# CPSC 340: Machine Learning and Data Mining

# Admin

- Tutorials today.
- Office hours tomorrow
- Assignment 2 due Friday.

- Steps of k-means++:
  - 1. Select initial mean  $\mu_1$ , from among the object  $x_i$ .
  - 2. Compute distance  $d_{ic}$  of object  $x_i$  to each mean  $\mu_c$ .

$$d_{ic} = ||X_i - M_c|| = \sqrt{\frac{2}{2}(x_{ij} - M_c)^2}$$

3. For each object set d<sub>i</sub> to the minimum distance across all clusters c.

$$d_i = \min \{ d_i \}$$

- 4. Choose next mean by sampling proportional to  $(d_i)^2$ .
- 4. Choose next mean by sampling proportion to (o),  $p_i \neq d_i^2 = p_i = \frac{d_i^2}{\frac{1}{2}d_i^2}$ 5. Stop when we have k means, otherwise return to 2.
- Expected approximation ratio is O(log(k)).























• K-means clusters are formed by the intersection of half-spaces.



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- Intersection of half-spaces forms a convex set:
  - Line between any two points in the set stays in the set.





### K-Means with Non-Convex Clusters



### K-Means with Non-Convex Clusters



K-means cannot separate non-convex

ttps://corelifesciences.com/human-long-non-coding-rna-expression-microarray-service.htm

### K-Means with Non-Convex Clusters



# **Application: Elephant Range Map**

- Find habitat area of African elephants.
  - Useful for assessing/protecting population.
- Build clusters from observations of locations.
- Clusters are non-convex:
  - affected by vegetation, relief, rivers, water access.
- We do not want a partition:
  - Some regions should not have a cluster.



# Motivation for Density-Based Clustering

- **Density-based clustering** is a non-parametric clustering method:
  - Clusters are defined by connected dense regions.
    - Become more complicated the more data we have.
  - Data points in non-dense regions are not assigned a cluster.



http://www.defenders.org/elephant/basic-facts

# **Other Potential Applications**

- Where are high crime regions of a city?
- Where should taxis patrol?
- Where does Iguodala make/miss shots?
- Which products are similar to this one?
- Which pictures are in the same place?
- Where can protein 'dock'?

https://en.wikipedia.org/wiki/Cluster\_analysis https://www.flickr.com/photos/dbarefoot/420194128/ http://letsgowarriors.com/replacing-jarrett-jack/2013/10/04/ http://www.dbs.informatik.uni-muenchen.de/Forschung/KDD/Clus





DRE IGUODALA



- **Density-based clustering** algorithm (DBSCAN) has two parameters:
  - Radius: minimum distance between points to be considered 'close'.
  - MinPoints: number of 'close' points needed to define a cluster.



- Pseudocode for DBSCAN:
  - For each example x<sub>i</sub>:
    - If x<sub>i</sub> is already assigned to a cluster, do nothing.
    - If  $x_i$  is not core point (less than minPoints neighbours with distance  $\leq$  'r'), do nothing.
    - If x<sub>i</sub> is a core point, expand cluster.
  - Expand cluster function:
    - Assign all x<sub>i</sub> within distance 'r' of core point x<sub>i</sub> to cluster.
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boundary point to boundary point to

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### **Density-Based Clustering Issues**

• Some points are not assigned to a cluster.

– Good or bad, depending on the application.

- Sensitive to the choice of radius and minPoints.
- Ambiguity of 'non-core' (boundary) points:
  They could be assigned more than once.
- Other than this ambiguity, not sensitive to initialization.
- Assigning new points to clusters is expensive.
- In high-dimensions, need a lot of points to 'fill' the space.

# Summary

- 1. K-means++: randomized initialization with good expected performance.
- Shape of K-means clusters: intersection of half-spaces => convex sets.
- **3.** Density-based clustering: useful for finding non-convex connected clusters.
- 4. DBSCAN algorithm: assign points in dense regions to same cluster.
- Next time:
  - Dealing with clusters of different densities.