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 $\mathbf{\bar{x}}_b$ in the training data, and use $\mathbf{x} = (\mathbf{x}_g, \mathbf{\bar{x}}_b)$.
 - 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} .
- \bullet The observations in ${\mathcal Z}$ 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

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

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

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

as the expected value of the complete-data loglikelihood 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

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

• 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

 \bullet 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 $\Theta = (\alpha_1, \dots, \alpha_m, \theta_1, \dots, \theta_m)$ such that $\alpha_j \ge 0$ and $\sum_{j=1}^m \alpha_j = 1$.

- $\alpha_1, \ldots, \alpha_m$ are called the mixing parameters.
- $p_j(\mathbf{x}|\boldsymbol{\theta_j})$, $j = 1, \dots, m$ 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 $p(\mathbf{x}|\mathbf{\Theta})$.
- \bullet The log-likelihood function of Θ becomes

$$\log L(\boldsymbol{\Theta}|\mathcal{X}) = \log \prod_{i=1}^{n} p(\mathbf{x}_i|\boldsymbol{\Theta}) = \sum_{i=1}^{n} \log \left(\sum_{j=1}^{m} \alpha_j p_j(\mathbf{x}_i|\boldsymbol{\theta}_j) \right).$$

• 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 X as incomplete and define hidden variables Y = {y_i}ⁿ_{i=1} where y_i corresponds to which mixture component generated the data vector 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

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

- Assume we have the initial parameter estimates $\Theta^{(g)} = (\alpha_1^{(g)}, \dots, \alpha_m^{(g)}, \boldsymbol{\theta}_1^{(g)}, \dots, \boldsymbol{\theta}_m^{(g)}).$
- Compute

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

and

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

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

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

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

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

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

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

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

where

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

• We can obtain analytical expressions for $m{ heta}_{j}$ for the special case of a Gaussian mixture where $m{ heta}_{j}=(m{\mu}_{j}, m{\Sigma}_{j})$ and

$$p_{j}(\mathbf{x}|\boldsymbol{\theta}_{j}) = p_{j}(\mathbf{x}|\boldsymbol{\mu}_{j}, \boldsymbol{\Sigma}_{j})$$
$$= \frac{1}{(2\pi)^{d/2} |\boldsymbol{\Sigma}_{j}|^{1/2}} \exp\left[-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu}_{j})^{T} \boldsymbol{\Sigma}_{j}^{-1}(\mathbf{x}-\boldsymbol{\mu}_{j})\right].$$

• 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:
 Σ_j = σ²I

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

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

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

• Covariance models continued:

• $\Sigma_j = \operatorname{diag}(\{\sigma_{jk}^2\}_{k=1}^d)$

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

 $\blacktriangleright \Sigma_j = \Sigma$

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

• $\Sigma_j = {\sf arbitrary}$

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

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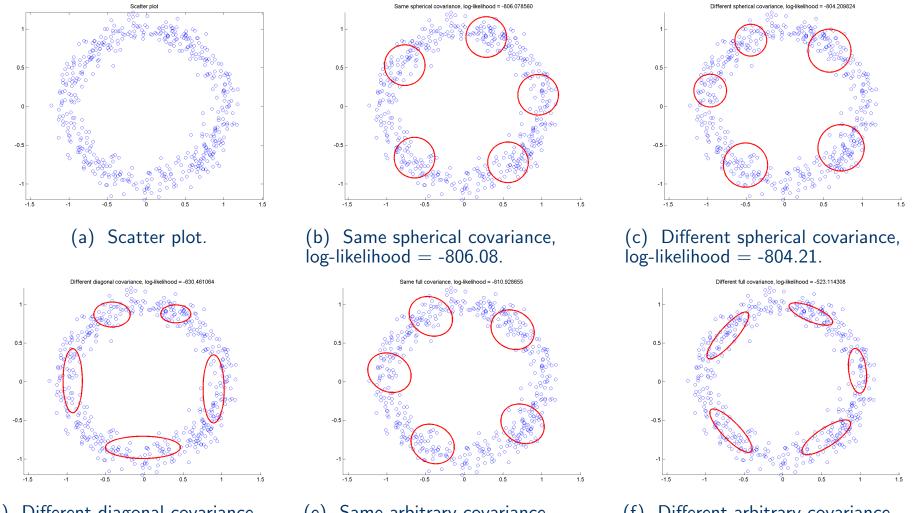
• Summary:

- 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

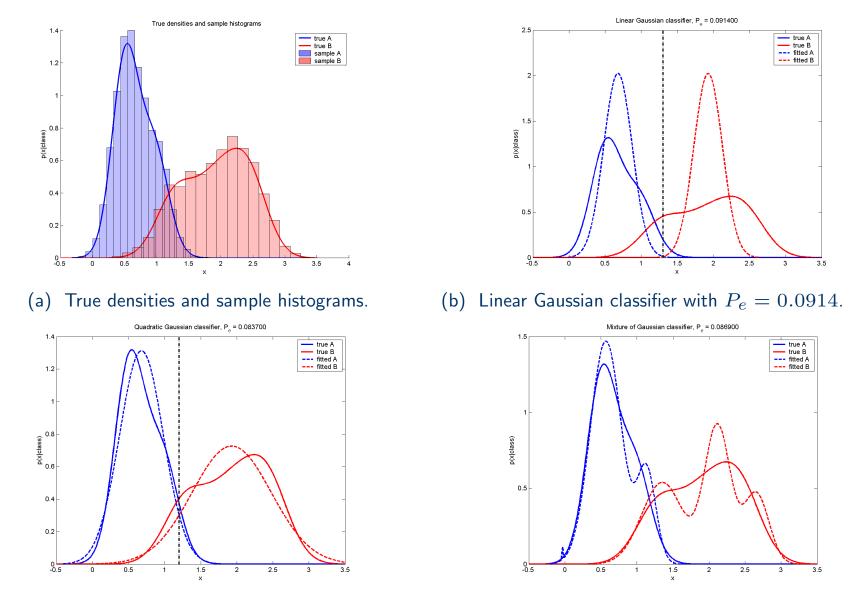


(d) Different diagonal covariance, log-likelihood = -630.46.

(e) Same arbitrary covariance, log-likelihood = -810.93.

(f) Different arbitrary covariance, log-likelihood = -523.11.

Figure 1: Fitting mixtures of 5 Gaussians to data from a circular distribution.



(c) Quadratic Gaussian classifier with $P_e = 0.0837$.

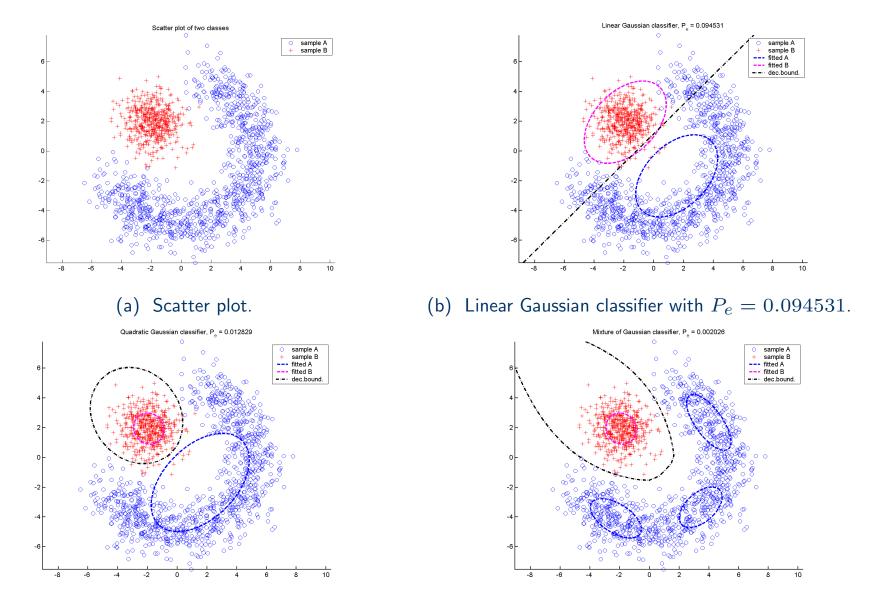
(d) Mixture of Gaussian classifier with $P_e = 0.0869$.

Figure 2: 1-D Bayesian classification examples where the data for each class come from a mixture of three Gaussians. Bayes error is $P_e = 0.0828$.

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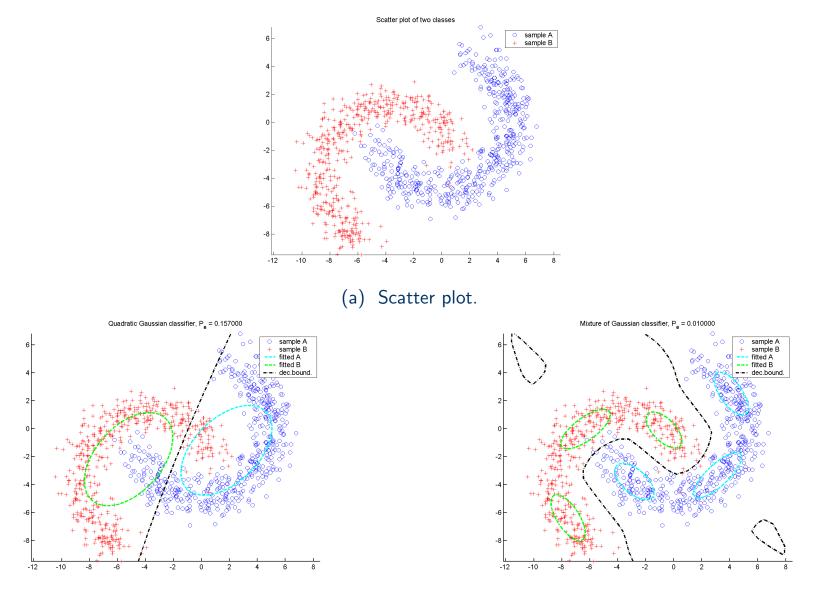
(c) Quadratic Gaussian classifier with $P_e = 0.012829$. (d) Mixture of Gaussian classifier with $P_e = 0.002026$.

Figure 3: 2-D Bayesian classification examples where the data for the classes come from a banana shaped distribution and a bivariate Gaussian.

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(b) Quadratic Gaussian classifier with $P_e = 0.1570$. (c) Quadratic Gaussian classifier with $P_e = 0.0100$.

Figure 4: 2-D Bayesian classification examples where the data for each class come from a banana shaped distribution.

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