessellation / Voronoi set over the dataset



K-Mean clustering (again) (3/3)

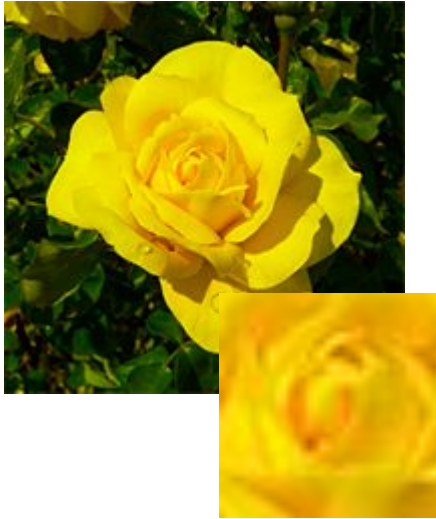
The previous algorithm is called “**Batch K-Means**”, or simply “**K-Means**”, because it considers the **whole the dataset at each iteration**.

Batch K-Means is not only **sensible to outliers** and **initialization**, it is also **very slow** to compute on large datasets. (I got OOM errors with the *Twin it!* poster!!)

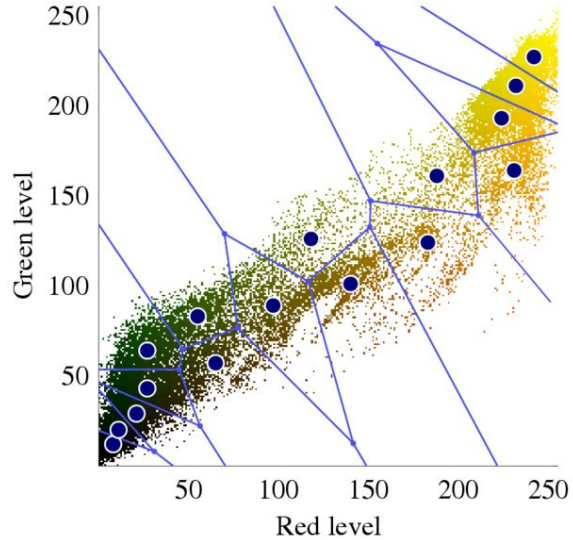
It is possible to avoid this speed / memory issue by **randomly sampling the dataset at each step**.

- Results are only slightly worse
- Speed and memory requirements make it usable on bigger datasets
- This approach is call “**Online K-Means**” or “**MiniBatch K-Means**”

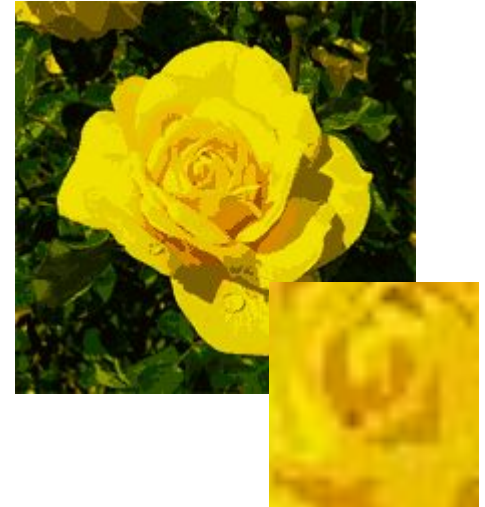
Application: Color quantization



Original RG image
(no blue channel)



Color quantization showing
original colors, target colors
and boundaries (Voronoi
cells here)



Color-indexed image
(no dithering)

Many clustering techniques to play with!

https://scikit-learn.org/stable/auto_examples/cluster/plot_cluster_comparison.html

