

ISyE 6416: Computational Statistics
Spring 2017

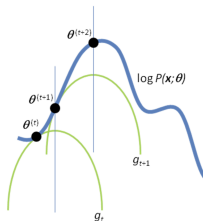
**Lecture 7: EM algorithm and
Gaussian Mixture Model**

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Expectation-Maximization (EM) Algorithm

- ▶ an algorithm to a maximum likelihood estimator in **non-ideal** case: missing data, indirect observations
 - ▶ missing data
 - ▶ clustering (unknown label)
 - ▶ hidden-states in HMM
 - ▶ latent factors
- ▶ replace one difficult likelihood maximization with a sequence of easier maximizations
- ▶ in the limit, the answer to the original problem



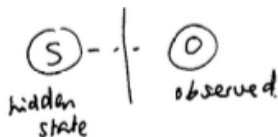
Supplementary Figure 1 Convergence of the EM algorithm. Starting from initial parameters $\theta^{(0)}$, the E-step of the EM algorithm constructs a function $\hat{g}_t$ that lower-bounds the objective function $\log P(x; \theta)$. In the M-step, $\theta^{(t+1)}$ is computed as the maximum of $\hat{g}_t$. In the next E-step, a new lower-bound $\hat{g}_{t+1}$ is constructed; maximization of $\hat{g}_{t+1}$ in the next M-step gives $\theta^{(t+2)}$, etc.

Applications of EM

- ▶ Data clustering in machine learning
- ▶ Natural language processing (Baum-Welch algorithm to fit hidden Markov model)
- ▶ Imputing missing data



General set-up



- ▶ need the distribution of S (or sometimes jointly S and O), but we only observe directly through O

$$S \sim f(S|\theta)$$

parameter θ

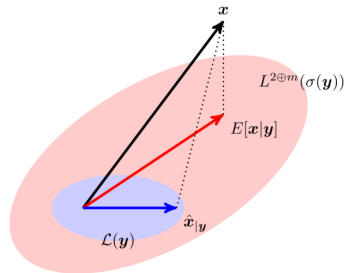
- ▶ Suppose mapping $\chi(o)$ finds all values of S that leads to O
- ▶ distribution of O : $g(O|\theta) = \int_{\chi(o)} f(S|\theta) ds$
- ▶ EM: $\max_{\theta} g(O|\theta)$

Deriving EM

- ▶ Introduce

$$Q(\theta; \theta') = \mathbb{E}\{\log f(S|\theta) | \theta', O\}$$

- ▶ The expectation uses the conditional distribution of S given O and assumed value of parameter θ'



Rhymer's Notes

Intuition

Given O , the “best guess” we could have for S , is its conditional expectation with respect to $S|O, \theta$ (notion of projection); but the computation of expectation involves parameter values. We take a guess, and improve in next round.

E-M algorithm

- ▶ **E-step:** compute expectation of the log-likelihood (observed) incomplete data $\mathbf{y}$, missing data $\mathbf{x}$, complete data $(\mathbf{y}, \mathbf{x})$

$$Q(\theta; \theta') = \mathbb{E}\{\log L(\theta|\mathbf{y}, \mathbf{x})|\theta', \mathbf{y}\}$$

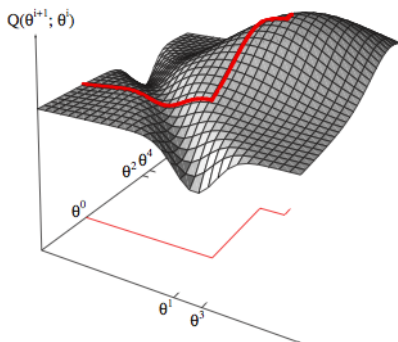
- ▶ **M-step:** compute maximum likelihood using the expectation in previous step

E-step $\Rightarrow$ M-step $\Rightarrow$ E-step $\Rightarrow$ M-step $\Rightarrow$

- ▶ stop until $\|\theta_{k+1} - \theta_k\| < \epsilon$ or $|Q(\theta_{k+1}|\theta_k) - Q(\theta_k|\theta_{k-1})| < \epsilon$

Algorithm 1 (Expectation-Maximization)

```
1 begin initialize  $\theta^0, T, i = 0$   
2   do  $i \leftarrow i + 1$   
3     E step : compute  $Q(\theta; \theta^i)$   
4     M step :  $\theta^{i+1} \leftarrow \arg \max_{\theta} Q(\theta; \theta^i)$   
5     until  $Q(\theta^{i+1}; \theta^i) - Q(\theta^i; \theta^{i-1}) \leq T$   
6   return  $\hat{\theta} \leftarrow \theta^{i+1}$   
7 end
```



Example: EM for missing data

$$n = 4, p = 2$$

$$x_1 = (0, 2)^T, x_2 = (1, 0)^T, x_3 = (2, 2)^T, x_4 = (*, 4)^T$$

Assume they are i.i.d. samples from Gaussian

$$\mathcal{N}([\mu_1, \mu_2]^T, \begin{bmatrix} \sigma_1^2 & 0 \\ 0 & \sigma_2^2 \end{bmatrix})$$

Use EM algorithm to impute the missing data *.

(Cont.) Example: missing data

- Initialization: $\theta_0 = (0, 0, 1, 1)^T$, i.e., mean $[0, 0]^T$ and covariance I_2 .
- E-step

$$\begin{aligned} Q(\theta|\theta_0) &= \mathbb{E}_{x_{41}} [\log p(x|\theta_0, x_1, x_2, x_3, x_{41}, x_{42}) | x_1, x_2, x_3, x_{42}] \\ &= \sum_{i=1}^3 \log p(x_i|\theta) \\ &\quad + \int \log(p([x_{41}, 4]^T|\theta) \cdot p([x_{41}, 4])|\theta_0) dx_{41} \\ &= \sum_{i=1}^3 \log p(x_i|\theta) - \frac{(1 + \mu_1^2)}{2\sigma_1^2} - \frac{(4 - \mu_2)^2}{2\sigma_2^2} - \log(2\pi\sigma_1\sigma_2) \end{aligned}$$

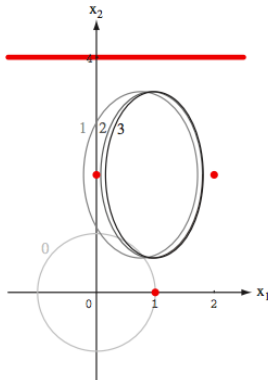
- M-step

$$\theta_1 = \arg \max_{\theta} Q(\theta|\theta_0)$$

(Cont.) Example: missing data - iterations

$$\theta_1 = \begin{bmatrix} 0.75 \\ 2.0 \\ 0.938 \\ 2.0 \end{bmatrix} \Rightarrow \mu_1 = \begin{bmatrix} 0.75 \\ 2.0 \end{bmatrix} \quad \Sigma_1 = \begin{bmatrix} 0.938 & 0 \\ 0 & 2.0 \end{bmatrix}$$

$$\theta_2 = \begin{bmatrix} 1.0 \\ 2.0 \\ 0.667 \\ 2.0 \end{bmatrix}$$



The absent-minded biologist

197 animals

Distributed into 4 categories



125

18

20

34

Multinomial model of 5 category with parameter θ

$$\left(\frac{1}{2}, \frac{\theta}{4}, \frac{1-\theta}{4}, \frac{1-\theta}{4}, \frac{\theta}{4}\right)$$

Can we figure out the number of Monkey A based on the data?

(Cont.) The absent-minded biologist

- ▶ data $y = (125, 18, 20, 34)$
- ▶ now assume $y_1 = y_{11} + y_{12} = 125$
- ▶ Likelihood function

$$f(y|\theta) = \frac{n!}{y_{11}!y_{12}!y_2!y_3!y_4!} \left(\frac{1}{2}\right)^{y_{11}} \left(\frac{\theta}{4}\right)^{y_{12}} \left(\frac{1}{4} - \frac{\theta}{4}\right)^{y_2} \left(\frac{1}{4} - \frac{\theta}{4}\right)^{y_3} \left(\frac{\theta}{4}\right)^{y_4}$$

- ▶ log-likelihood

$$\ell(\theta|y) \propto (\textcolor{red}{y}_{12} + y_4) \log \theta + (y_2 + y_3) \log(1 - \theta)$$

- ▶ y_{12} unknown, cannot directly maximize $\ell(\theta|y)$

(Cont.) The absent-minded biologist: set-up EM

$$\begin{aligned}Q(\theta|\theta') &= \mathbb{E}_{y_{12}}[(y_{12} + y_4) \log \theta + (y_2 + y_3) \log(1 - \theta) | y_1, \dots, y_4, \theta'] \\&= (\mathbb{E}_{y_{12}}[y_{12} | y_1, \theta'] + y_4) \log \theta + (y_2 + y_3) \log(1 - \theta)\end{aligned}$$

Conditional distribution of y_{12} given y_1 : Binomial $(y_1, \frac{\theta'/4}{\theta'/4+1/2})$

$$\mathbb{E}_{y_{12}}[y_{12} | y_1, \theta'] = \frac{y_1 \theta'}{2 + \theta'} := y_{12}^{\theta'},$$

E-step:

$$Q(\theta|\theta') = (y_{12}^{\theta'} + y_4) \log \theta + (y_2 + y_3) \log(1 - \theta)$$

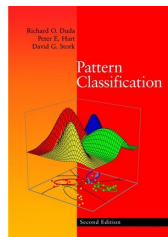
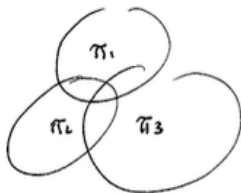
M-step: $\theta_{k+1} = \arg \max Q(\theta|\theta_k) = \frac{y_{12}^{(\theta_k)} + y_4}{y_{12}^{(\theta_k)} + y_2 + y_3 + y_4}$

Fitting Gaussian mixture model (GMM)

$$x_i \sim \sum_{i=1}^C \pi_i \phi(x_i | \mu_i, \Sigma_i)$$

ϕ : density of multi-variate normal

- ▶ parameters $\{\mu_i, \Sigma_i, \pi_i, C\}$
- ▶ assume C is known.
- ▶ observed data $\{x_1, \dots, x_n\}$
- ▶ complete data $\{(x_1, y_1), \dots, (x_n, y_n)\}$
 y_n : "label" for each sample, missing.



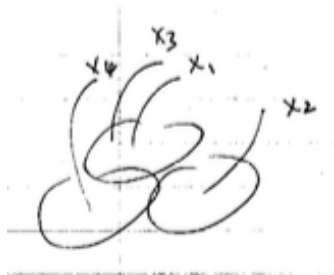
EM for GMM

- ▶ If we know the label information y_i , likelihood function can be easily written

$$\pi_{y_i} \phi(x_i | \mu_{y_i}, \Sigma_{y_i})$$

- ▶ now y_i unknown, compute its expectation with respect to the set of parameters

$$Q(\theta | \theta') = \mathbb{E} \left[\sum_{i=1}^n \log \pi_{y_i} + \log \phi(x_i | \mu_{y_i}, \Sigma_{y_i}) | x_i, \theta' \right]$$



E-step

- ▶ $(\pi_i^{(k)}, \mu_i^{(k)}, \Sigma_i^{(k)})$ parameter values in the k th iteration
- ▶ we need $y_i|x_i$, posterior distribution of label, given observation x_i

$$p_{i,c} := p(y_i = c|x_i) \propto \pi_c^{(k)} \phi(x_i|\mu_i^{(k)}, \Sigma_i^{(k)})$$

$$\text{and } \sum_{c=1}^C p(y_i = c|x_i) = 1$$

$$\begin{aligned} Q(\theta|\theta_k) &= \sum_{i=1}^n \mathbb{E}[\log \pi_{y_i} + \log \phi(x_i|\mu_{y_i}, \Sigma_{y_i}|x_i, \theta_k)] \\ &= \sum_{i=1}^n \sum_{c=1}^C p_{i,c} \log \pi_c + \sum_{i=1}^n \sum_{c=1}^C p_{i,c} \log \phi(x_i|\mu_c, \Sigma_c) \end{aligned}$$

Q: where is θ ?

M-step

- Maximize $Q(\theta|\theta_k)$ with respect to π_c, μ_c, Σ_c (note that they can be maximized separately)

$$\theta_{k+1} = \arg \max_{\theta} Q(\theta|\theta_k)$$

- note that $\sum_{c=1}^C \pi_c = 1$

$\vdots$

in the end

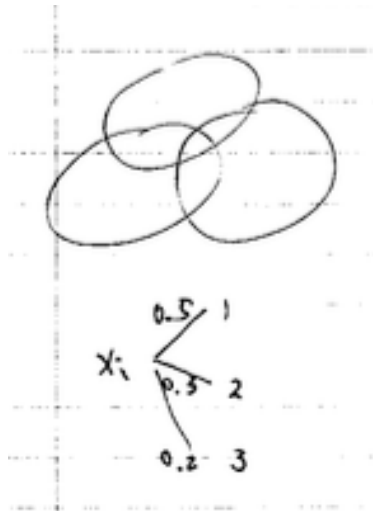
$$\mu_c^{(k+1)} = \frac{\sum_{i=1}^n p_{i,c} x_i}{\sum_{i=1}^n p_{i,c}}$$

$$\Sigma_c^{(k+1)} = \frac{\sum_{i=1}^n p_{i,c} (x_i - \mu_c^{(k+1)})(x_i - \mu_c^{(k+1)})^T}{\sum_{i=1}^n p_{i,c}}$$

$$\pi_c^{(k+1)} = \frac{1}{n} \sum_{i=1}^n p_{i,c}$$

Interpretation

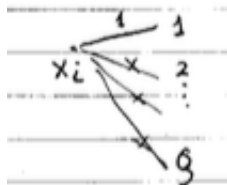
- ▶ $p_{i,c}$: probability of each sample belong to computer c
- ▶ $\pi_c^{(k+1)}$: count the expected number of samples belong to component c
- ▶ soft-assignment: x_i belong to component c with assignment probability $p_{i,c}$
- ▶ $\mu_c^{(k+1)}$: “average” centroid using soft assignment
- ▶ $\Sigma_c^{(k+1)}$: “average” covariance using soft assignment



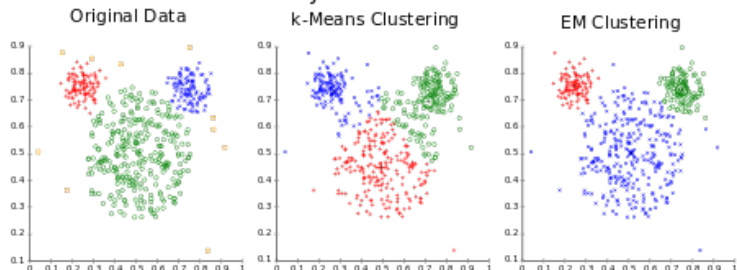
k-means

- ▶ “hard” assignment
- ▶ EM algorithm can also be used for clustering: in the end, $p_{i,c}$ can be viewed as a soft label for each sample
- ▶ convert into hard label:

$$\hat{c}_i = \arg \max_{c=1}^C p_{i,c}$$



Different cluster analysis results on "mouse" data set:



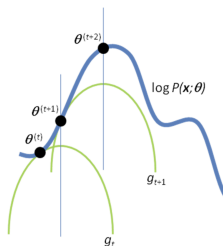
Properties of EM

- ▶ EM algorithm converges to local maximum
- ▶ Heuristic: escaping the local maximum through a random start
- ▶ EM works on improving $Q(\theta|\theta')$ rather than directly improving $\log f(x|\theta)$
- ▶ one can show that improvement on $Q(\theta|\theta')$ improves $\log f(x|\theta)$
- ▶ EM works well with exponential family
 - ▶ E-step: sum of expectations of the sufficient statistics
 - ▶ M-step: maximizing a linear function

usually possible to derive closed-form update

Convergence of EM

- ▶ Proof by A. Dempster, N. Laird and D. Rubin in 1977, later generalized by J. Wu in 1983.
- ▶ Basic idea: find a sequence of quadratic lower bounds for the likelihood function
- ▶ EM monotonically increases the observed data log likelihood



Supplementary Figure 1 Convergence of the EM algorithm. Starting from initial parameters $\theta^{(0)}$, the E-step of the EM algorithm constructs a function $\hat{g}_t$ that lower-bounds the objective function $\log P(\mathbf{x}; \theta)$. In the M-step, $\theta^{(t+1)}$ is computed as the maximum of $\hat{g}_t$. In the next E-step, a new lower-bound $\hat{g}_{t+1}$ is constructed; maximization of $\hat{g}_{t+1}$ in the next M-step gives $\theta^{(t+2)}$, etc.

$$\ell(\theta_{k+1}) \geq Q(\theta_{k+1}; \theta_k) \geq Q(\theta_k; \theta_k) = \ell(\theta_k)$$