

# Parametric Models

## Part II: Expectation-Maximization and Mixture Density Estimation

Selim Aksoy  
Bilkent University  
Department of Computer Engineering  
saksoy@cs.bilkent.edu.tr

# 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.
  - ▶ Or, substitute  $\mathbf{x}_b$  by their average  $\bar{\mathbf{x}}_b$  in the training data, and use  $\mathbf{x} = (\mathbf{x}_g, \bar{\mathbf{x}}_b)$ .
  - ▶ Or, marginalize the posterior over the missing features, and use the resulting posterior

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

# Expectation-Maximization

- 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 method of finding the maximum likelihood estimates of the parameters of a distribution from training data.

# Expectation-Maximization

- 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.

# Expectation-Maximization

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

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

# Expectation-Maximization

- We can define a new likelihood function

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

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).

# Expectation-Maximization

- Define

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

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

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

- This is called the *E-step*.

# Expectation-Maximization

- Then, the expectation can be maximized by finding optimum values for the new parameters  $\Theta$  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

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

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

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

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

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

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

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

# Mixture Density Estimation via EM

- 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

$$\begin{aligned}\log L(\Theta | \mathcal{X}, \mathcal{Y}) &= \log p(\mathcal{X}, \mathcal{Y} | \Theta) \\ &= \sum_{i=1}^n \log(p(\mathbf{x}_i | y_i, \theta_i) p(y_i | \theta_i)) \\ &= \sum_{i=1}^n \log(\alpha_{y_i} p_{y_i}(\mathbf{x}_i | \theta_{y_i}))\end{aligned}$$

# Mixture Density Estimation via EM

- Assume we have the initial parameter estimates  $\Theta^{(g)} = (\alpha_1^{(g)}, \dots, \alpha_m^{(g)}, \theta_1^{(g)}, \dots, \theta_m^{(g)})$ .

- Compute

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

and

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

# Mixture Density Estimation via EM

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

$$\begin{aligned} Q(\Theta, \Theta^{(g)}) &= \sum_y \log p(\mathcal{X}, y | \Theta) p(y | \mathcal{X}, \Theta^{(g)}) \\ &= \sum_{j=1}^m \sum_{i=1}^n \log(\alpha_j p_j(\mathbf{x}_i | \theta_j)) p(j | \mathbf{x}_i, \Theta^{(g)}) \\ &= \sum_{j=1}^m \sum_{i=1}^n \log(\alpha_j) p(j | \mathbf{x}_i, \Theta^{(g)}) \\ &\quad + \sum_{j=1}^m \sum_{i=1}^n \log(p_j(\mathbf{x}_i | \theta_j)) p(j | \mathbf{x}_i, \Theta^{(g)}) \end{aligned}$$

# Mixture Density Estimation via EM

- We can maximize the two sets of summations for  $\alpha_j$  and  $\theta_j$  independently because they are not related.
- The expression for  $\alpha_j$  can be computed as

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

# Mixture of Gaussians

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

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

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

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



# Mixture of Gaussians

- We consider five models for the covariance matrix  $\Sigma_j$ :
  - ▶  $\Sigma_j = \sigma^2 \mathbf{I}$

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

- ▶  $\Sigma_j = \sigma_j^2 \mathbf{I}$

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

# Mixture of Gaussians

- Covariance models continued:

- ▶  $\Sigma_j = \text{diag}(\{\sigma_{jk}^2\}_{k=1}^d)$

$$\sigma_{jk}^2 = \frac{\sum_{i=1}^n p(j|\mathbf{x}_i, \Theta^{(g)}) (\mathbf{x}_{ik} - \mu_{jk})^2}{\sum_{i=1}^n p(j|\mathbf{x}_i, \Theta^{(g)})}$$

- ▶  $\Sigma_j = \Sigma$

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

- ▶  $\Sigma_j = \text{arbitrary}$

$$\Sigma_j = \frac{\sum_{i=1}^n p(j|\mathbf{x}_i, \Theta^{(g)}) (\mathbf{x}_i - \mu_j) (\mathbf{x}_i - \mu_j)^T}{\sum_{i=1}^n p(j|\mathbf{x}_i, \Theta^{(g)})}$$

# Mixture of Gaussians

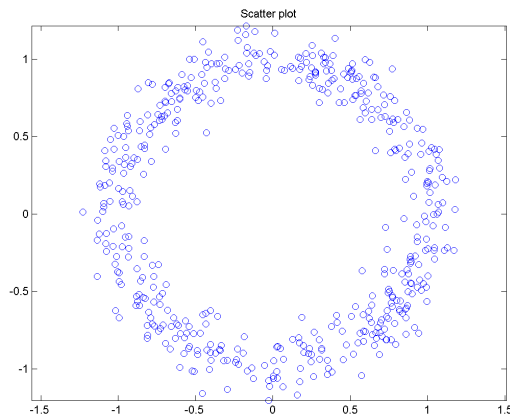
- Summary:
  - ▶ Estimates for  $\alpha_j$ ,  $\mu_j$  and  $\Sigma_j$  perform both expectation and maximization steps simultaneously.
  - ▶ EM iterations proceed by using the current estimates as the initial estimates for the next iteration.
  - ▶ The priors are computed from the proportion of examples 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.

# Mixture of Gaussians

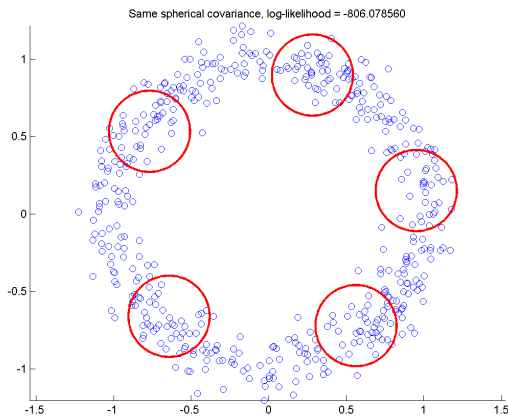
- Questions:
  - ▶ How can we find the number of components in the mixture?
  - ▶ How can we find the initial estimates for  $\Theta$ ?
  - ▶ How do we know when to stop the iterations?
    - Stop if the change in log-likelihood between two iterations is less than a threshold.
    - Or, use a threshold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

# Examples

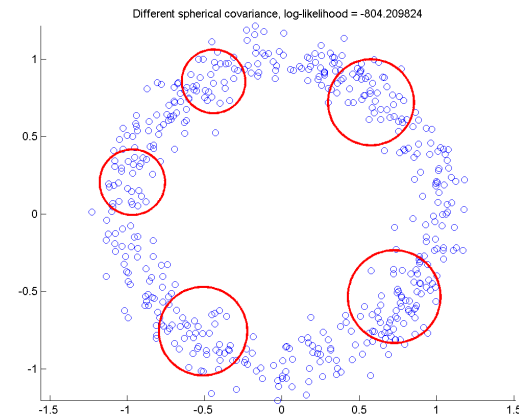
- Mixture of Gaussians examples
- 1-D Bayesian classification examples
- 2-D Bayesian classification examples



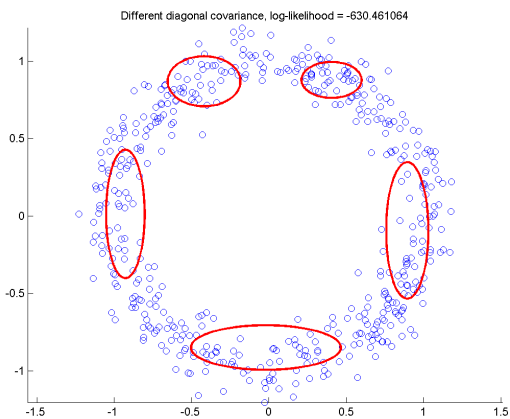
(a) Scatter plot.



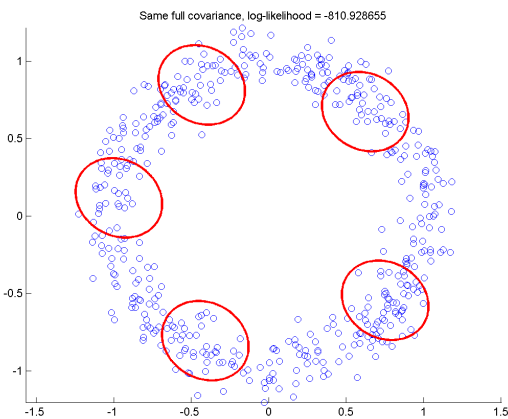
(b) Same spherical covariance, log-likelihood = -806.08.



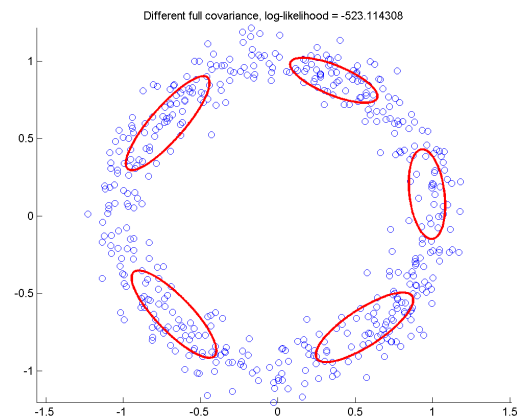
(c) Different spherical covariance, log-likelihood = -804.21.



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

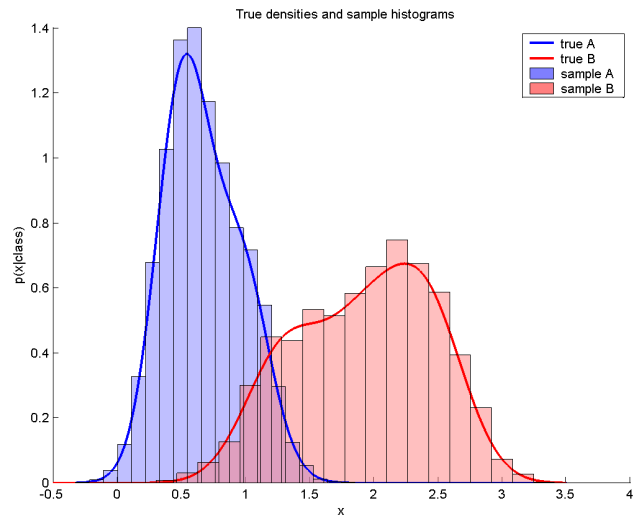


(e) Same arbitrary covariance, log-likelihood = -810.93.

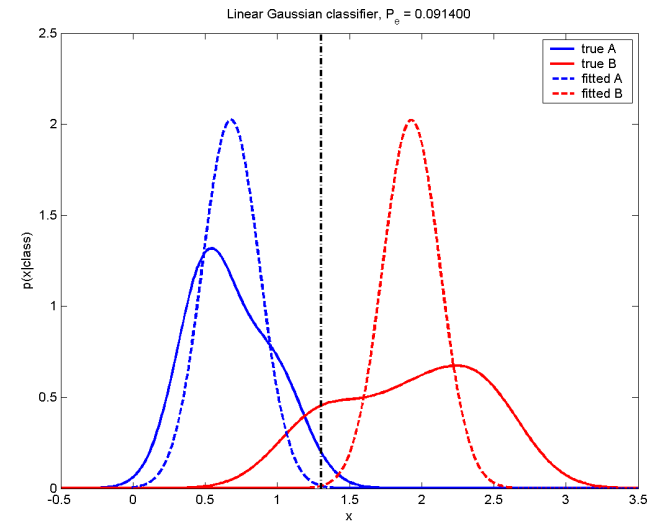


(f) Different arbitrary covariance, log-likelihood = -523.11.

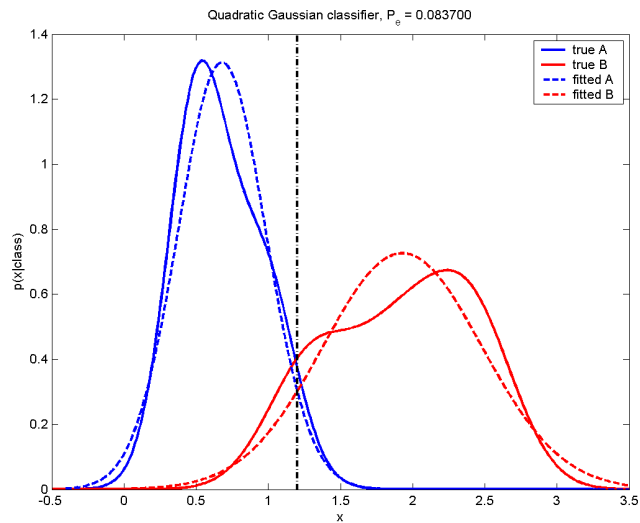
Figure 1: Fitting mixtures of 5 Gaussians to data from a circular distribution.



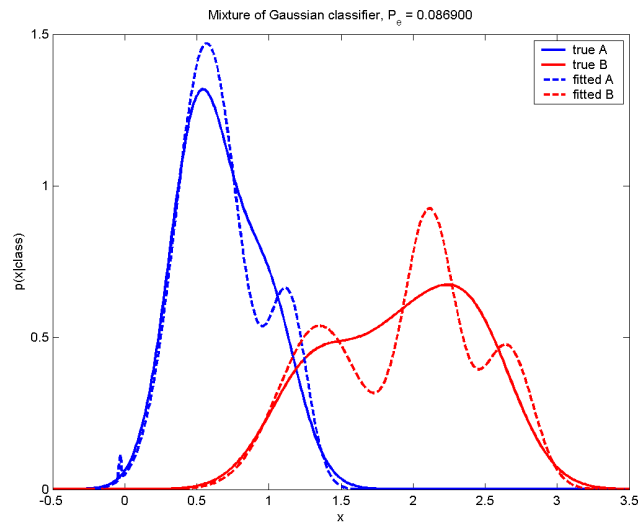
(a) True densities and sample histograms.



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



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

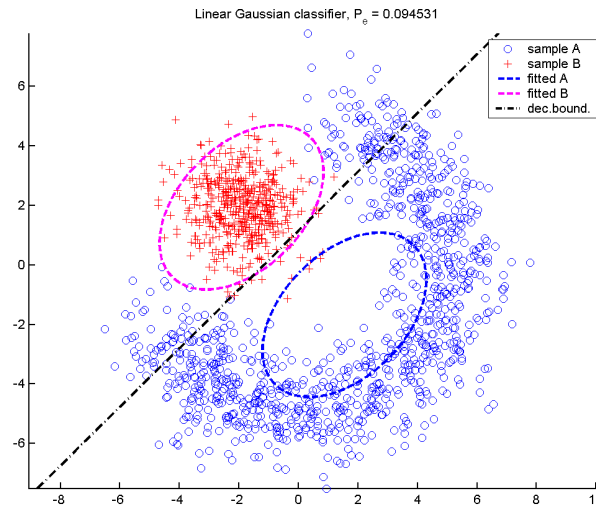


(d) Mixture of Gaussian classifier with  $P_e = 0.0869$ .

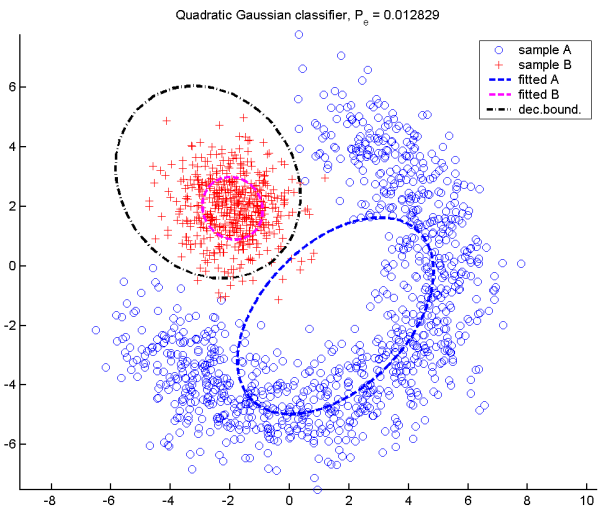
Figure 2: 1-D Bayesian classification examples where the data for each class come from a mixture of three Gaussians. Bayes error is  $P_e = 0.0828$ .



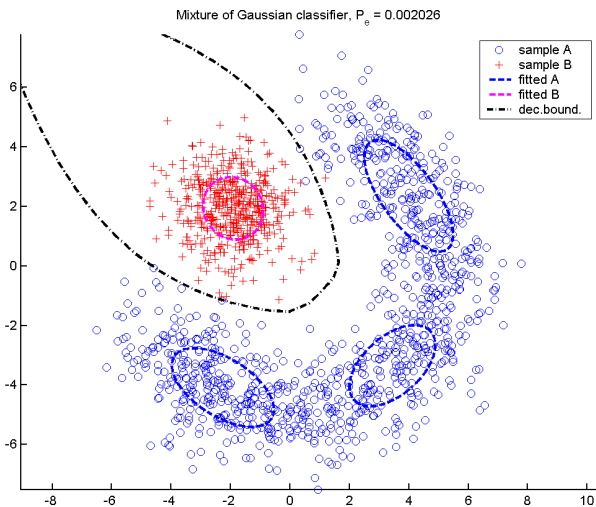
(a) Scatter plot.



(b) Linear Gaussian classifier with  $P_e = 0.094531$ .



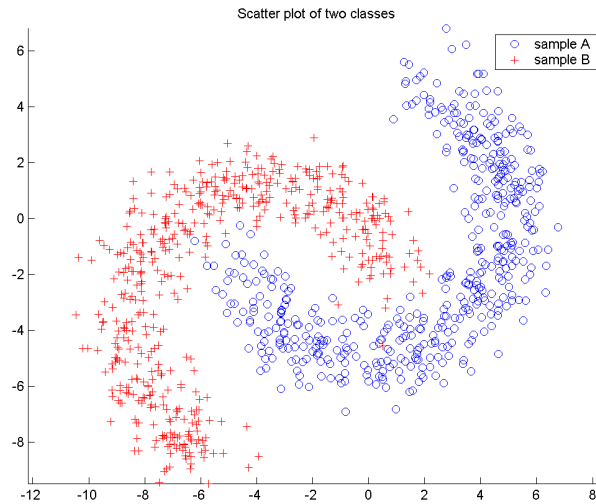
(c) Quadratic Gaussian classifier with  $P_e = 0.012829$ .



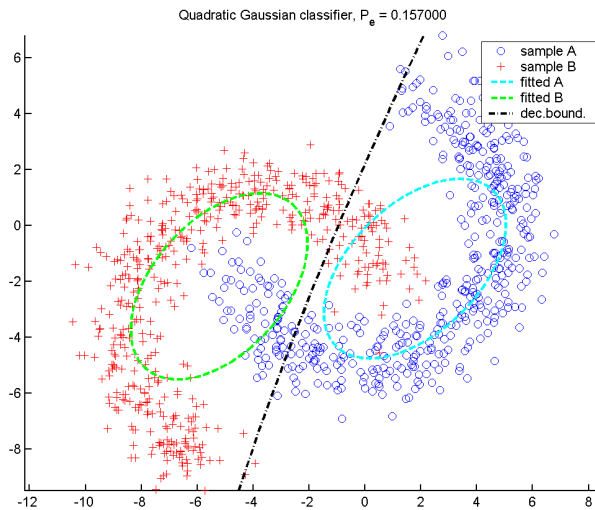
(d) Mixture of Gaussian classifier with  $P_e = 0.002026$ .

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

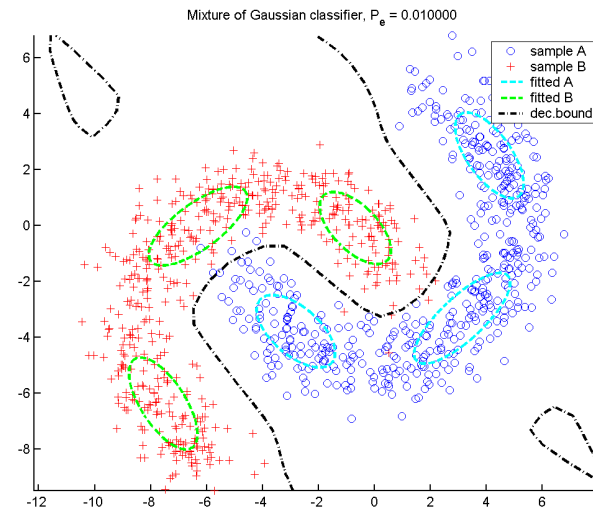




(a) Scatter plot.



(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.