CS 412 Introduction to Machine Learning

Gaussian Mixture Model (GMM)

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Gaussian Mixture Model (GMM)

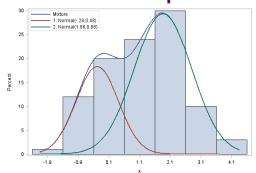
- Gaussian mixture distribution is written as
 - a linear superposition of κ Gaussian components:

$$p(x) = \sum_{k=1}^{K} \pi_k N(x \mid \mu_k, \Sigma_k)$$

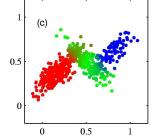
- Represent k by a κ -dimensional binary variable z
 - Using 1-of-K representation (one-hot vector)
 - Let $z = z_1,...,z_K$ whose elements are

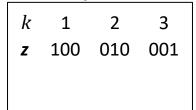
$$z_k \in \{0,1\} \text{ and } \sum_{k} z_k = 1$$

K possible states of z corresponding to K components



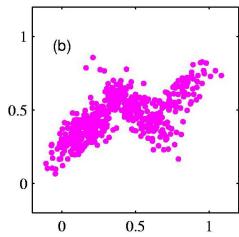
$$k$$
 1 2 z 10 01 π_k 0.4 0.6 μ_k -28 1.86 σ_k 0.48 0.88





Joint Distribution

- Define joint distribution of latent variable and observed variable
 - $-p(x,z)=p(x|z) \cdot p(z)$
 - x is observed variable: a feature vector
 - z is the hidden variable: cluster assignment
 - Prior prob. distribution p(z)
 - Likelihood prob. distribution p(x|z)



Specifying the prior prob. p(z)

- Associate a probability with each component z_k
 - Denote $p(z_k = 1) = \pi_k$ where parameters $\{\pi_k\}$ satisfy $0 \le \pi_k \le 1$ and $\sum_k \pi_k = 1$
- Because z uses 1-of-κ it follows that

$$p(\mathbf{z}) = \prod_{k=1}^K \pi_k^{z_k}$$

With one component $p(z_1) = \pi_1^{z_1}$

With two components $p(z_1, z_2) = \pi_1^{z_1} \pi_2^{z_2}$

Specifying the Likelihood prob. p(x|z)

For a particular component (value of z)

$$p(x \mid z_k = 1) = N(x \mid \mu_k, \Sigma_k)$$

• Thus p(x|z) can be written in the form

$$p(x \mid z) = \prod_{k=1}^{K} N \left(x \mid \mu_{k} \Sigma_{k} \right)^{z_{k}}$$

All product terms except for one equal one

Marginal distribution p(x)

- The joint distribution p(x,z) is given by p(z)p(x|z)
- Thus marginal distribution of x is obtained by summing over all possible states of z to give

$$p(x) = \sum_{z} p(z)p(x \mid z) = \sum_{z} \prod_{k=1}^{K} \pi_{k}^{z_{k}} N \left(x \mid \mu_{k}, \Sigma_{k} \right)^{x_{k}} = \sum_{k=1}^{K} \pi_{k} N \left(x \mid \mu_{k}, \Sigma_{k} \right)$$
Circle 7. = (0.1)

- Since z_k ∈ {0,1}
- This is the standard form of a Gaussian mixture

Gaussian Mixture Model (GMM)

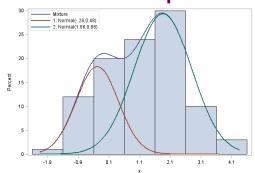
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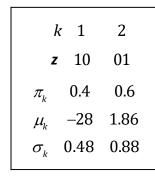
$$p(x) = \sum_{k=1}^{K} \pi_k N(x \mid \mu_k, \Sigma_k)$$

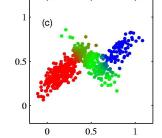
- Represent κ by a κ -dimensional binary variable z
 - Using 1-of-K representation (one-hot vector)
 - Let $z = z_1,...,z_K$ whose elements are

$$z_k \in \{0,1\} \text{ and } \sum_{k} z_k = 1$$

K possible states of z corresponding to K components







k	1	2	3
Z	100	010	001

Synthesizing data from mixture

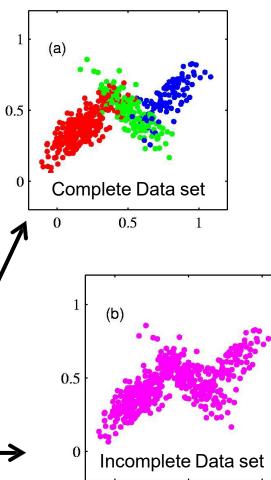
Generate sample of z, called zˆ

 Then generate a value for x from conditional p(x| z[^])

 Samples from p(x,z) are plotted according to value of x and colored with value of z

• Samples from marginal p(x) obtained by ignoring values of z

500 points from three Gaussians



0

0.5

Learning: expectation maximization (EM)

Another conditional probability (Responsibility)

- In EM $p(z \mid x)$ plays a role (posterior in classification)
- The probability $p(z_k=1 \mid x)$ is denoted $\gamma(z_k)$
 - From Bayes theorem

$$\gamma(z_k) = p(z_k = 1 \mid x) = \frac{p(z_k = 1)p(x \mid z_k = 1)}{\sum_{j=1}^{K} p(z_j = 1)p(x \mid z_j = 1)}$$
$$= \frac{\pi_k N(x \mid \mu_k, \Sigma_k)}{\sum_{j=1}^{K} \pi_j N(x \mid \mu_k, \Sigma_j)}$$

- View $p(z_k = 1) = \pi_k$ as prior probability of component k and $\gamma(z_k) = p(z_k = 1 | x)$ as the posterior probability

Maximum Likelihood for GMM

• We wish to model data set $\{x_1,...x_N\}$ using a mixture of Gaussians (N items each of dimension D)

Find maximum likelihood parameters π_k , μ_k , Σ_k

Likelihood Function for GMM

Mixture density function is

$$p(\mathbf{x}) = \sum_{\mathbf{z}} p(\mathbf{z}) p(\mathbf{x} \mid \mathbf{z}) = \sum_{k=1}^{K} \pi_{k} N\left(\mathbf{x} \mid \mu_{k}, \Sigma_{k}\right)$$

Therefore Likelihood function is

$$p(X \mid \pi, \mu, \Sigma) = \prod_{n=1}^{N} \left\{ \sum_{k=1}^{K} \pi_{k} N(x_{n} \mid \mu_{k}, \Sigma_{k}) \right\}$$
Product is over the N i.i.d. samples

Therefore log-likelihood function is

$$\ln p(X \mid \pi, \mu, \Sigma) = \sum_{n=1}^{N} \ln \left\{ \sum_{k=1}^{K} \pi_{k} N(x_{n} \mid \mu_{k}, \Sigma_{k}) \right\}$$

Which we wish to maximize

A more difficult problem than for a single Gaussian

Maximization of Log-Likelihood

$$\ln p(X \mid \pi, \mu, \Sigma) = \sum_{n=1}^{N} \ln \left\{ \sum_{k=1}^{K} \pi_k N(x_n \mid \mu_k, \Sigma_k) \right\}$$

Goal is to estimate the three sets of parameters

$$\pi_k$$
, μ_k , Σ_k

- By taking derivatives in turn w.r.t each while keeping others constant
- But there are no closed-form solutions
- While a gradient-based optimization is possible, we consider the iterative EM algorithm

EM for Gaussian Mixtures

- EM is a method for finding maximum likelihood solutions for models with latent variables
- Begin with log-likelihood function

$$\ln p(X \mid \pi, \mu, \Sigma) = \sum_{n=1}^{N} \ln \left\{ \sum_{k=1}^{K} \pi_{k} N(\mathbf{x}_{n} \mid \mu_{k}, \Sigma_{k}) \right\}$$

– We wish to find π, μ, Σ that maximize this quantity

$$Q(\theta,\theta^0) = E_{Z|X,\theta^0} \left[\log(P(X,Z|\theta)) \right] = \sum_Z P(Z|X,\theta^0) \log(P(X,Z|\theta))$$

Instead take derivatives in turn of Q w.r.t O

- Means μ_k and set to zero
- covariance matrices \sum_k and set to zero
- mixing coefficients π_k and set to zero

K-MEANS ALGORITHM REMINDER

- 1. Initialize means $\mu_{\scriptscriptstyle k}$
 - E Step: Assign each point to a cluster
 - 3. M Step: Given clusters, refine mean $\,\mu_k^{}$ of each cluster k
- Stop when change in means is small

EXPECTATION MAXIMIZATION (EM) FOR GAUSSIAN MIXTURES

- 1. Initialize Gaussian* parameters: means μ_k , covariances Σ_k and mixing coefficients π_k
 - 2. **E Step:** Assign each point X_n an assignment (z_{nk}) for each cluster (z_{nk})

0.5 0.5

- 3. **M Step:** Given scores, adjust μ_k , π_k , Σ_k for each cluster k
- Evaluate likelihood. If likelihood or parameters converge, stop.

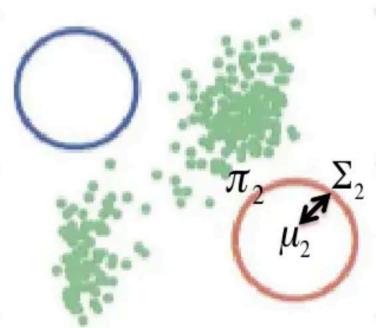
^{*}There are k Gaussians

1. Initialize μ_k , Σ_k π_k , one for each Gaussian k

Tip! Use K-means result to initialize:

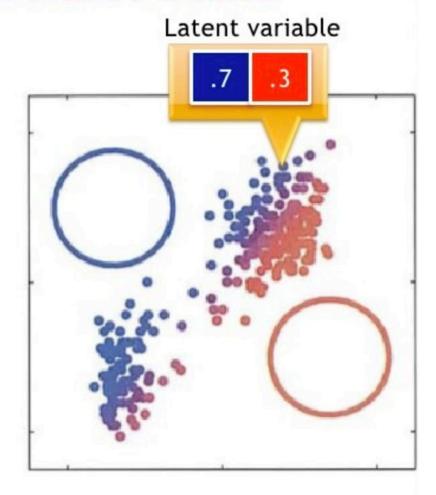
$$\mu_k \leftarrow \mu_k$$

$$\sum_k \leftarrow \text{cov}(cluster(K))$$
 $\pi_k \leftarrow \frac{\text{Number of points in k}}{\text{Total number of points}}$



2. E Step: For each point X_n, determine its assignment score to each Gaussian k:

$$\gamma(z_{nk}) = \frac{\pi_k \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j)}$$



 $\gamma(z_{nk})$ is called a "responsibility": how much is this Gaussian k responsible for this point X_n ?

3. **M Step:** For each Gaussian k, update parameters using new $\gamma(z_{nk})$

Mean of Gaussian k

$$\mu_k^{\text{new}} = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) \mathbf{x}_n$$

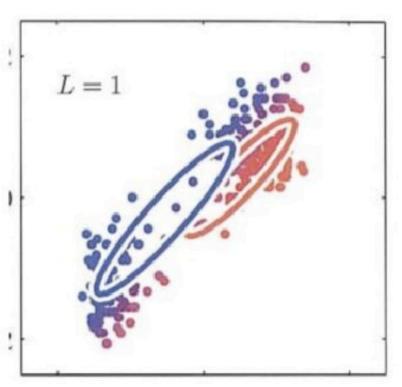
$$N_k = \sum_{n=1}^N \gamma(z_{nk})$$

Responsibility

for this Xn

Find the mean that "fits" the assignment scores best

3. **M Step:** For each Gaussian k, update parameters using new $\gamma(z_{nk})$



Covariance matrix of Gaussian k

$$\Sigma_k^{\text{new}} = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) \left(\mathbf{x}_n - \boldsymbol{\mu}_k^{\text{new}} \right) \left(\mathbf{x}_n - \boldsymbol{\mu}_k^{\text{new}} \right)^{\text{T}}$$

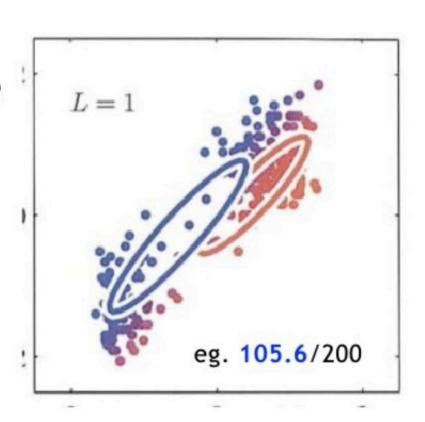
Just calculated this!

3. **M Step:** For each Gaussian k, update parameters using new $\gamma(z_{nk})$

Mixing Coefficient for Gaussian k

$$\pi_k^{\text{new}} = \frac{N_k}{N}$$

Total # of points



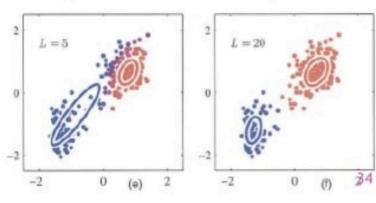
$$N_k = \sum_{n=1}^N \gamma(z_{nk})$$

 Evaluate log likelihood. If likelihood or parameters converge, stop. Else go to Step 2 (E step).

$$\ln p(\mathbf{X}|\boldsymbol{\mu}, \boldsymbol{\Sigma}, \boldsymbol{\pi}) = \sum_{n=1}^{N} \ln \left\{ \sum_{k=1}^{K} \underline{\pi_k} \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \right\}$$

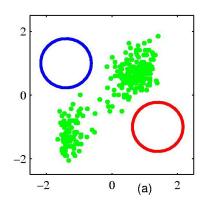
Likelihood is the probability that the data X was generated by the parameters you found.

ie. Correctness!

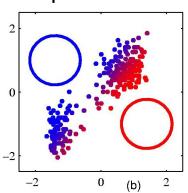


EM Example

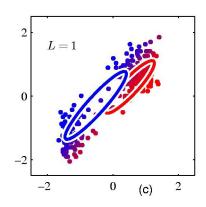
Data points and Initial mixture model



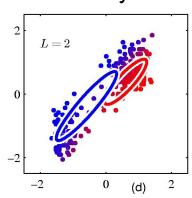
Initial E step Determine responsibilities



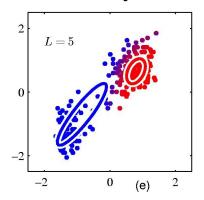
After first M step Re-evaluate Parameters



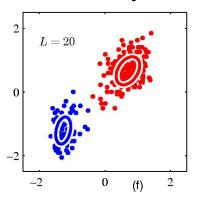
After 2 cycles



After 5 cycles

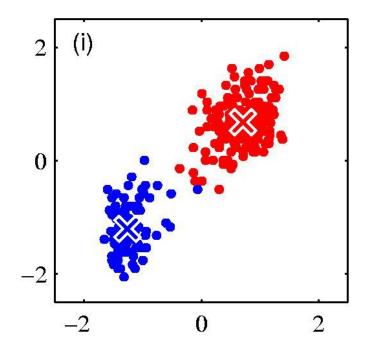


After 20 cycles

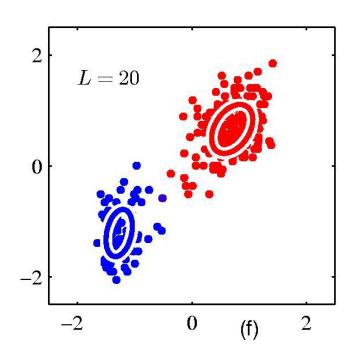


Comparison with K-Means

K-means result



E-M result



Practical Issues with EM

- Takes many more iterations than K-means
 - Each cycle requires significantly more comparison
- Common to run K-means first in order to find suitable initialization
- Covariance matrices can be initialized to covariances of clusters found by K-means
- EM is not guaranteed to find global maximum of log likelihood function

Attributes:

weights_: array-like of shape (n_components,)

The weights of each mixture components.

means_: array-like of shape (n_components, n_features)

The mean of each mixture component.

covariances_: array-like

The covariance of each mixture component.

Summary of EM for GMM

- Given a Gaussian mixture model
- Goal is to maximize the likelihood function w.r.t. the parameters (means, covariances and mixing coefficients)

Step1: Initialize the means μ_k covariances Σ_k and mixing coefficients π_k and evaluate initial value of log-likelihood

EM continued

 Step 2: E step: Evaluate responsibilities using current parameter values

$$\gamma(z_k) = \frac{\pi_k N(x_n \mid \mu_k, \Sigma_k)}{\sum_{j=1}^K \pi_j N(x_n \mid \mu_j, \Sigma_j))}$$

 Step 3: M Step: Re-estimate parameters using current responsibilities

$$\mu_k^{\text{new}} = \frac{1}{N_k} \sum_{n=1}^N \gamma(z_{nk}) \mathbf{x}_n$$

$$\sum_{k}^{\text{new}} = \frac{1}{N_k} \sum_{n=1}^{N} \gamma(z_{nk}) (\mathbf{x}_n - \mu_k^{\text{new}}) (\mathbf{x}_n - \mu_k^{\text{new}})^T$$

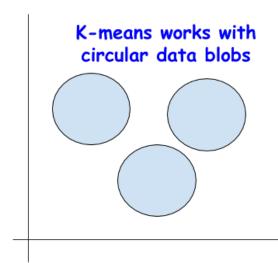
$$\pi_k^{\text{new}} = \frac{N_k}{N} \quad \text{where} \quad N_k = \sum_{n=1}^N \gamma(z_{nk})$$

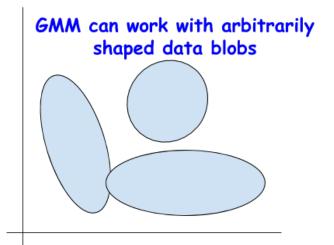
EM Continued

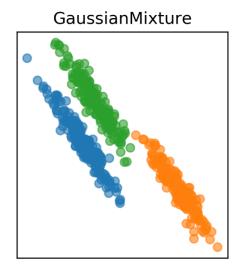
Step 4: Evaluate the log likelihood

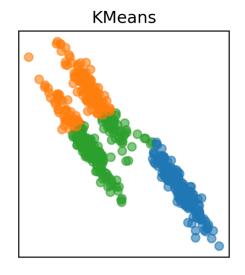
$$\ln p(X \mid \pi, \mu, \Sigma) = \sum_{n=1}^{N} \ln \left\{ \sum_{k=1}^{K} \pi_{k} N(\mathbf{x}_{n} \mid \mu_{k}, \Sigma_{k}) \right\}$$

- And check for convergence of either parameters or log likelihood
- If convergence not satisfied return to Step 2









Different cluster analysis results on "mouse" data set:

