The M step involves maximization of this function with respect to  $\theta$ , keeping  $\theta^{\text{old}}$ , and hence  $\gamma_{nk}$ , fixed. Maximization with respect to  $\pi_k$  can be done in the usual way, with a Lagrange multiplier to enforce the summation constraint  $\sum_k \pi_k = 1$ , giving the familiar result

$$\pi_k = \frac{1}{N} \sum_{n=1}^{N} \gamma_{nk}.$$
 (14.50)

To determine the  $\{\mathbf{w}_k\}$ , we note that the  $Q(\boldsymbol{\theta}, \boldsymbol{\theta}^{\text{old}})$  function comprises a sum over terms indexed by k each of which depends only on one of the vectors  $\mathbf{w}_k$ , so that the different vectors are decoupled in the M step of the EM algorithm. In other words, the different components interact only via the responsibilities, which are fixed during the M step. Note that the M step does not have a closed-form solution and must be solved iteratively using, for instance, the iterative reweighted least squares (IRLS) algorithm. The gradient and the Hessian for the vector  $\mathbf{w}_k$  are given by

$$\nabla_k Q = \sum_{n=1}^N \gamma_{nk} (t_n - y_{nk}) \phi_n$$
 (14.51)

$$\mathbf{H}_{k} = -\nabla_{k}\nabla_{k}Q = \sum_{n=1}^{N} \gamma_{nk} y_{nk} (1 - y_{nk}) \boldsymbol{\phi}_{n} \boldsymbol{\phi}_{n}^{\mathrm{T}}$$
(14.52)

where  $\nabla_k$  denotes the gradient with respect to  $\mathbf{w}_k$ . For fixed  $\gamma_{nk}$ , these are independent of  $\{\mathbf{w}_j\}$  for  $j \neq k$  and so we can solve for each  $\mathbf{w}_k$  separately using the IRLS algorithm. Thus the M-step equations for component k correspond simply to fitting a single logistic regression model to a weighted data set in which data point n carries a weight  $\gamma_{nk}$ . Figure 14.10 shows an example of the mixture of logistic regression models applied to a simple classification problem. The extension of this model to a mixture of softmax models for more than two classes is straightforward.

# 14.5.3 Mixtures of experts

In Section 14.5.1, we considered a mixture of linear regression models, and in Section 14.5.2 we discussed the analogous mixture of linear classifiers. Although these simple mixtures extend the flexibility of linear models to include more complex (e.g., multimodal) predictive distributions, they are still very limited. We can further increase the capability of such models by allowing the mixing coefficients themselves to be functions of the input variable, so that

$$p(\mathbf{t}|\mathbf{x}) = \sum_{k=1}^{K} \pi_k(\mathbf{x}) p_k(\mathbf{t}|\mathbf{x}).$$
 (14.53)

This is known as a *mixture of experts* model (Jacobs *et al.*, 1991) in which the mixing coefficients  $\pi_k(\mathbf{x})$  are known as *gating* functions and the individual component densities  $p_k(\mathbf{t}|\mathbf{x})$  are called *experts*. The notion behind the terminology is that different components can model the distribution in different regions of input space (they

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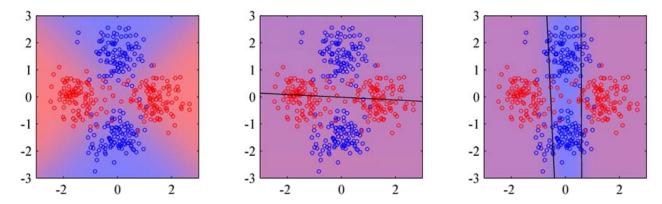


Figure 14.10 Illustration of a mixture of logistic regression models. The left plot shows data points drawn from two classes denoted red and blue, in which the background colour (which varies from pure red to pure blue) denotes the true probability of the class label. The centre plot shows the result of fitting a single logistic regression model using maximum likelihood, in which the background colour denotes the corresponding probability of the class label. Because the colour is a near-uniform purple, we see that the model assigns a probability of around 0.5 to each of the classes over most of input space. The right plot shows the result of fitting a mixture of two logistic regression models, which now gives much higher probability to the correct labels for many of the points in the blue class.

are 'experts' at making predictions in their own regions), and the gating functions determine which components are dominant in which region.

The gating functions  $\pi_k(\mathbf{x})$  must satisfy the usual constraints for mixing coefficients, namely  $0 \leqslant \pi_k(\mathbf{x}) \leqslant 1$  and  $\sum_k \pi_k(\mathbf{x}) = 1$ . They can therefore be represented, for example, by linear softmax models of the form (4.104) and (4.105). If the experts are also linear (regression or classification) models, then the whole model can be fitted efficiently using the EM algorithm, with iterative reweighted least squares being employed in the M step (Jordan and Jacobs, 1994).

Such a model still has significant limitations due to the use of linear models for the gating and expert functions. A much more flexible model is obtained by using a multilevel gating function to give the *hierarchical mixture of experts*, or *HME* model (Jordan and Jacobs, 1994). To understand the structure of this model, imagine a mixture distribution in which each component in the mixture is itself a mixture distribution. For simple unconditional mixtures, this hierarchical mixture is trivially equivalent to a single flat mixture distribution. However, when the mixing coefficients are input dependent, this hierarchical model becomes nontrivial. The HME model can also be viewed as a probabilistic version of *decision trees* discussed in Section 14.4 and can again be trained efficiently by maximum likelihood using an EM algorithm with IRLS in the M step. A Bayesian treatment of the HME has been given by Bishop and Svensén (2003) based on variational inference.

We shall not discuss the HME in detail here. However, it is worth pointing out the close connection with the *mixture density network* discussed in Section 5.6. The principal advantage of the mixtures of experts model is that it can be optimized by EM in which the M step for each mixture component and gating model involves a convex optimization (although the overall optimization is nonconvex). By contrast, the advantage of the mixture density network approach is that the component

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densities and the mixing coefficients share the hidden units of the neural network. Furthermore, in the mixture density network, the splits of the input space are further relaxed compared to the hierarchical mixture of experts in that they are not only soft, and not constrained to be axis aligned, but they can also be nonlinear.

## **Exercises**

- 14.1 (\*\*) www Consider a set models of the form  $p(\mathbf{t}|\mathbf{x}, \mathbf{z}_h, \boldsymbol{\theta}_h, h)$  in which  $\mathbf{x}$  is the input vector,  $\mathbf{t}$  is the target vector, h indexes the different models,  $\mathbf{z}_h$  is a latent variable for model h, and  $\boldsymbol{\theta}_h$  is the set of parameters for model h. Suppose the models have prior probabilities p(h) and that we are given a training set  $\mathbf{X} = \{\mathbf{x}_1, \dots, \mathbf{x}_N\}$  and  $\mathbf{T} = \{\mathbf{t}_1, \dots, \mathbf{t}_N\}$ . Write down the formulae needed to evaluate the predictive distribution  $p(\mathbf{t}|\mathbf{x}, \mathbf{X}, \mathbf{T})$  in which the latent variables and the model index are marginalized out. Use these formulae to highlight the difference between Bayesian averaging of different models and the use of latent variables within a single model.
- **14.2** (\*) The expected sum-of-squares error  $E_{\rm AV}$  for a simple committee model can be defined by (14.10), and the expected error of the committee itself is given by (14.11). Assuming that the individual errors satisfy (14.12) and (14.13), derive the result (14.14).
- **14.3** (\*) www By making use of Jensen's inequality (1.115), for the special case of the convex function  $f(x) = x^2$ , show that the average expected sum-of-squares error  $E_{\text{AV}}$  of the members of a simple committee model, given by (14.10), and the expected error  $E_{\text{COM}}$  of the committee itself, given by (14.11), satisfy

$$E_{\text{COM}} \leqslant E_{\text{AV}}.$$
 (14.54)

- **14.4** ( $\star\star$ ) By making use of Jensen's in equality (1.115), show that the result (14.54) derived in the previous exercise hods for any error function E(y), not just sum-of-squares, provided it is a convex function of y.
- 14.5  $(\star \star)$  www Consider a committee in which we allow unequal weighting of the constituent models, so that

$$y_{\text{COM}}(\mathbf{x}) = \sum_{m=1}^{M} \alpha_m y_m(\mathbf{x}). \tag{14.55}$$

In order to ensure that the predictions  $y_{\text{COM}}(\mathbf{x})$  remain within sensible limits, suppose that we require that they be bounded at each value of  $\mathbf{x}$  by the minimum and maximum values given by any of the members of the committee, so that

$$y_{\min}(\mathbf{x}) \leqslant y_{\text{COM}}(\mathbf{x}) \leqslant y_{\max}(\mathbf{x}).$$
 (14.56)

Show that a necessary and sufficient condition for this constraint is that the coefficients  $\alpha_m$  satisfy

$$\alpha_m \geqslant 0, \qquad \sum_{m=1}^{M} \alpha_m = 1. \tag{14.57}$$