CSCI567 Machine Learning (Fall 2021)

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U of Southern California

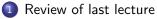
Oct 28, 2021

HW3: discuss solutions today

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HW4: to be released, due on Tue, 11/09





Clustering 2



Gaussian mixture models

Outline



Review of last lecture

Clustering



General training algorithm for decision trees

DecisionTreeLearning(Examples, Features)

- if Examples have the same class, return a leaf with this class
- else if Features is empty, return a leaf with the majority class
- else if Examples is empty, return a leaf with majority class of parent

else

find the best feature A to split (e.g. based on conditional entropy)

Tree \leftarrow a root with test on A

For each value a of A:

Child \leftarrow **DecisionTreeLearning**(Examples with A = a, Features \{A}) add **Child** to **Tree** as a new branch

• return Tree

The AdaBoost Algorithm

Given a training set S and a base algorithm \mathcal{A} , initialize D_1 to be uniform

For $t = 1, \ldots, T$

- obtain a weak classifier $h_t \leftarrow \mathcal{A}(S, D_t)$
- calculate the importance of h_t as

$$\beta_t = \frac{1}{2} \ln \left(\frac{1 - \epsilon_t}{\epsilon_t} \right) \qquad \qquad (\beta_t > 0 \Leftrightarrow \epsilon_t < 0.5)$$

where $\epsilon_t = \sum_{n:h_t(\boldsymbol{x}_n) \neq y_n} D_t(n)$ is the weighted error of h_t .

• update distributions

$$D_{t+1}(n) \propto D_t(n) e^{-\beta_t y_n h_t(\boldsymbol{x}_n)} = \begin{cases} D_t(n) e^{-\beta_t} & \text{if } h_t(x_n) = y_n \\ D_t(n) e^{\beta_t} & \text{else} \end{cases}$$

Output the final classifier $H(\boldsymbol{x}) = \operatorname{sgn}\left(\sum_{t=1}^{T} \beta_t h_t(\boldsymbol{x})\right)$

Outline



2 Clustering

- Problem setup
- K-means algorithm
- Initialization and Convergence



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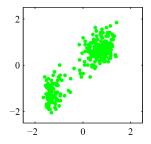
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Today's focus: **clustering**, an important unsupervised learning problem

Clustering: informal definition

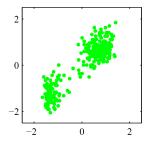
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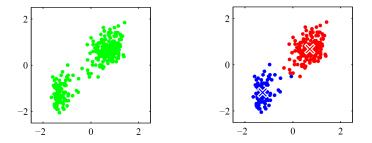


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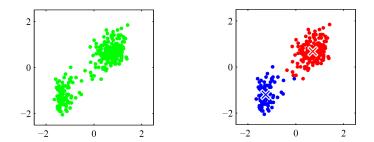
Output: group the data into some clusters, which means

- assign each point to a specific cluster
- find the center (representative/prototype/...) of each cluster



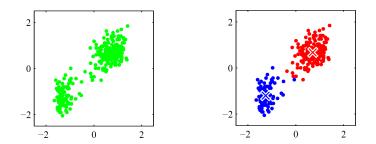
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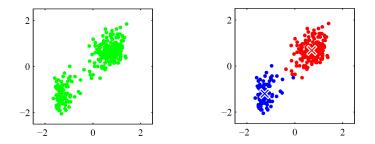


Clustering: formal definition

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Output: group the data into K clusters, which means

• find assignment $\gamma_{nk} \in \{0,1\}$ for each data point $n \in [N]$ and $k \in [K]$ s.t. $\sum_{k \in [K]} \gamma_{nk} = 1$ for any fixed n



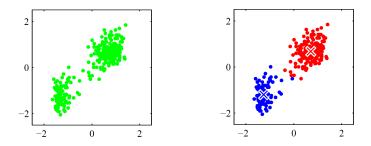
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- find the cluster centers $\mu_1, \ldots, \mu_K \in \mathbb{R}^{\mathsf{D}}$



Many applications

- recognize communities in a social network
- group similar customers in market research
- image segmentation
- accelerate other algorithms (e.g. NNC as in programing projects)

• . . .

One example

image compression:

- each pixel is a point
- perform clustering over these points
- replace each point by the center of the cluster it belongs to









Original image

 $\mathsf{Large}\ K \longrightarrow \mathsf{Small}\ K$

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Still, we can turn it into an optimization problem, e.g. through the popular "K-means" objective: find γ_{nk} and μ_k to minimize

$$F(\{\gamma_{nk}\}, \{\mu_k\}) = \sum_{n=1}^{N} \sum_{k=1}^{K} \gamma_{nk} \| \boldsymbol{x}_n - \boldsymbol{\mu}_k \|_2^2$$

i.e. the sum of squared distances of each point to its center.

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A closer look

The first step

$$\min_{\{\gamma_{nk}\}} F\left(\{\gamma_{nk}\}, \{\boldsymbol{\mu}_k\}\right) = \min_{\{\gamma_{nk}\}} \sum_n \sum_k \gamma_{nk} \|\boldsymbol{x}_n - \boldsymbol{\mu}_k\|_2^2$$

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is simply to assign each x_n to the closest μ_k , i.e.

$$\gamma_{nk} = \mathbb{I}\left[k = \operatorname*{argmin}_{c} \|\boldsymbol{x}_{n} - \boldsymbol{\mu}_{c}\|_{2}^{2}\right]$$

for all $k \in [K]$ and $n \in [N]$.

A closer look

The second step

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is simply to average the points of each cluster (hence the name)

$$\boldsymbol{\mu}_k = \frac{\sum_{n:\gamma_{nk}=1} \boldsymbol{x}_n}{|\{n:\gamma_{nk}=1\}|} = \frac{\sum_n \gamma_{nk} \boldsymbol{x}_n}{\sum_n \gamma_{nk}}$$

for each $k \in [K]$.

The K-means algorithm

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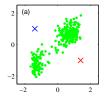
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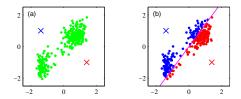
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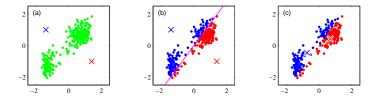
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Step 3 Return to Step 1 if not converged

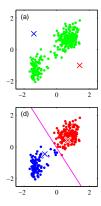
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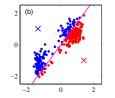


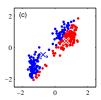




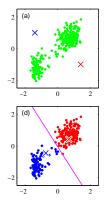
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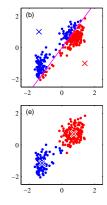


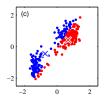




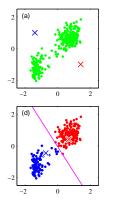
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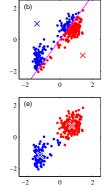


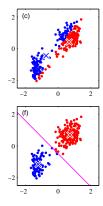




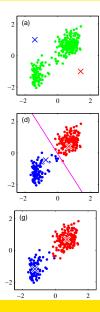
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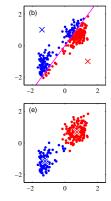


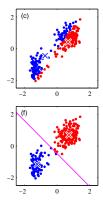




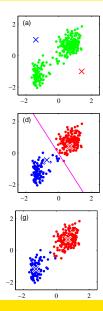
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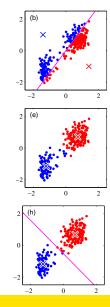


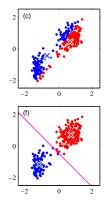




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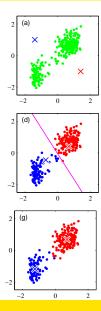


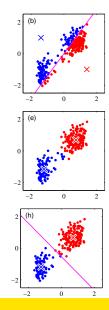


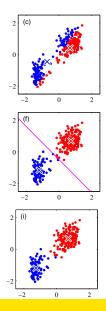


K-means algorithm

An example







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Initialization matters for convergence.

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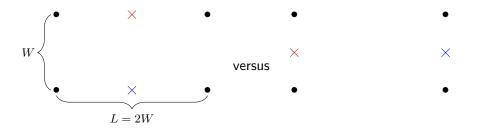
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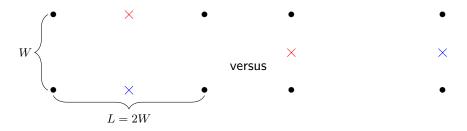
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- it could take exponentially many iterations to converge
- and it *might not converge to the global minimum* of the K-means objective

Simple example: 4 data points, 2 clusters, 2 different initializations

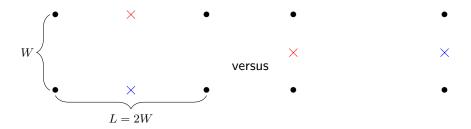


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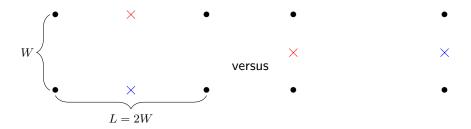
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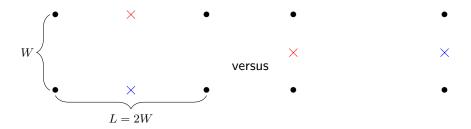
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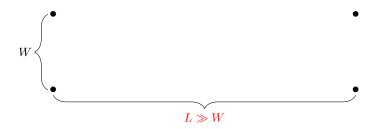
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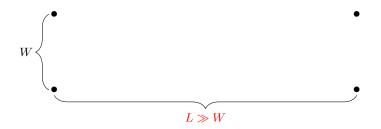
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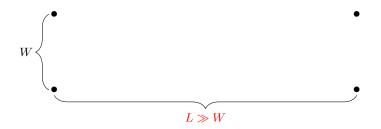
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- left has K-means objective $L^2 = 4W^2$
- right has K-means objective W^2 , 4 times better than left!
- in fact, left is local minimum, and right is global minimum.



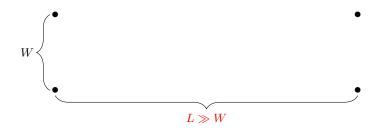


• moreover, local minimum can be *arbitrarily worse* if we increase L



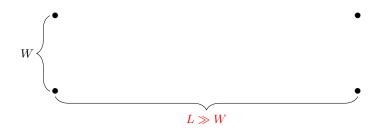
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• so *initialization matters a lot* for K-means



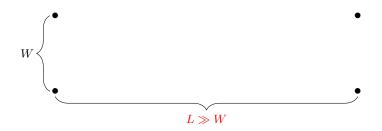
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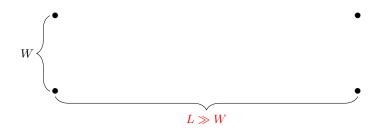


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- randomly pick K points as initial centers: fails with 1/3 probability
- or randomly assign each point to a cluster, then average: similarly fail with a constant probability
- or more sophisticated approaches: K-means++ guarantees to find a solution that in expectation is at most O(log K) times of the optimal

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For $k = 2, \ldots, K$

• randomly pick the k-th center μ_k such that

$$\Pr[oldsymbol{\mu}_k = oldsymbol{x}_n] \propto \min_{j=1,...,k-1} \|oldsymbol{x}_n - oldsymbol{\mu}_j\|_2^2$$

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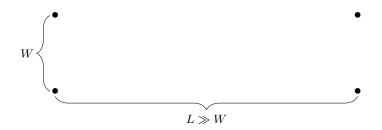
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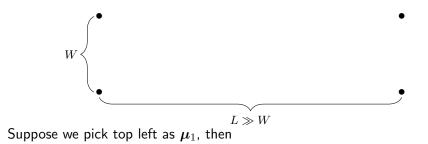
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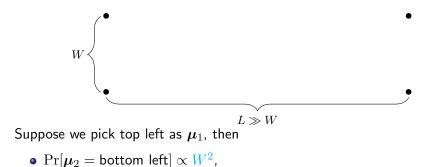
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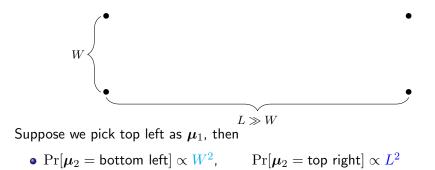
$$\Pr[oldsymbol{\mu}_k = oldsymbol{x}_n] \propto \min_{j=1,...,k-1} \|oldsymbol{x}_n - oldsymbol{\mu}_j\|_2^2$$

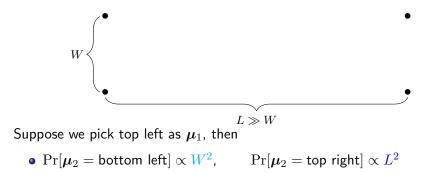
Intuitively this spreads out the initial centers.



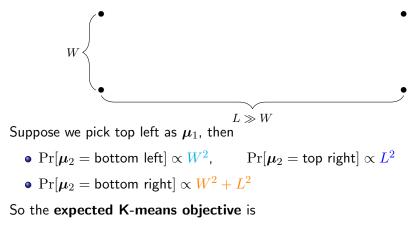




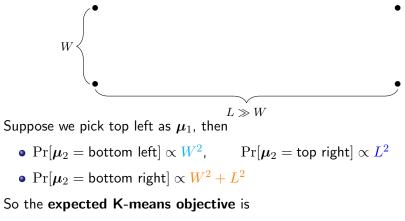




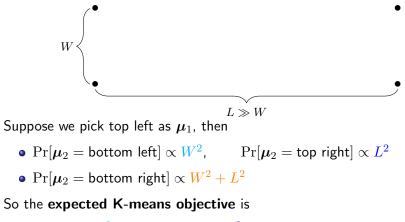
•
$$\Pr[\mu_2 = \text{bottom right}] \propto W^2 + L^2$$



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that is, at most 1.5 times of the optimal.

Summary for K-means

K-means is alternating minimization for the K-means objective.

The initialization matters a lot for the convergence.

K-means++ uses a theoretically (and often empirically) better initialization.

Outline



Clustering

- Gaussian mixture models
 - Motivation and Model
 - EM algorithm
 - EM applied to GMMs

Gaussian mixture models

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To solve GMM, we will introduce a powerful method for learning probabilistic model: **Expectation–Maximization (EM) algorithm**

For classification, we discussed the sigmoid model to "explain" how the labels are generated.

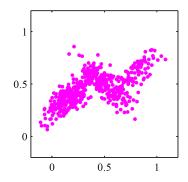
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That is, each point is an independent sample of $\boldsymbol{x} \sim p$.

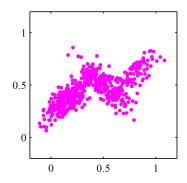


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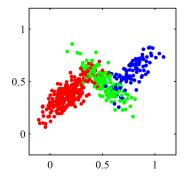
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What probabilistic model generates data like this?



GMM is a natural model to explain such data

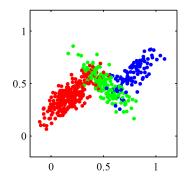
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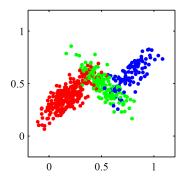
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Hence the name "Gaussian mixture model".

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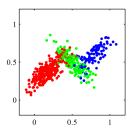
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 \boldsymbol{x} and z are both random variables drawn from the model

• x is observed

• z is unobserved/latent

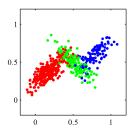
An example



The conditional distributions are

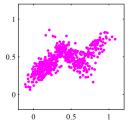
$$\begin{split} p(\boldsymbol{x} \mid z = \mathsf{red}) &= N(\boldsymbol{x} \mid \boldsymbol{\mu}_1, \boldsymbol{\Sigma}_1) \\ p(\boldsymbol{x} \mid z = \mathsf{blue}) &= N(\boldsymbol{x} \mid \boldsymbol{\mu}_2, \boldsymbol{\Sigma}_2) \\ p(\boldsymbol{x} \mid z = \mathsf{green}) &= N(\boldsymbol{x} \mid \boldsymbol{\mu}_3, \boldsymbol{\Sigma}_3) \end{split}$$

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The marginal distribution is

$$\begin{split} p(\pmb{x}) &= p(\mathsf{red}) N(\pmb{x} \mid \pmb{\mu}_1, \pmb{\Sigma}_1) + p(\mathsf{blue}) N(\pmb{x} \mid \pmb{\mu}_2, \pmb{\Sigma}_2) \\ &+ p(\mathsf{green}) N(\pmb{x} \mid \pmb{\mu}_3, \pmb{\Sigma}_3) \end{split}$$

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- both learn the cluster centers μ_k 's
- ullet in addition, GMM learns cluster weight ω_k and covariance $oldsymbol{\Sigma}_k$, thus
 - we can predict probability of seeing a new point
 - we can generate synthetic data

How to learn these parameters?

An obvious attempt is maximum-likelihood estimation (MLE): find

$$\underset{\boldsymbol{\theta}}{\operatorname{argmax}} \ln \prod_{n=1}^{N} p(\boldsymbol{x}_{n} ; \boldsymbol{\theta}) = \underset{\boldsymbol{\theta}}{\operatorname{argmax}} \sum_{n=1}^{N} \ln p(\boldsymbol{x}_{n} ; \boldsymbol{\theta}) \triangleq \underset{\boldsymbol{\theta}}{\operatorname{argmax}} P(\boldsymbol{\theta})$$

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One solution is to still apply GD/SGD, but a much more effective approach is the **Expectation–Maximization (EM) algorithm**.

Step 0 Initialize $\omega_k, \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k$ for each $k \in [K]$

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$$\omega_k = rac{\sum_n \gamma_{nk}}{N}$$
 $\mu_k = rac{\sum_n \gamma_{nk} \boldsymbol{x}_n}{\sum_n \gamma_{nk}}$

$$\boldsymbol{\Sigma}_k = rac{1}{\sum_n \gamma_{nk}} \sum_n \gamma_{nk} (\boldsymbol{x}_n - \boldsymbol{\mu}_k) (\boldsymbol{x}_n - \boldsymbol{\mu}_k)^{\mathrm{T}}$$

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We will see how this is a special case of EM.

Generate 50 data points from a mixture of 2 Gaussians with

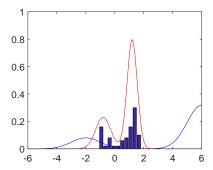
- $\omega_1 = 0.3, \mu_1 = -0.8, \Sigma_1 = 0.52$
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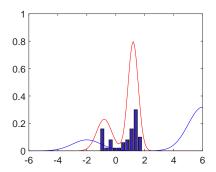
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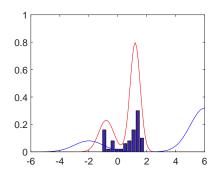
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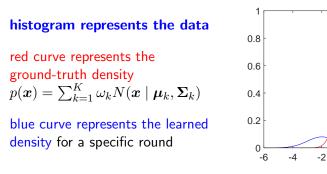
blue curve represents the learned density for a specific round

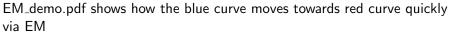


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6

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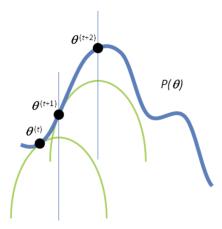
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Again, directly solving the objective is intractable.

High level idea

Keep maximizing a lower bound of P that is more manageable



Derivation of EM

Finding the lower bound of *P*:

$$\ln p(\boldsymbol{x} ; \boldsymbol{\theta}) = \ln \frac{p(\boldsymbol{x}, z ; \boldsymbol{\theta})}{p(z | \boldsymbol{x} ; \boldsymbol{\theta})}$$

(true for any z)

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(Jensen's inequality)

$$= \mathbb{E}_{z \sim q} \left[\ln p(\boldsymbol{x}, z ; \boldsymbol{\theta}) \right] + H(q)$$

Therefore, we obtain a lower bound for the log-likelihood function

$$P(\boldsymbol{\theta}) = \sum_{n=1}^{N} \ln p(\boldsymbol{x}_n ; \boldsymbol{\theta})$$

$$\geq \sum_{n=1}^{N} \left(\mathbb{E}_{z_n \sim q_n} \left[\ln p(\boldsymbol{x}_n, z_n ; \boldsymbol{\theta}) \right] + H(q_n) \right) = F(\boldsymbol{\theta}, \{q_n\})$$

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This holds for any $\{q_n\}$, so how do we choose? Naturally, the one that maximizes the lower bound (i.e. the tightest lower bound)!

Equivalently, this is the same as alternatingly maximizing F over $\{q_n\}$ and θ (similar to K-means).

Maximizing over $\{q_n\}$

Fix $\boldsymbol{\theta}^{(t)}$, the solution to

$$\operatorname*{argmax}_{q_n} \mathbb{E}_{z_n \sim q_n} \left[\ln p(\boldsymbol{x}_n, z_n ; \boldsymbol{\theta}^{(t)}) \right] + H(q_n)$$

is $q_n^{(t)}$ s.t. $q_n^{(t)}(z_n) = p(z_n \mid \pmb{x}_n \ ; \pmb{\theta}^{(t)})$

i.e., the *posterior distribution of* z_n given x_n and $\theta^{(t)}$. (Verified in HW4)

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So at $\boldsymbol{\theta}^{(t)}$, we found the tightest lower bound $F\left(\boldsymbol{\theta}, \{q_n^{(t)}\}\right)$:

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$$F\left(\boldsymbol{\theta}, \{q_n^{(t)}\}\right) \leq P(\boldsymbol{\theta})$$
 for all $\boldsymbol{\theta}$.
• $F\left(\boldsymbol{\theta}^{(t)}, \{q_n^{(t)}\}\right) = P(\boldsymbol{\theta}^{(t)})$ (verify yourself by going through Slide 40)

Fix $\{q_n^{(t)}\}$, maximize over $\boldsymbol{\theta}$:

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Q is the (expected) complete likelihood and is usually more tractable.

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$$P(\theta) = \sum_{n=1}^{N} \ln p(\boldsymbol{x}_n; \theta)$$

EM algorithm

General EM algorithm

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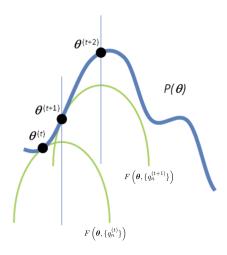
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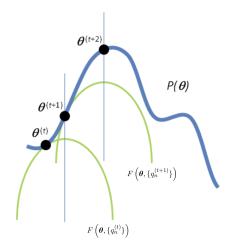
Step 2 (M-Step) update the model parameter via Maximization

$$\boldsymbol{\theta}^{(t+1)} \leftarrow \operatorname*{argmax}_{\boldsymbol{\theta}} Q(\boldsymbol{\theta} ; \boldsymbol{\theta}^{(t)})$$

Step 3 $t \leftarrow t + 1$ and return to Step 1 if not converged

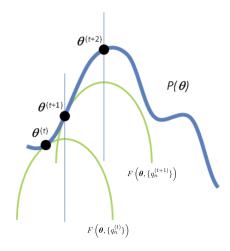


 $P(\pmb{\theta})$ is non-concave, but $Q(\pmb{\theta}; \pmb{\theta}^{(t)})$ often is concave and easy to maximize.



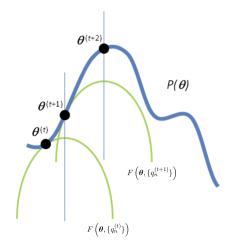
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$$P(\boldsymbol{\theta}^{(\mathsf{t+1})}) \ge F\left(\boldsymbol{\theta}^{(\mathsf{t+1})}; \{q_n^{(t)}\}\right)$$



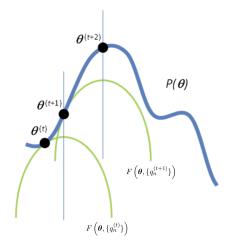
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So EM always increases the objective value and will converge to some local maximum (similar to K-means).

E-Step:

$$q_n^{(t)}(z_n = k) = p\left(z_n = k \mid \boldsymbol{x}_n ; \boldsymbol{\theta}^{(t)}\right)$$
$$\propto p\left(\boldsymbol{x}_n, z_n = k ; \boldsymbol{\theta}^{(t)}\right)$$

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This computes the "soft assignment" $\gamma_{nk} = q_n^{(t)}(z_n = k)$, i.e. conditional probability of x_n belonging to cluster k.

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$$\operatorname{argmax}_{\boldsymbol{\theta}} Q(\boldsymbol{\theta}, \boldsymbol{\theta}^{(t)}) = \operatorname{argmax}_{\boldsymbol{\theta}} \sum_{n=1}^{N} \mathbb{E}_{z_n \sim q_n^{(t)}} \left[\ln p(\boldsymbol{x}_n, z_n ; \boldsymbol{\theta}) \right]$$

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To find each μ_k, Σ_k , solve

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$$\operatorname{argmax}_{\boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k} \sum_{n=1}^N \gamma_{nk} \ln N(\boldsymbol{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$$

Solutions to previous two problems are very natural, for each k

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i.e. (weighted) fraction of examples belonging to cluster k

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You will verify some of these in HW4.

EM for learning GMMs:

Step 0 Initialize $\omega_k, \mu_k, \Sigma_k$ for each $k \in [K]$

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GMM is a soft version of K-means and it provides a probabilistic interpretation of the data, which means we can predict and generate data after learning.