### 272 8. Model Inference and Averaging

that is, the prior probability mass function is proportional to  $\prod_{\ell=1}^{L} w_{\ell}^{a-1}$ . Then the posterior density of w is

$$w \sim \mathrm{Di}_L(a1 + N\hat{w}),\tag{8.33}$$

where N is the sample size. Letting  $a \to 0$  to obtain a noninformative prior gives

$$w \sim \mathrm{Di}_L(N\hat{w}). \tag{8.34}$$

Now the bootstrap distribution, obtained by sampling with replacement from the data, can be expressed as sampling the category proportions from a multinomial distribution. Specifically,

$$N\hat{w}^* \sim \operatorname{Mult}(N, \hat{w}), \tag{8.35}$$

where  $\operatorname{Mult}(N, \hat{w})$  denotes a multinomial distribution, having probability mass function  $\binom{N}{N\hat{w}_1^*, \dots, N\hat{w}_\ell^*} \prod \hat{w}_\ell^{N\hat{w}_\ell^*}$ . This distribution is similar to the posterior distribution above, having the same support, same mean, and nearly the same covariance matrix. Hence the bootstrap distribution of  $S(\hat{w}^*)$  will closely approximate the posterior distribution of S(w).

In this sense, the bootstrap distribution represents an (approximate) nonparametric, noninformative posterior distribution for our parameter. But this bootstrap distribution is obtained painlessly—without having to formally specify a prior and without having to sample from the posterior distribution. Hence we might think of the bootstrap distribution as a "poor man's" Bayes posterior. By perturbing the data, the bootstrap approximates the Bayesian effect of perturbing the parameters, and is typically much simpler to carry out.

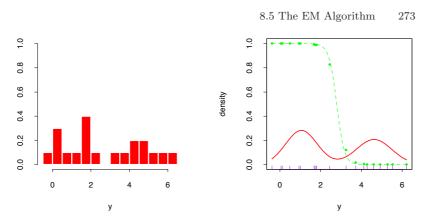
# 8.5 The EM Algorithm

The EM algorithm is a popular tool for simplifying difficult maximum likelihood problems. We first describe it in the context of a simple mixture model.

### 8.5.1 Two-Component Mixture Model

In this section we describe a simple mixture model for density estimation, and the associated EM algorithm for carrying out maximum likelihood estimation. This has a natural connection to Gibbs sampling methods for Bayesian inference. Mixture models are discussed and demonstrated in several other parts of the book, in particular Sections 6.8, 12.7 and 13.2.3.

The left panel of Figure 8.5 shows a histogram of the 20 fictitious data points in Table 8.1.



**FIGURE 8.5.** Mixture example. (Left panel:) Histogram of data. (Right panel:) Maximum likelihood fit of Gaussian densities (solid red) and responsibility (dotted green) of the left component density for observation y, as a function of y.

**TABLE 8.1.** Twenty fictitious data points used in the two-component mixture example in Figure 8.5.

-0.39	0.12	0.94	1.67	1.76	2.44	3.72	4.28	4.92	5.53
0.06	0.48	1.01	1.68	1.80	3.25	4.12	4.60	5.28	6.22

We would like to model the density of the data points, and due to the apparent bi-modality, a Gaussian distribution would not be appropriate. There seems to be two separate underlying regimes, so instead we model Y as a mixture of two normal distributions:

$$Y_1 \sim N(\mu_1, \sigma_1^2), 
 Y_2 \sim N(\mu_2, \sigma_2^2), 
 Y = (1 - \Delta) \cdot Y_1 + \Delta \cdot Y_2,$$
(8.36)

where  $\Delta \in \{0, 1\}$  with  $Pr(\Delta = 1) = \pi$ . This generative representation is explicit: generate a  $\Delta \in \{0, 1\}$  with probability  $\pi$ , and then depending on the outcome, deliver either  $Y_1$  or  $Y_2$ . Let  $\phi_{\theta}(x)$  denote the normal density with parameters  $\theta = (\mu, \sigma^2)$ . Then the density of Y is

$$g_Y(y) = (1 - \pi)\phi_{\theta_1}(y) + \pi\phi_{\theta_2}(y).$$
(8.37)

Now suppose we wish to fit this model to the data in Figure 8.5 by maximum likelihood. The parameters are

$$\theta = (\pi, \theta_1, \theta_2) = (\pi, \mu_1, \sigma_1^2, \mu_2, \sigma_2^2).$$
(8.38)

The log-likelihood based on the N training cases is

$$\ell(\theta; \mathbf{Z}) = \sum_{i=1}^{N} \log[(1-\pi)\phi_{\theta_1}(y_i) + \pi \phi_{\theta_2}(y_i)].$$
(8.39)

### 274 8. Model Inference and Averaging

Direct maximization of  $\ell(\theta; \mathbf{Z})$  is quite difficult numerically, because of the sum of terms inside the logarithm. There is, however, a simpler approach. We consider unobserved latent variables  $\Delta_i$  taking values 0 or 1 as in (8.36): if  $\Delta_i = 1$  then  $Y_i$  comes from model 2, otherwise it comes from model 1. Suppose we knew the values of the  $\Delta_i$ 's. Then the log-likelihood would be

$$\ell_{0}(\theta; \mathbf{Z}, \mathbf{\Delta}) = \sum_{i=1}^{N} \left[ (1 - \Delta_{i}) \log \phi_{\theta_{1}}(y_{i}) + \Delta_{i} \log \phi_{\theta_{2}}(y_{i}) \right] \\ + \sum_{i=1}^{N} \left[ (1 - \Delta_{i}) \log(1 - \pi) + \Delta_{i} \log \pi \right], \quad (8.40)$$

and the maximum likelihood estimates of  $\mu_1$  and  $\sigma_1^2$  would be the sample mean and variance for those data with  $\Delta_i = 0$ , and similarly those for  $\mu_2$ and  $\sigma_2^2$  would be the sample mean and variance of the data with  $\Delta_i = 1$ . The estimate of  $\pi$  would be the proportion of  $\Delta_i = 1$ .

Since the values of the  $\Delta_i$ 's are actually unknown, we proceed in an iterative fashion, substituting for each  $\Delta_i$  in (8.40) its expected value

$$\gamma_i(\theta) = \mathcal{E}(\Delta_i | \theta, \mathbf{Z}) = \Pr(\Delta_i = 1 | \theta, \mathbf{Z}), \tag{8.41}$$

also called the *responsibility* of model 2 for observation *i*. We use a procedure called the EM algorithm, given in Algorithm 8.1 for the special case of Gaussian mixtures. In the *expectation* step, we do a soft assignment of each observation to each model: the current estimates of the parameters are used to assign responsibilities according to the relative density of the training points under each model. In the *maximization* step, these responsibilities are used in weighted maximum-likelihood fits to update the estimates of the parameters.

A good way to construct initial guesses for  $\hat{\mu}_1$  and  $\hat{\mu}_2$  is simply to choose two of the  $y_i$  at random. Both  $\hat{\sigma}_1^2$  and  $\hat{\sigma}_2^2$  can be set equal to the overall sample variance  $\sum_{i=1}^{N} (y_i - \bar{y})^2 / N$ . The mixing proportion  $\hat{\pi}$  can be started at the value 0.5.

Note that the actual maximizer of the likelihood occurs when we put a spike of infinite height at any one data point, that is,  $\hat{\mu}_1 = y_i$  for some i and  $\hat{\sigma}_1^2 = 0$ . This gives infinite likelihood, but is not a useful solution. Hence we are actually looking for a good local maximum of the likelihood, one for which  $\hat{\sigma}_1^2, \hat{\sigma}_2^2 > 0$ . To further complicate matters, there can be more than one local maximum having  $\hat{\sigma}_1^2, \hat{\sigma}_2^2 > 0$ . In our example, we ran the EM algorithm with a number of different initial guesses for the parameters, all having  $\hat{\sigma}_k^2 > 0.5$ , and chose the run that gave us the highest maximized likelihood. Figure 8.6 shows the progress of the EM algorithm in maximizing the log-likelihood. Table 8.2 shows  $\hat{\pi} = \sum_i \hat{\gamma}_i / N$ , the maximum likelihood estimate of the proportion of observations in class 2, at selected iterations of the EM procedure.

Algorithm 8.1 EM Algorithm for Two-component Gaussian Mixture.

- 1. Take initial guesses for the parameters  $\hat{\mu}_1, \hat{\sigma}_1^2, \hat{\mu}_2, \hat{\sigma}_2^2, \hat{\pi}$  (see text).
- 2. Expectation Step: compute the responsibilities

$$\hat{\gamma}_i = \frac{\hat{\pi}\phi_{\hat{\theta}_2}(y_i)}{(1-\hat{\pi})\phi_{\hat{\theta}_1}(y_i) + \hat{\pi}\phi_{\hat{\theta}_2}(y_i)}, \ i = 1, 2, \dots, N.$$
(8.42)

3. Maximization Step: compute the weighted means and variances:

$$\hat{\mu}_{1} = \frac{\sum_{i=1}^{N} (1 - \hat{\gamma}_{i}) y_{i}}{\sum_{i=1}^{N} (1 - \hat{\gamma}_{i})}, \qquad \hat{\sigma}_{1}^{2} = \frac{\sum_{i=1}^{N} (1 - \hat{\gamma}_{i}) (y_{i} - \hat{\mu}_{1})^{2}}{\sum_{i=1}^{N} (1 - \hat{\gamma}_{i})},$$
$$\hat{\mu}_{2} = \frac{\sum_{i=1}^{N} \hat{\gamma}_{i} y_{i}}{\sum_{i=1}^{N} \hat{\gamma}_{i}}, \qquad \hat{\sigma}_{2}^{2} = \frac{\sum_{i=1}^{N} \hat{\gamma}_{i} (y_{i} - \hat{\mu}_{2})^{2}}{\sum_{i=1}^{N} \hat{\gamma}_{i}},$$

and the mixing probability  $\hat{\pi} = \sum_{i=1}^{N} \hat{\gamma}_i / N$ .

4. Iterate steps 2 and 3 until convergence.

**TABLE 8.2.** Selected iterations of the EM algorithm for mixture example.

Iteration	$\hat{\pi}$
1	0.485
5	0.493
10	0.523
15	0.544
20	0.546

The final maximum likelihood estimates are

$$\hat{\mu}_1 = 4.62,$$
  $\hat{\sigma}_1^2 = 0.87,$   
 $\hat{\mu}_2 = 1.06,$   $\hat{\sigma}_2^2 = 0.77,$   
 $\hat{\pi} = 0.546.$ 

The right panel of Figure 8.5 shows the estimated Gaussian mixture density from this procedure (solid red curve), along with the responsibilities (dotted green curve). Note that mixtures are also useful for supervised learning; in Section 6.7 we show how the Gaussian mixture model leads to a version of radial basis functions.

276 8. Model Inference and Averaging

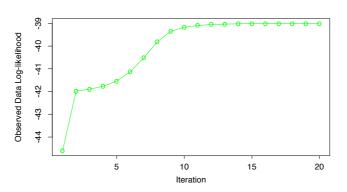


FIGURE 8.6. EM algorithm: observed data log-likelihood as a function of the iteration number.

## 8.5.2 The EM Algorithm in General



The above procedure is an example of the EM (or Baum–Welch) algorithm for maximizing likelihoods in certain classes of problems. These problems are ones for which maximization of the likelihood is difficult, but made easier by enlarging the sample with latent (unobserved) data. This is called *data augmentation*. Here the latent data are the model memberships  $\Delta_i$ . In other problems, the latent data are actual data that should have been observed but are missing.

Algorithm 8.2 gives the general formulation of the EM algorithm. Our observed data is  $\mathbf{Z}$ , having log-likelihood  $\ell(\theta; \mathbf{Z})$  depending on parameters  $\theta$ . The latent or missing data is  $\mathbf{Z}^m$ , so that the complete data is  $\mathbf{T} = (\mathbf{Z}, \mathbf{Z}^m)$  with log-likelihood  $\ell_0(\theta; \mathbf{T})$ ,  $\ell_0$  based on the complete density. In the mixture problem  $(\mathbf{Z}, \mathbf{Z}^m) = (\mathbf{y}, \Delta)$ , and  $\ell_0(\theta; \mathbf{T})$  is given in (8.40).

In our mixture example,  $E(\ell_0(\theta'; \mathbf{T}) | \mathbf{Z}, \hat{\theta}^{(j)})$  is simply (8.40) with the  $\Delta_i$  replaced by the responsibilities  $\hat{\gamma}_i(\hat{\theta})$ , and the maximizers in step 3 are just weighted means and variances.

We now give an explanation of why the EM algorithm works in general. Since

$$\Pr(\mathbf{Z}^m | \mathbf{Z}, \theta') = \frac{\Pr(\mathbf{Z}^m, \mathbf{Z} | \theta')}{\Pr(\mathbf{Z} | \theta')},$$
(8.44)

we can write

$$\Pr(\mathbf{Z}|\theta') = \frac{\Pr(\mathbf{T}|\theta')}{\Pr(\mathbf{Z}^m|\mathbf{Z},\theta')}.$$
(8.45)

In terms of log-likelihoods, we have  $\ell(\theta'; \mathbf{Z}) = \ell_0(\theta'; \mathbf{T}) - \ell_1(\theta'; \mathbf{Z}^m | \mathbf{Z})$ , where  $\ell_1$  is based on the conditional density  $\Pr(\mathbf{Z}^m | \mathbf{Z}, \theta')$ . Taking conditional expectations with respect to the distribution of  $\mathbf{T} | \mathbf{Z}$  governed by parameter  $\theta$  gives

$$\ell(\theta'; \mathbf{Z}) = \mathrm{E}[\ell_0(\theta'; \mathbf{T}) | \mathbf{Z}, \theta] - \mathrm{E}[\ell_1(\theta'; \mathbf{Z}^m | \mathbf{Z}) | \mathbf{Z}, \theta]$$