Parametric Models Part II: Expectation-Maximization and Mixture Density Estimation

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Missing Features

- Suppose that we have a Bayesian classifier that uses the feature vector x but a subset x_g of x are observed and the values for the remaining features x_b are missing.
- ▶ How can we make a decision?
 - Throw away the observations with missing values.
 - Or, substitute \mathbf{x}_b by their average $\bar{\mathbf{x}}_b$ in the training data, and use $\mathbf{x} = (\mathbf{x}_q, \bar{\mathbf{x}}_b)$.
 - Or, marginalize the posterior over the missing features, and use the resulting posterior

$$P(w_i|\mathbf{x}_g) = \frac{\int P(w_i|\mathbf{x}_g, \mathbf{x}_b) p(\mathbf{x}_g, \mathbf{x}_b) d\mathbf{x}_b}{\int p(\mathbf{x}_g, \mathbf{x}_b) d\mathbf{x}_b}.$$



- We can also extend maximum likelihood techniques to allow learning of parameters when some training patterns have missing features.
- ► The Expectation-Maximization (EM) algorithm is a general iterative method of finding the maximum likelihood estimates of the parameters of a distribution from training data.

- ▶ There are two main applications of the EM algorithm:
 - ► Learning when the data is incomplete or has missing values.
 - Optimizing a likelihood function that is analytically intractable but can be simplified by assuming the existence of and values for additional but missing (or hidden) parameters.
- ► The second problem is more common in pattern recognition applications.

- Assume that the observed data \mathcal{X} is generated by some distribution.
- Assume that a complete dataset $\mathcal{Z} = (\mathcal{X}, \mathcal{Y})$ exists as a combination of the observed but incomplete data \mathcal{X} and the missing data \mathcal{Y} .
- ► The observations in Z are assumed to be i.i.d. from the joint density

$$p(\mathbf{z}|\mathbf{\Theta}) = p(\mathbf{x}, \mathbf{y}|\mathbf{\Theta}) = p(\mathbf{y}|\mathbf{x}, \mathbf{\Theta})p(\mathbf{x}|\mathbf{\Theta}).$$



We can define a new likelihood function

$$L(\mathbf{\Theta}|\mathcal{Z}) = L(\mathbf{\Theta}|\mathcal{X}, \mathcal{Y}) = p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta})$$

called the complete-data likelihood where $L(\Theta|\mathcal{X})$ is referred to as the incomplete-data likelihood.

- ► The EM algorithm:
 - ► First, finds the expected value of the complete-data log-likelihood using the current parameter estimates (expectation step).
 - ► Then, maximizes this expectation (maximization step).



Define

$$Q(\boldsymbol{\Theta}, \boldsymbol{\Theta}^{(i-1)}) = E[\log p(\mathcal{X}, \mathcal{Y} | \boldsymbol{\Theta}) \, | \, \mathcal{X}, \boldsymbol{\Theta}^{(i-1)}]$$

as the expected value of the complete-data log-likelihood w.r.t. the unknown data $\mathcal Y$ given the observed data $\mathcal X$ and the current parameter estimates $\Theta^{(i-1)}$.

The expected value can be computed as

$$E\left[\log p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta})|\mathcal{X}, \mathbf{\Theta}^{(i-1)}\right] = \int \log p(\mathcal{X}, \mathbf{y}|\mathbf{\Theta}) \, p(\mathbf{y}|\mathcal{X}, \mathbf{\Theta}^{(i-1)}) \, d\mathbf{y}.$$

► This is called the *E-step*.



► Then, the expectation can be maximized by finding optimum values for the new parameters Θ as

$$\Theta^{(i)} = \arg \max_{\Theta} Q(\Theta, \Theta^{(i-1)}).$$

- ► This is called the *M-step*.
- ► These two steps are repeated iteratively where each iteration is guaranteed to increase the log-likelihood.
- ► The EM algorithm is also guaranteed to converge to a local maximum of the likelihood function.

Generalized Expectation-Maximization

▶ Instead of maximizing $Q(\Theta, \Theta^{(i-1)})$, the *Generalized* Expectation-Maximization algorithm finds some set of parameters $\Theta^{(i)}$ that satisfy

$$Q(\mathbf{\Theta}^{(i)}, \mathbf{\Theta}^{(i-1)}) > Q(\mathbf{\Theta}, \mathbf{\Theta}^{(i-1)})$$

at each iteration.

► Convergence will not be as rapid as the EM algorithm but it allows greater flexibility to choose computationally simpler steps.

Mixture Densities

▶ A mixture model is a linear combination of m densities

$$p(\mathbf{x}|\mathbf{\Theta}) = \sum_{j=1}^{m} \alpha_j p_j(\mathbf{x}|\boldsymbol{\theta_j})$$

where $\Theta = (\alpha_1, \dots, \alpha_m, \boldsymbol{\theta_1}, \dots, \boldsymbol{\theta_m})$ such that $\alpha_j \geq 0$ and $\sum_{i=1}^m \alpha_i = 1$.

- $ightharpoonup \alpha_1, \ldots, \alpha_m$ are called the mixing parameters.
- ▶ $p_j(\mathbf{x}|\boldsymbol{\theta_j})$, j = 1, ..., m are called the component densities.



Mixture Densities

- ▶ Suppose that $\mathcal{X} = \{\mathbf{x_1}, \dots, \mathbf{x_n}\}$ is a set of observations i.i.d. with distribution $p(\mathbf{x}|\Theta)$.
- ▶ The log-likelihood function of ⊖ becomes

$$\log L(\boldsymbol{\Theta}|\mathcal{X}) = \log \prod_{i=1}^n p(\mathbf{x_i}|\boldsymbol{\Theta}) = \sum_{i=1}^n \log \bigg(\sum_{j=1}^m \alpha_j p_j(\mathbf{x_i}|\boldsymbol{\theta_j}) \bigg).$$

• We cannot obtain an analytical solution for Θ by simply setting the derivatives of $\log L(\Theta|\mathcal{X})$ to zero because of the logarithm of the sum.

- ▶ Consider \mathcal{X} as incomplete and define hidden variables $\mathcal{Y} = \{y_i\}_{i=1}^n$ where y_i corresponds to which mixture component generated the data vector \mathbf{x}_i .
- ▶ In other words, $y_i = j$ if the i'th data vector was generated by the j'th mixture component.
- ► Then, the log-likelihood becomes

$$\begin{split} \log L(\boldsymbol{\Theta}|\mathcal{X}, \mathcal{Y}) &= \log p(\mathcal{X}, \mathcal{Y}|\boldsymbol{\Theta}) \\ &= \sum_{i=1}^n \log(p(\mathbf{x}_{\boldsymbol{i}}|y_i, \boldsymbol{\theta}_{\boldsymbol{i}})p(y_i|\boldsymbol{\theta}_{\boldsymbol{i}})) \\ &= \sum_{i=1}^n \log(\alpha_{y_i} p_{y_i}(\mathbf{x}_{\boldsymbol{i}}|\boldsymbol{\theta}_{\boldsymbol{y}_i})). \end{split}$$



Assume we have the initial parameter estimates

$$\boldsymbol{\Theta}^{(g)} = (\alpha_1^{(g)}, \dots, \alpha_m^{(g)}, \boldsymbol{\theta}_1^{(g)}, \dots, \boldsymbol{\theta}_m^{(g)}).$$

▶ Compute

$$p(y_i|\mathbf{x}_i, \mathbf{\Theta}^{(g)}) = \frac{\alpha_{y_i}^{(g)} p_{y_i}(\mathbf{x}_i|\boldsymbol{\theta}_{y_i}^{(g)})}{p(\mathbf{x}_i|\mathbf{\Theta}^{(g)})} = \frac{\alpha_{y_i}^{(g)} p_{y_i}(\mathbf{x}_i|\boldsymbol{\theta}_{y_i}^{(g)})}{\sum_{j=1}^m \alpha_j^{(g)} p_j(\mathbf{x}_i|\boldsymbol{\theta}_j^{(g)})}$$

and

$$p(\mathcal{Y}|\mathcal{X}, \mathbf{\Theta}^{(g)}) = \prod_{i=1}^{n} p(y_i|\mathbf{x}_i, \mathbf{\Theta}^{(g)}).$$



▶ Then, $Q(\Theta, \Theta^{(g)})$ takes the form

$$Q(\mathbf{\Theta}, \mathbf{\Theta}^{(g)}) = \sum_{\mathbf{y}} \log p(\mathcal{X}, \mathbf{y} | \mathbf{\Theta}) p(\mathbf{y} | \mathcal{X}, \mathbf{\Theta}^{(g)})$$

$$= \sum_{j=1}^{m} \sum_{i=1}^{n} \log(\alpha_{j} p_{j}(\mathbf{x}_{i} | \boldsymbol{\theta}_{j})) p(j | \mathbf{x}_{i}, \mathbf{\Theta}^{(g)})$$

$$= \sum_{j=1}^{m} \sum_{i=1}^{n} \log(\alpha_{j}) p(j | \mathbf{x}_{i}, \mathbf{\Theta}^{(g)})$$

$$+ \sum_{i=1}^{m} \sum_{j=1}^{n} \log(p_{j}(\mathbf{x}_{i} | \boldsymbol{\theta}_{j})) p(j | \mathbf{x}_{i}, \mathbf{\Theta}^{(g)}).$$

- ▶ We can maximize the two sets of summations for α_j and θ_j independently because they are not related.
- ▶ The estimate for α_i can be computed as

$$\hat{\alpha}_j = \frac{1}{n} \sum_{i=1}^n p(j|\mathbf{x}_i, \mathbf{\Theta}^{(g)})$$

where

$$p(j|\mathbf{x}_i, \mathbf{\Theta}^{(g)}) = \frac{\alpha_j^{(g)} p_j(\mathbf{x}_i | \boldsymbol{\theta}_j^{(g)})}{\sum_{t=1}^m \alpha_t^{(g)} p_t(\mathbf{x}_i | \boldsymbol{\theta}_t^{(g)})}.$$



• We can obtain analytical expressions for θ_j for the special case of a Gaussian mixture where $\theta_j=(\mu_j,\Sigma_j)$ and

$$\begin{split} p_j(\mathbf{x}|\boldsymbol{\theta_j}) &= p_j(\mathbf{x}|\boldsymbol{\mu_j}, \boldsymbol{\Sigma_j}) \\ &= \frac{1}{(2\pi)^{d/2} |\boldsymbol{\Sigma_j}|^{1/2}} \, \exp\left[-\frac{1}{2}(\mathbf{x} - \boldsymbol{\mu_j})^T \boldsymbol{\Sigma_j^{-1}} (\mathbf{x} - \boldsymbol{\mu_j})\right]. \end{split}$$

▶ Equating the partial derivative of $Q(\Theta, \Theta^{(g)})$ with respect to μ_j to zero gives

$$\hat{\boldsymbol{\mu}}_{\boldsymbol{j}} = \frac{\sum_{i=1}^n p(j|\mathbf{x}_{\boldsymbol{i}}, \boldsymbol{\Theta}^{(g)}) \mathbf{x}_{\boldsymbol{i}}}{\sum_{i=1}^n p(j|\mathbf{x}_{\boldsymbol{i}}, \boldsymbol{\Theta}^{(g)})}.$$



• We consider five models for the covariance matrix Σ_j :

$$\mathbf{\Sigma}_{i} = \sigma^{2} \mathbf{I}$$

$$\hat{\sigma}^2 = \frac{1}{nd} \sum_{j=1}^m \sum_{i=1}^n p(j|\mathbf{x}_i, \mathbf{\Theta}^{(g)}) \|\mathbf{x}_i - \hat{\boldsymbol{\mu}}_j\|^2$$

$$\mathbf{\Sigma}_{j} = \sigma_{j}^{2} \mathbf{I}$$

$$\hat{\sigma}_j^2 = \frac{\sum_{i=1}^n p(j|\mathbf{x}_i, \mathbf{\Theta}^{(g)}) \|\mathbf{x}_i - \hat{\boldsymbol{\mu}}_j\|^2}{d\sum_{i=1}^n p(j|\mathbf{x}_i, \mathbf{\Theta}^{(g)})}$$



- Covariance models continued:
 - $\qquad \boldsymbol{\Sigma_j} = \mathrm{diag}(\{\sigma_{jk}^2\}_{k=1}^d)$

$$\hat{\sigma}_{jk}^2 = \frac{\sum_{i=1}^n p(j|\mathbf{x_i}, \boldsymbol{\Theta}^{(g)}) (\mathbf{x_{ik}} - \boldsymbol{\hat{\mu}_{jk}})^2}{\sum_{i=1}^n p(j|\mathbf{x_i}, \boldsymbol{\Theta}^{(g)})}$$

 $\Sigma_i = \Sigma$

$$\hat{\mathbf{\Sigma}} = \frac{1}{n} \sum_{i=1}^{m} \sum_{i=1}^{n} p(j|\mathbf{x}_i, \mathbf{\Theta}^{(g)}) (\mathbf{x}_i - \hat{\boldsymbol{\mu}}_j) (\mathbf{x}_i - \hat{\boldsymbol{\mu}}_j)^T$$

• Σ_i = arbitrary

$$\hat{\boldsymbol{\Sigma}}_{\boldsymbol{j}} = \frac{\sum_{i=1}^n p(j|\mathbf{x}_{\boldsymbol{i}}, \boldsymbol{\Theta}^{(g)}) (\mathbf{x}_{\boldsymbol{i}} - \hat{\boldsymbol{\mu}}_{\boldsymbol{j}}) (\mathbf{x}_{\boldsymbol{i}} - \hat{\boldsymbol{\mu}}_{\boldsymbol{j}})^T}{\sum_{i=1}^n p(j|\mathbf{x}_{\boldsymbol{i}}, \boldsymbol{\Theta}^{(g)})}$$



Summary:

- ▶ Estimates for α_j , μ_j and Σ_j perform both expectation and maximization steps simultaneously.
- ► EM iterations proceed by using the current estimates as the initial estimates for the next iteration.
- ► The priors are computed from the proportion of examples belonging to each mixture component.
- ► The means are the component centroids.
- The covariance matrices are calculated as the sample covariance of the points associated with each component.

Questions:

- How can we find the number of components in the mixture?
- ▶ How can we find the initial estimates for Θ ?
- How do we know when to stop the iterations?
 - Stop if the change in log-likelihood between two iterations is less than a threshold.
 - Or, use a threshold for the number of iterations.

Examples

- Mixture of Gaussians examples
- 1-D Bayesian classification examples
- ▶ 2-D Bayesian classification examples

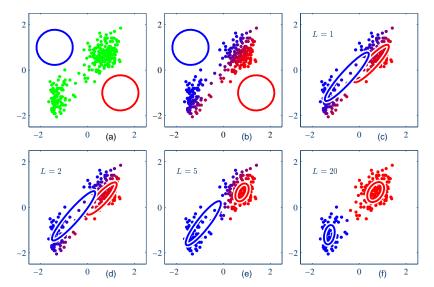
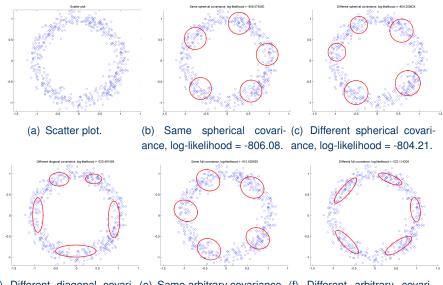


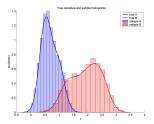
Figure 1: Illustration of the EM algorithm iterations for a mixture of two Gaussians.



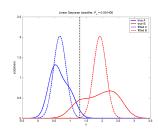


(d) Different diagonal covari- (e) Same arbitrary covariance, (f) Different arbitrary covariance, log-likelihood = -630.46. log-likelihood = -810.93. ance, log-likelihood = -523.11.

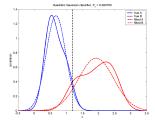
Figure 2: Fitting mixtures of 5 Gaussians to data from a circular distributio



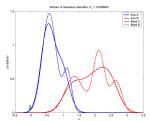
(a) True densities and sample histograms.



(b) Linear Gaussian classifier with P_e = 0.0914.

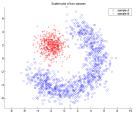


(c) Quadratic Gaussian classifier with $P_e = 0$ (d) Mixture of Gaussian classifier with $P_e = 0$ 0.0837.

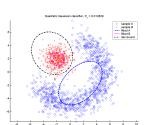


0.0869.

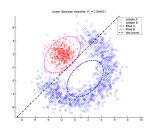
Figure 3: 1-D Bayesian classification examples where the data for each class come from a mixture of three Gaussians. Bayes error is $P_e = 0.0828$.



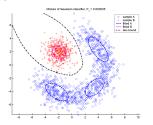




(c) Quadratic Gaussian classifier with $P_e=$ (d) Mixture of Gaussian classifier with $P_e=$ 0.012829.



(b) Linear Gaussian classifier with P_e = 0.094531.



0.002026.

Figure 4: 2-D Bayesian classification examples where the data for the classes come from a banana shaped distribution and a bivariate Gaussian

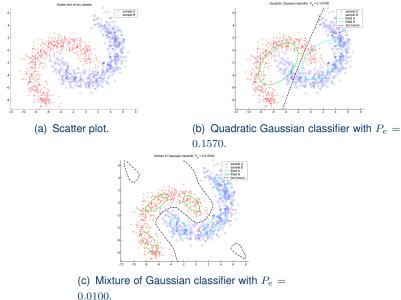


Figure 5: 2-D Bayesian classification examples where the <u>data</u> for each class come from a banana shaped distribution.

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