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 \mathbf{x} but a subset \mathbf{x}_g of \mathbf{x} are observed and the values for the remaining features \mathbf{x}_b are missing.
- How can we make a decision?
 - ▶ Throw away the observations with missing values.
 - $lackbox{ Or, substitute } \mathbf{x}_b$ 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

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

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- 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.

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- 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.

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- ullet 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} .
- ullet The observations in ${\mathcal Z}$ are assumed to be i.i.d. from the joint density

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

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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 $L(\mathbf{\Theta}|\mathcal{X})$ 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).

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Define

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

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 $\mathbf{\Theta}^{(i-1)}$.

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*.

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ullet Then, the expectation can be maximized by finding optimum values for the new parameters $oldsymbol{\Theta}$ as

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

- 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.

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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.

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Mixture Densities

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

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

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Mixture Densities

- Suppose that $\mathcal{X} = \{\mathbf{x}_1, \dots, \mathbf{x}_n\}$ is a set of observations i.i.d. with distribution $p(\mathbf{x}|\mathbf{\Theta})$.
- ullet The log-likelihood function of ullet 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.

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- 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

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

$$= \sum_{i=1}^{n} \log(p(\mathbf{x}_{i}|y_{i}, \boldsymbol{\theta}_{i})p(y_{i}|\boldsymbol{\theta}_{i}))$$

$$= \sum_{i=1}^{n} \log(\alpha_{y_{i}}p_{y_{i}}(\mathbf{x}_{i}|\boldsymbol{\theta}_{y_{i}})).$$

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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

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

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

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

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

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

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

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- 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

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

where

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

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ullet 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 $Q(\mathbf{\Theta},\mathbf{\Theta}^{(g)})$ with respect to $\pmb{\mu_j}$ to zero gives

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

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• We consider five models for the covariance matrix Σ_j :

$$\Sigma_j = \sigma^2 \mathbf{I}$$

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

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

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

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- Covariance models continued:
 - $\mathbf{\Sigma_{j}} = \operatorname{diag}(\{\sigma_{jk}^2\}_{k=1}^d)$

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

lacksquare $\Sigma_j = \Sigma$

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

 $ar{\Sigma}_j = \mathsf{arbitrary}$

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

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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.

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- 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.

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Examples

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

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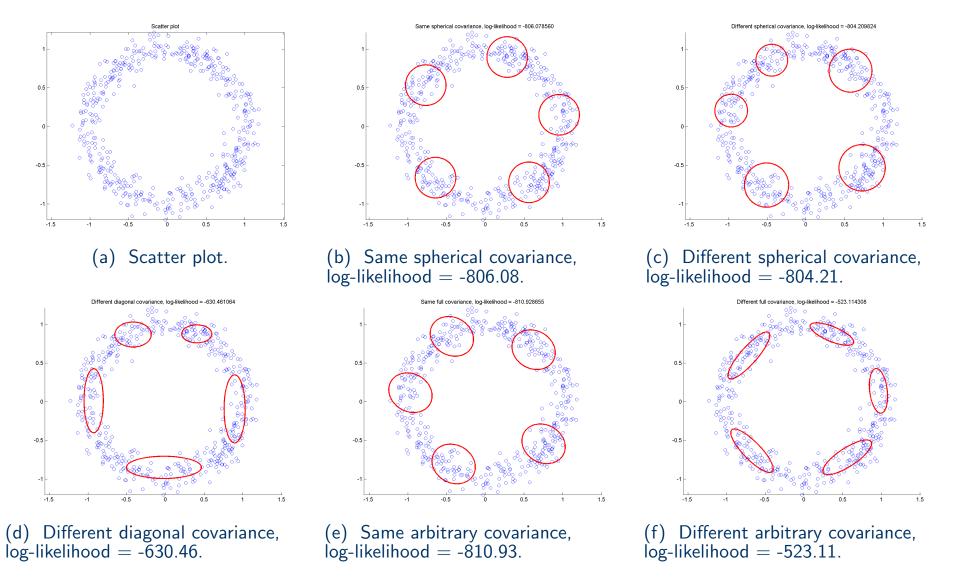
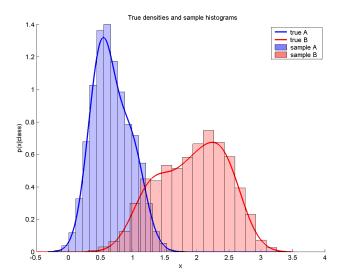
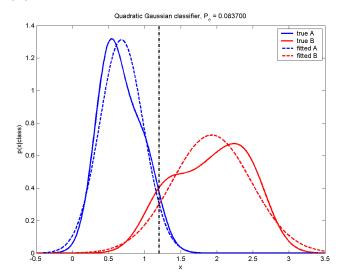


Figure 1: Fitting mixtures of 5 Gaussians to data from a circular distribution.

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(a) True densities and sample histograms.



Linear Gaussian classifier, P_e = 0.091400

2.5

1.5

2

1.5

0.5

0.5

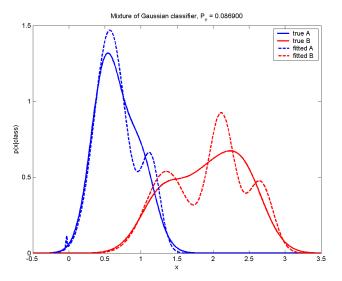
1.5

2

2.5

3.5

(b) Linear Gaussian classifier with $P_e = 0.0914$.

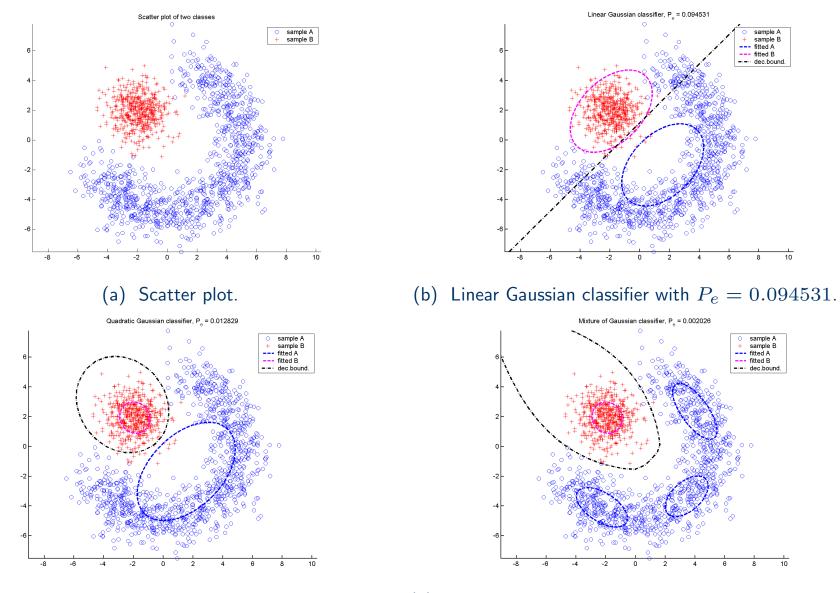


(c) Quadratic Gaussian classifier with $P_e = 0.0837$.

(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 $P_e=0.0828$.

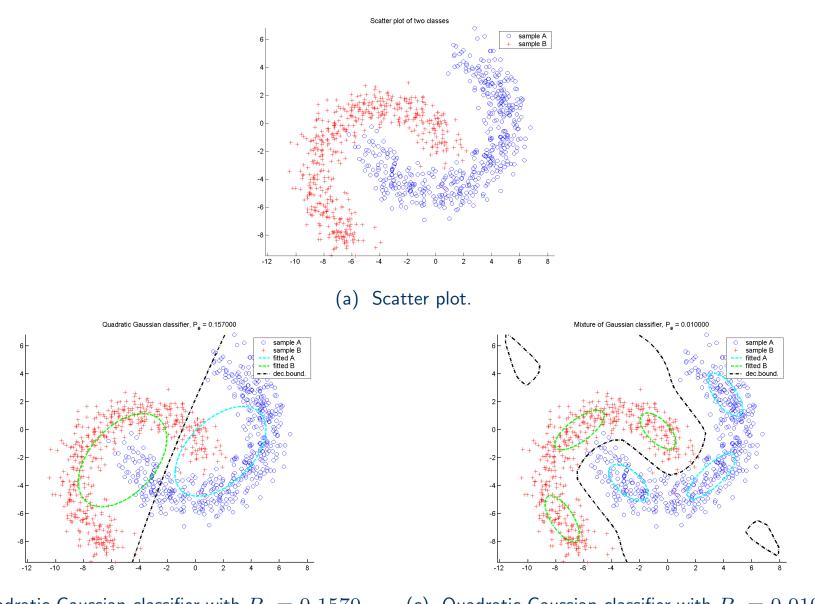
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(c) Quadratic Gaussian classifier with $P_e=0.012829$. (d) Mixture of Gaussian classifier with $P_e=0.002026$.

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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