机器学习引论

彭玺

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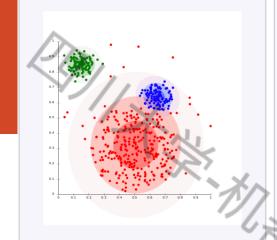
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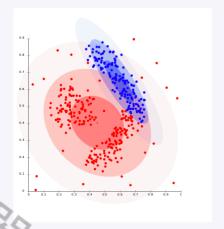
提纲

- . Review
- 二 . k-means clustering
- 三 . k-medoids clustering
- 四. Mixture of Gaussian

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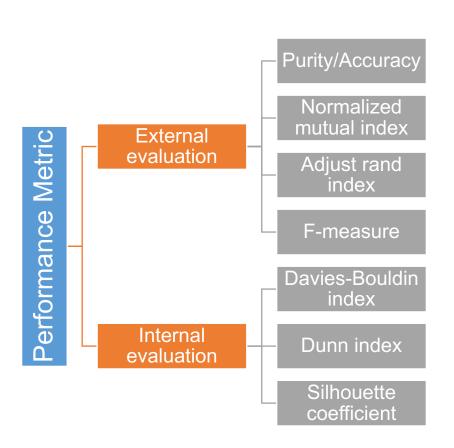
Problem Statement:

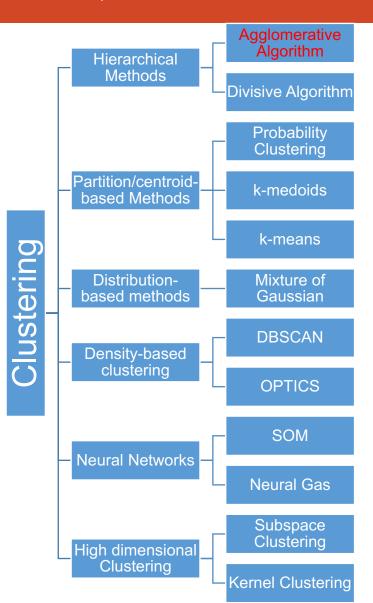
- Given a set of data points, group them into multiple clusters so that:
 - points within each cluster are similar to each other
 - points from different clusters are dissimilar

Challenges 1 (key problem of clustering analysis): The major difficulty is that the label is unknown so that the within-/between- class scatter is unavailable.

Challenges 2 (high-dimensional clustering analysis): Usually, points are in a high-dimensional space, and similarity is defined using a distance measure

■ Euclidean, Cosine, Jaccard, edit distance, ...





- Agglomerative (Bottom-up)
 - Compute all pair-wise patternpattern similarity coefficients
 - Place each of *n* patterns into a class of its own
 - Merge the two most similar clusters into one
 - Replace the two clusters into the new cluster
 - Re-compute inter-cluster similarity scores w.r.t. the new cluster
 - Repeat the above step until there are *k* clusters left (*k* can be 1)

Agglomerative (Bottom up)



Agglomerative (Bottom up)

1st iteration



Agglomerative (Bottom up)

 2^{nd} iteration

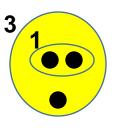






Agglomerative (Bottom up)

3rd iteration

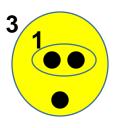




• •

Agglomerative (Bottom up)

4th iteration









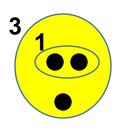


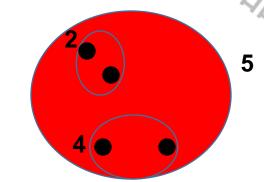


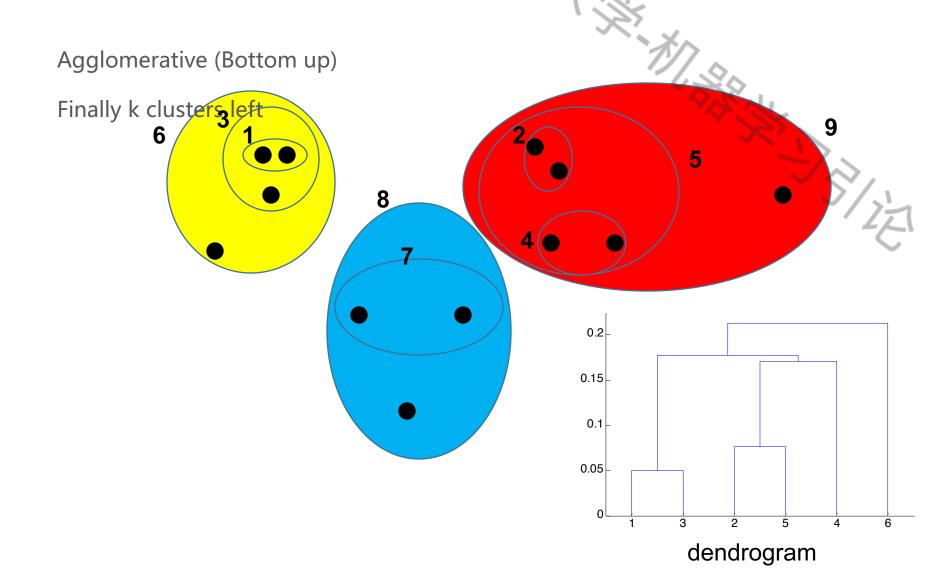


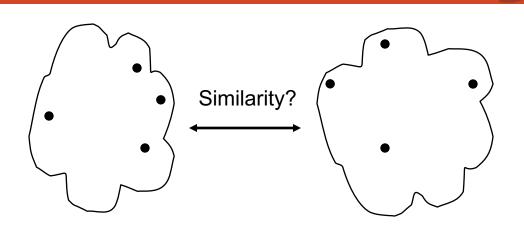
Agglomerative (Bottom up)

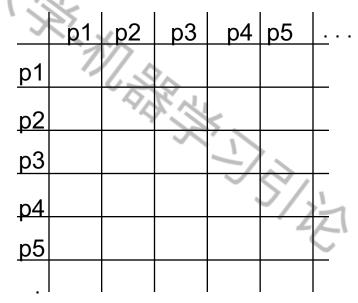
5th iteration



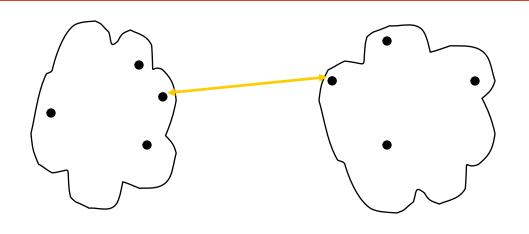


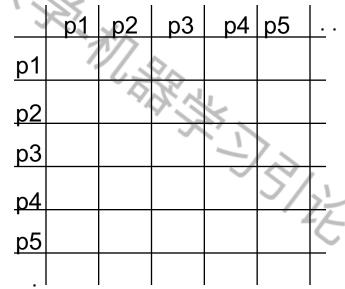




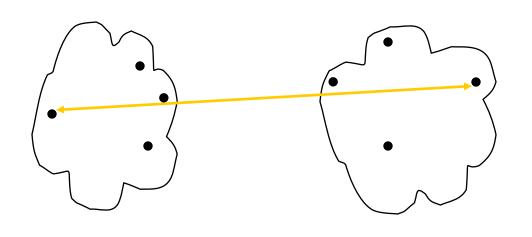


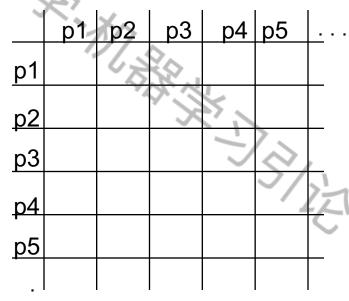
- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
 - Ward's Method uses squared error



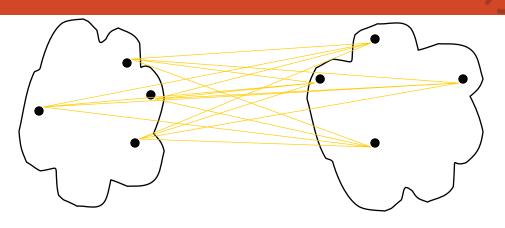


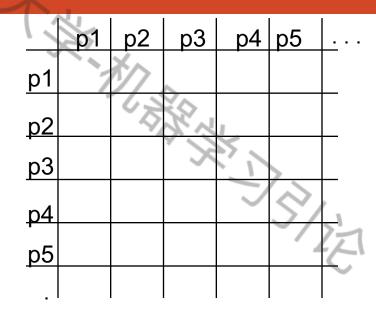
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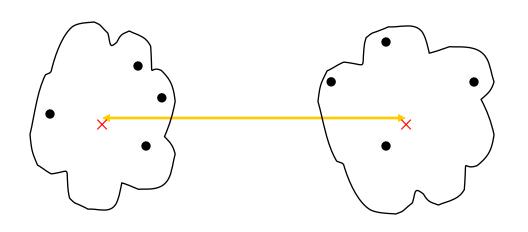


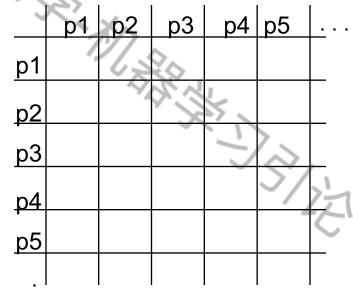
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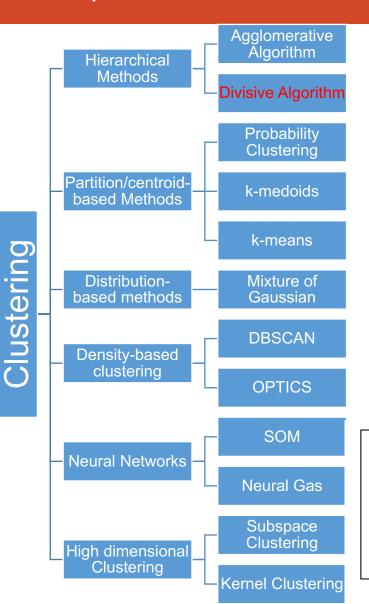


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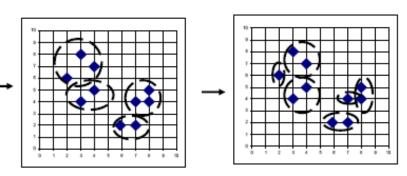


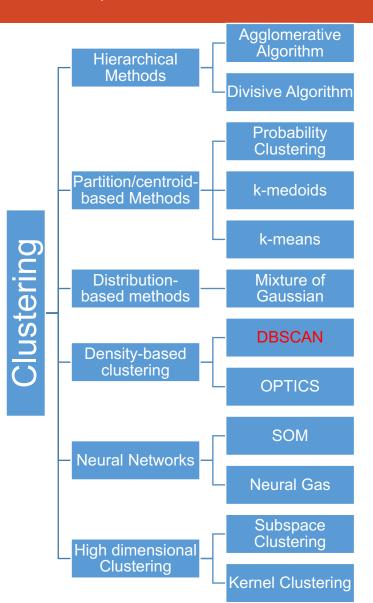


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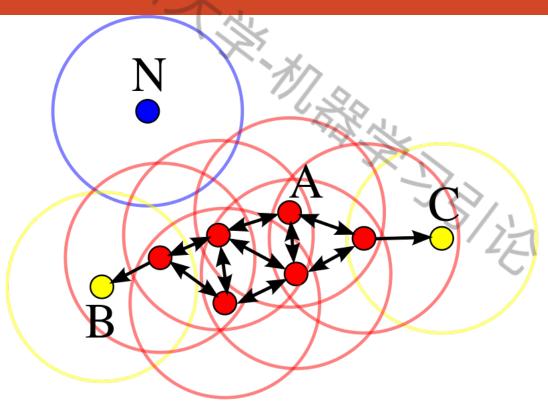


- Divisive (Top-down)
 - Start at the top with all patterns in one cluster
 - one clusterThe cluster is split using a flat clustering algorithm
 - This procedure is applied recursively until each pattern is in its own singleton cluster

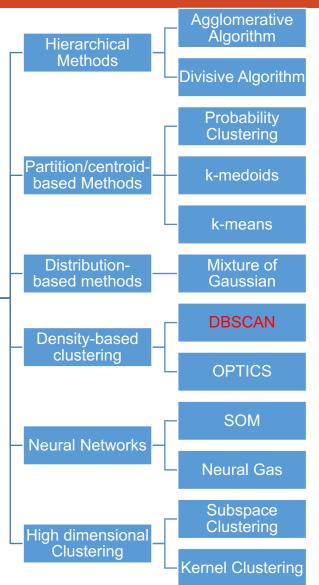




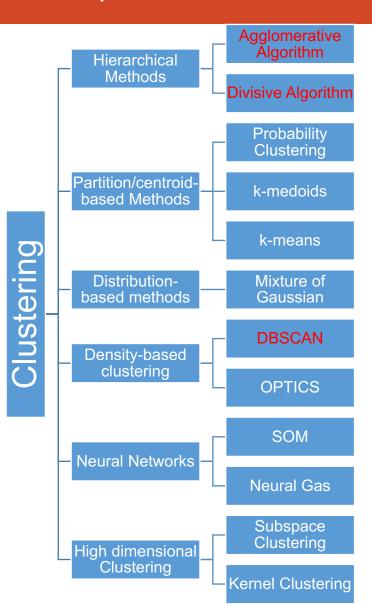
- Important Questions:
 - How do we measure density?
 - What is a dense region?
- Density at point p: number of points within a circle of radius Eps
- Dense Region: A circle of radius Eps that contains at least MinPts points



In this diagram, minPts = 4. Point A and the other red points are core points, because the area surrounding these points in an ε radius contain at least 4 points (including the point itself). Because they are all reachable from one another, they form a single cluster. Points B and C are not core points, but are reachable from A (via other core points) and thus belong to the cluster as well. Point N is a noise point that is neither a core point nor directly-reachable.



- Label points as core, border and noise
- Eliminate noise points
- For every core point p that has not been assigned to a cluster
 - Create a new cluster with the point p and all the points that are density-connected to p.
- Assign border points to the cluster of the closest core point.



No objective function is directly minimized, i.e., there are not learning based methods.

提纲

- . Review
- ☐ . k-means clustering
- 三 . k-medoids clustering
- 四. Mixture of Gaussian

k-means: 学习k个means(均值), where each mean corresponds to a cluster center. In other words, k-means achieves clustering by learning/finding k cluster centers.

- Given a set of data points, group them into multiple clusters so that:
 - **points** within each cluster are similar to each other $\min \sum_{j} \sum_{\mathbf{x}_i \in C_j} \|\mathbf{x}_i \mathbf{u}_j\|_2^2$
 - lacksquare points from different clusters are dissimilar $\max \sum_i \sum_j \|\mathbf{u}_i \mathbf{u}_j\|_2^2$

i.e. the cluster assignment (label) and centers are unknown.

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Solution: Iteratively learning clustering assignment and cluster centers so that

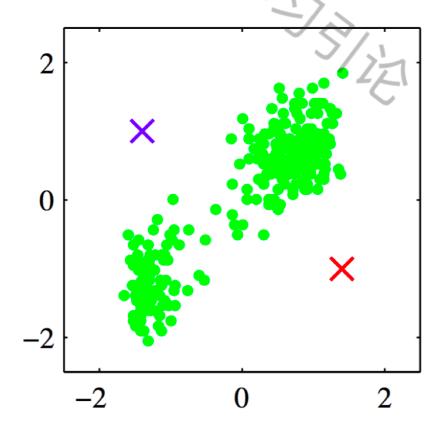
k-means: k个means(均值), clearly, each mean corresponds to a cluster center. In other words, k-means achieves clustering by learning/finding k cluster centers.

Solution: Iteratively learning clustering assignment and cluster centers so that

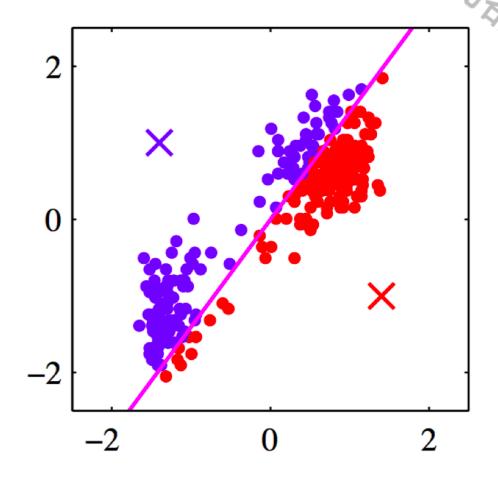
$$\min \sum_{j} \sum_{\mathbf{x}_i \in C_j} \|\mathbf{x}_i - \mathbf{u}_j\|_2^2 \quad \max \sum_{i} \sum_{j} \|\mathbf{u}_i - \mathbf{u}_j\|_2^2$$

Example:

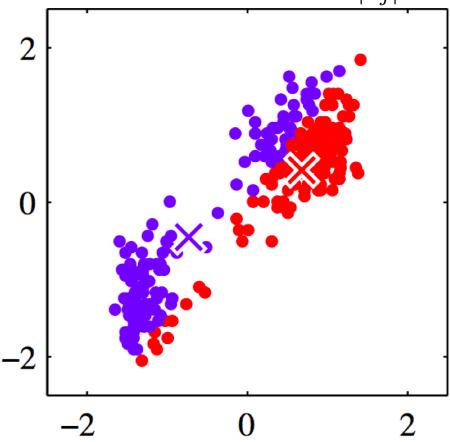
Iter1: randomly choose two points as cluster centers



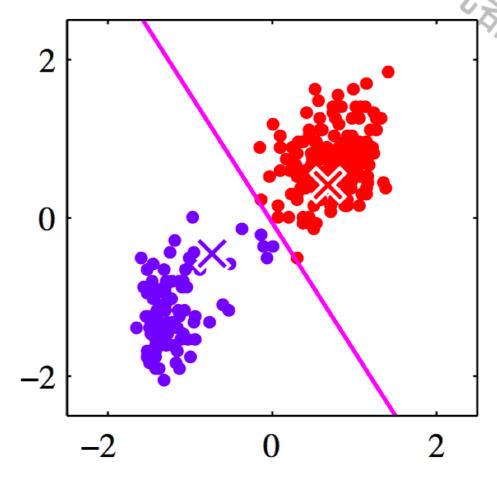
Iter1: Assign each point to closest center.



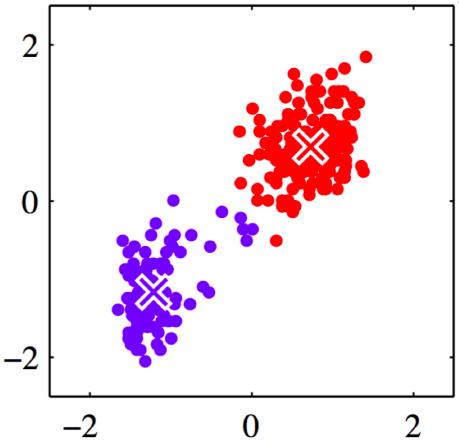
Iter2: Compute new class centers by $\mu_j = \frac{\sum_{\mathbf{x}_i \in \mathcal{C}_j} \mathbf{x}_i}{|\mathcal{C}_i|}$



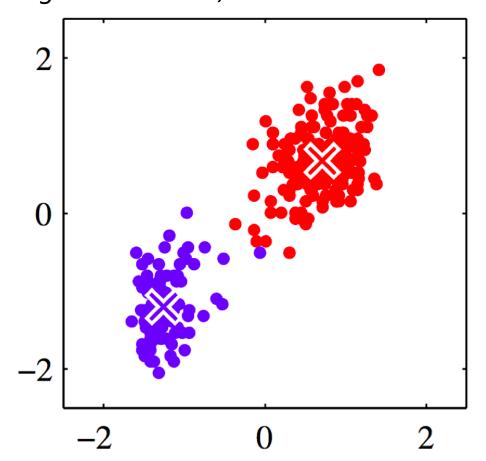
Iter2: Assign points to closest center.



Iter3: Compute cluster centers by $\mu_j = \frac{\sum_{\mathbf{x}_i \in \mathcal{C}_j} \mathbf{x}_i}{|\mathcal{C}_j|}$



Iterate until convergence (reach to the max iteration number or the loss is smaller than a given threshold).



- Dataset $\mathcal{D} = \{x_1, \dots, x_n\} \in \mathbf{R}^d$
- Goal (version 1): Partition data into k clusters.
- Goal (version 2): Partition \mathbb{R}^d into k regions.
- Let μ_1, \ldots, μ_k denote cluster centers.

- Dataset $\mathcal{D} = \{x_1, \dots, x_n\} \in \mathbf{R}^d$
- Goal (version 1): Partition data into k clusters.
- Goal (version 2): Partition \mathbb{R}^d into k regions.
- Let μ_1, \ldots, μ_k denote cluster centers.
- For each x_i , use a **one-hot encoding** to designate membership:

$$r_i = (0, 0, \dots, 0, 0, 1, 0, 0) \in \mathbf{R}^k$$

Let

$$r_{ic} = 1(x_i \text{ assigned to cluster } c).$$

Then

$$r_i = (r_{i1}, r_{i2}, \ldots, r_{ik}).$$

Find cluster centers and cluster assignments minimizing

$$J(r, \mu) = \sum_{i=1}^{n} \sum_{c=1}^{k} r_{ic} ||x_i - \mu_c||^2.$$

Find cluster centers and cluster assignments minimizing

$$J(r, \mu) = \sum_{i=1}^{n} \sum_{c=1}^{k} r_{ic} ||x_i - \mu_c||^2.$$

- Is objective function convex?
- What's the domain of *J*?

Find cluster centers and cluster assignments minimizing

$$J(r, \mu) = \sum_{i=1}^{n} \sum_{c=1}^{k} r_{ic} ||x_i - \mu_c||^2.$$

- Is objective function convex?
- What's the domain of *J*?
- $r \in \{0, 1\}^{n \times k}$, which is not a convex set...
- So domain of J is not convex \Longrightarrow J is not a convex function
- We should expect local minima.

• For fixed r (cluster assignments), minimizing over μ is easy:

$$J(r, \mu) = \sum_{i=1}^{n} \sum_{c=1}^{k} r_{ic} ||x_{i} - \mu_{c}||^{2}$$

$$= \sum_{c=1}^{k} \sum_{j=1}^{n} r_{ic} ||x_{i} - \mu_{c}||^{2}$$

$$= \int_{c} ||x_{i} - \mu_{c}||^{2}$$

$$J_{c}(\mu_{c}) = \sum_{\{i | x_{i} \text{ belongs to cluster } c\}} ||x_{i} - \mu_{c}||^{2}$$

• J_c is minimized by

$$\mu_c = \text{mean}(\{x_i \mid x_i \text{ belongs to cluster } c\})$$

• For fixed μ (cluster centers), minimizing over r is easy:

$$J(r, \mu) = \sum_{i=1}^{n} \sum_{c=1}^{k} r_{ic} ||x_i - \mu_c||^2$$

• For each *i*, exactly one of the following terms is nonzero:

$$r_{i1}||x_i - \mu_1||^2$$
, $r_{i2}||x_i - \mu_2||^2$, ..., $r_{ik}||x_i - \mu_k||^2$

Take

$$r_{ic} = 1(c = \underset{j}{\operatorname{arg\,min}} \|x_i - \mu_j\|^2)$$

• That is, assign x_i to cluster c with minimum distance

$$||x_i - \mu_c||^2$$

- We will use an alternating minimization algorithm:
 - ① Choose initial cluster centers $\mu = (\mu_1, \dots, \mu_k)$.
 - e.g. choose k randomly chosen data points
 - Repeat
 - For given cluster centers, find optimal cluster assignments:

$$r_{ic}^{\text{new}} = 1(c = \underset{i}{\text{arg min}} \|x_i - \mu_j\|^2)$$

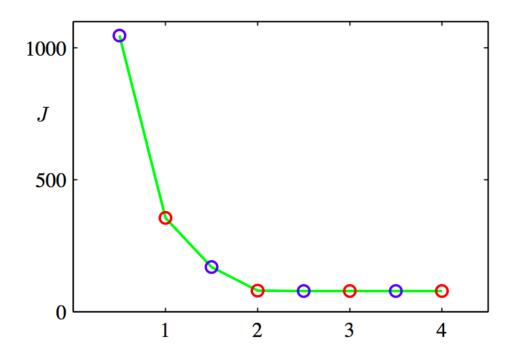
@ Given cluster assignments, find optimal cluster centers:

$$\mu_c^{\mathsf{new}} = \underset{m \in \mathsf{R}^d}{\mathsf{arg\,min}}; \sum_{\{i \mid r_{ic} = 1\}} \|x_i - \mu_c\|^2$$

Convergence:

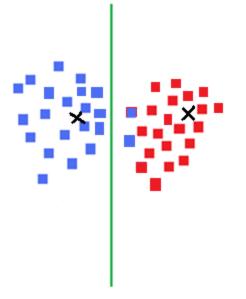
- Note: Objective value never increases in an update.
 - (Obvious: worst case, everything stays the same)
- Consider the sequence of objective values: J_1, J_2, J_3, \dots
 - monotonically decreasing
 - bounded below by zero
- Therefore, k-Means objective value converges to $\inf_t J_t$.
- Reminder: This is convergence to a local minimum.
- Best to repeat k-means several times, with different starting points

- Blue circles after "E" step: assigning each point to a cluster
- Red circles after "M" step: recomputing the cluster centers



- Disadvantages
 - Dependent on initialization
 - Select random seeds with at least D_{\min}
 - Or, run the algorithm many times

- Disadvantages
 - Dependent on initialization
 - Sensitive to outliers



提纲

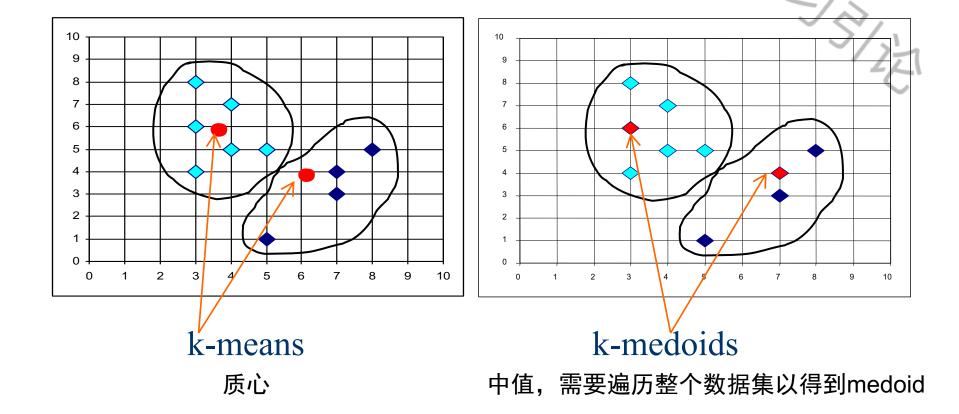
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三、k-medoids clustering

- *k*-means (MacQueen'67): Each cluster is represented by center of cluster
 - Sensitive to noise/outlier
- *k*-medoids (Kaufman & Rousseeuw'87): Each cluster is represented by one of the objects (medoid) in cluster
 - Robust to noise/outlier
 - keep the physical meaning of the dataset
 - Higher computational cost than *k*-means

三、k-medoids clustering

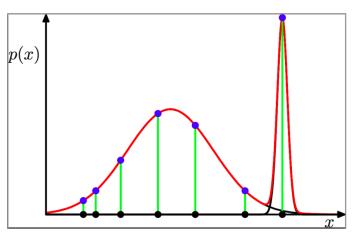
• k-medoids: Find k representative objects, called medoids



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Universal Approximation: any distribution could be represented by a MOG, namely, any data set is a MOG and each cluster corresponds to a Gaussian distribution.



1-dimensional
$$\mathcal{N}(x|\mu,\sigma^2) = \frac{1}{2\pi\sigma^2} \exp(-\frac{1}{2\sigma^2}(x-\mu)^2)$$

Multivariate Gaussian

$$\mathcal{N}(\mathbf{x}|\mu, \mathbf{\Sigma}) = \frac{1}{(2\pi)^{D/2}} \frac{1}{|\mathbf{\Sigma}|^{1/2}} \exp(-\frac{1}{2} (\mathbf{x} - \mu)^T \mathbf{\Sigma}^{-1} (\mathbf{x} - \mu))$$

Definition

A probability density p(x) represents a **mixture distribution** or **mixture model**, if we can write it as a **convex combination** of probability densities. That is,

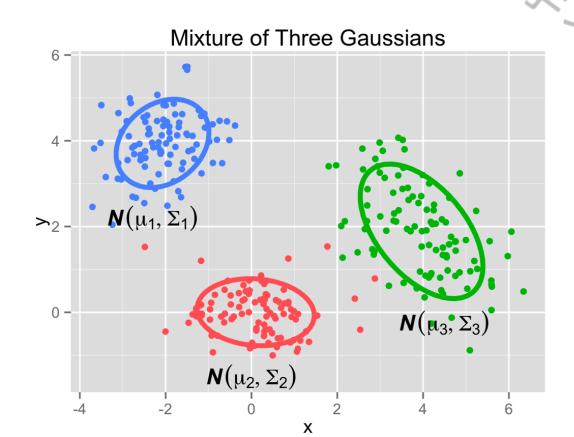
$$p(x) = \sum_{i=1}^{k} w_i p_i(x),$$

where $w_i \ge 0$, $\sum_{i=1}^k w_i = 1$, and each p_i is a probability density.

- In our Gaussian mixture model, X has a mixture distribution.
- More constructively, let S be a set of probability distributions:
 - \bullet Choose a distribution randomly from S.
 - Sample X from the chosen distribution.
- Then X has a mixture distribution.

- Let's consider a generative model for the data.
- Suppose
 - \bigcirc There are k clusters.
 - We have a probability density for each cluster.
- Generate a point as follows
 - ① Choose a random cluster $z \in \{1, 2, ..., k\}$.
 - $Z \sim \mathsf{Multi}(\pi_1, \ldots, \pi_k)$.
 - \bigcirc Choose a point from the distribution for cluster Z.
 - $X \mid Z = z \sim p(x \mid z).$

- **1** Choose $Z \in \{1, 2, 3\} \sim \text{Multi}(\frac{1}{3}, \frac{1}{3}, \frac{1}{3})$.
- ② Choose $X \mid Z = z \sim \mathcal{N}(X \mid \mu_z, \Sigma_z)$.



Cluster probabilities:

 $\pi = (\pi_1, \ldots, \pi_k)$

Cluster means:

 $\mu = (\mu_1, \ldots, \mu_k)$

Cluster covariance matrices:

 $\Sigma = (\Sigma_1, \dots \Sigma_k)$

Since we only observe X, we

四、Mixture of Gaussian

• The model likelihood for $\mathcal{D} = \{x_1, \dots, x_n\}$ is

$$L(\pi, \mu, \Sigma) = \prod_{i=1}^{n} p(x_i)$$

$$= \sum_{i=1}^{k} p(x_i)$$

$$= \sum_{j=1}^{k} \pi_z \mathcal{N}(x_j | \mu_z, \Sigma_z)$$

$$= \sum_{j=1}^{k} \pi_z \mathcal{N}(x_j | \mu_z, \Sigma_z).$$

• As usual, we'll take our objective function to be the log of this:

$$J(\pi, \mu, \Sigma) = \sum_{i=1}^{n} \log \left\{ \sum_{z=1}^{k} \pi_{z} \mathcal{N}(x_{i} \mid \mu_{z}, \Sigma_{z}) \right\}$$

$$\mathcal{N}(\mathbf{x}|\mu, \mathbf{\Sigma}) = \frac{1}{(2\pi)^{D/2}} \frac{1}{|\mathbf{\Sigma}|^{1/2}} \exp(-\frac{1}{2} (\mathbf{x} - \mu)^T \mathbf{\Sigma}^{-1} (\mathbf{x} - \mu))$$

- Let's start by considering the MLE for the Gaussian model.
- For data $\mathcal{D} = \{x_1, \dots, x_n\}$, the log likelihood is given by

$$\sum_{i=1}^{n} \log \mathcal{N}(x_i \mid \mu, \Sigma) = -\frac{nd}{2} \log (2\pi) - \frac{n}{2} \log |\Sigma| - \frac{1}{2} \sum_{i=1}^{n} (x_i - \mu)' \Sigma^{-1}(x_i - \mu).$$

With some calculus, we find that the MLE parameters are

$$\mu_{\text{MLE}} = \frac{1}{n} \sum_{i=1}^{n} x_{i}$$

$$\Sigma_{\text{MLE}} = \frac{1}{n} \sum_{i=1}^{n} (x_{i} - \mu_{\text{MLE}}) (x_{i} - \mu_{\text{MLE}})^{T}$$

- For GMM, If we knew the cluster assignment z_i for each x_i ,
 - we could compute the MLEs for each cluster.

• Denote the probability that observed value x_i comes from cluster j by

$$\gamma_i^j = \mathbb{P}(Z = j \mid X = x_i).$$

- The **responsibility** that cluster j takes for observation x_i .
- Computationally,

$$\gamma_{i}^{j} = \mathbb{P}(Z = j \mid X = x_{i}).$$

$$= p(Z = j, X = x_{i})/p(x)$$

$$= \frac{\pi_{j} \mathcal{N}(x_{i} \mid \mu_{j}, \Sigma_{j})}{\sum_{c=1}^{k} \pi_{c} \mathcal{N}(x_{i} \mid \mu_{c}, \Sigma_{c})}$$

- The vector $(\gamma_i^1, \dots, \gamma_i^k)$ is exactly the **soft assignment** for x_i .
- Let $n_c = \sum_{i=1}^n \gamma_i^c$ be the number of points "soft assigned" to cluster c.

- **1** Initialize parameters μ , Σ , π .
- 2 "E step". Evaluate the responsibilities using current parameters:

$$\gamma_i^j = \frac{\pi_j \mathcal{N}(x_i \mid \mu_j, \Sigma_j)}{\sum_{c=1}^k \pi_c \mathcal{N}(x_i \mid \mu_c, \Sigma_c)},$$

for i = 1, ..., n and j = 1, ..., k.

(3) "M step". Re-estimate the parameters using responsibilities:

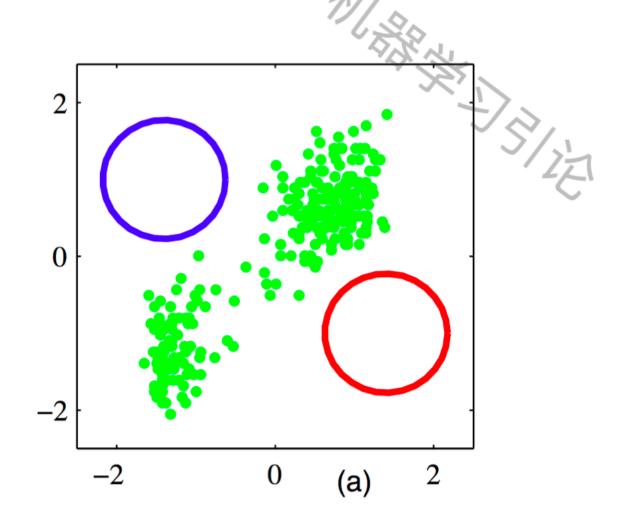
$$\mu_c^{\text{new}} = \frac{1}{n_c} \sum_{i=1}^n \gamma_i^c x_i$$

$$\Sigma_c^{\text{new}} = \frac{1}{n_c} \sum_{i=1}^n \gamma_i^c (x_i - \mu_{\text{MLE}}) (x_i - \mu_{\text{MLE}})^T$$

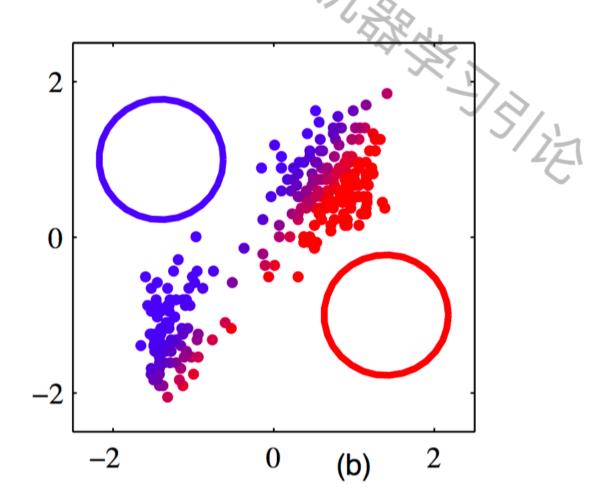
$$\pi_c^{\text{new}} = \frac{n_c}{n},$$

Repeat from Step 2, until log-likelihood converges.

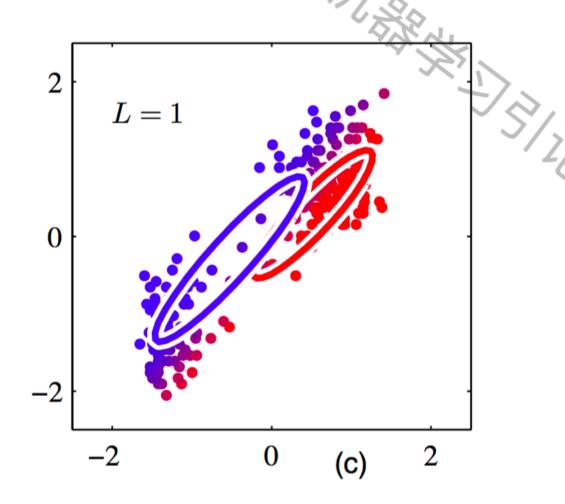
Initialization



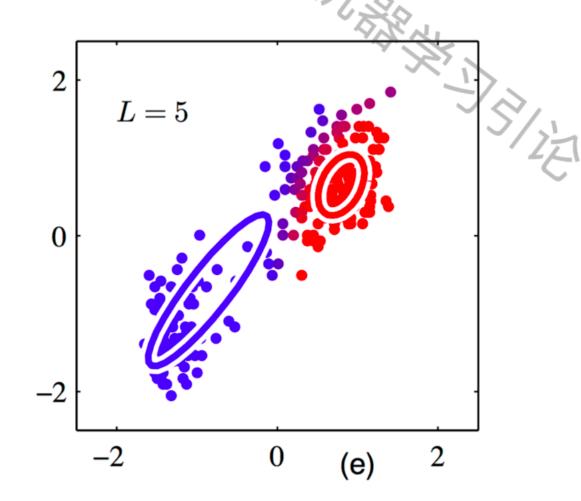
• First soft assignment:



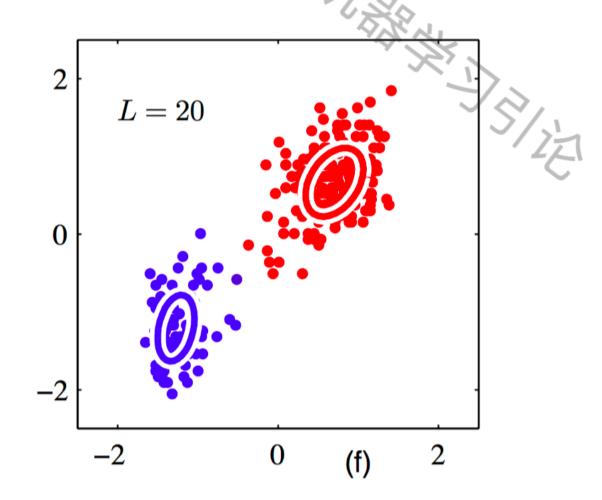
• First soft assignment:



• After 5 rounds of EM:



• After 20 rounds of EM:



k-means vs. MOG

- EM for GMM seems a little like k-means.
- In fact, there is a precise correspondence.
- First, fix each cluster covariance matrix to be $\sigma^2 I$.
- As we take $\sigma^2 \to 0$, the update equations converge to doing k-means.
- If you do a quick experiment yourself, you'll find
 - Soft assignments converge to hard assignments.
 - Has to do with the tail behavior (exponential decay) of Gaussian.

Test Questions:

- Write k-means in pseudocode?
- Write MOG in pseudocode?
- What the limitations of k-means besides that I given?
- What the limitations of MOG besides that I given?
- What the connections between k-means and MOG?
- What the difference between k-means and MOG?
- What the advantage of k-means/MOG over MOG/k-means?

Further reading:

[1] D. P. Kingma and M. Welling, "Auto-Encoding Variational Bayes," presented at the International Conference on Learning Representations, 2014.

Final Test

For a given data set (mnist test partition), achieving

- 1. a classification accuracy over 80% using the methods introduced in this course. Report the corresponding F-measure.
- 2. alternatively, a clustering accuracy over 58% using the methods introduced in this course. Report the corresponding NMI.

Requirements:

- Give the design details and explain why it as does
- Report the mean and std score
- Report the tuned parameters
- Report the hardware and used time cost

Q&A THANKS!