

## 8 Notes on EM Algorithm

### 8.1 EM Algorithm for Multinomial & Mixture Data

**General Example 1.** Suppose that for fixed integers  $1 \leq K < C$ , cell-counts  $\mathbf{X} = (X_1, \dots, X_K)$  are observed, and cell-counts  $\mathbf{Y} = (Y_{K+1}, \dots, Y_C)$  *cannot* be observed, where

$$\left( X_1, \dots, X_K, Y_{K+1}, \dots, Y_C \right) \sim \text{Multinomial}(n, p_j(\vartheta), j = 1, \dots, C)$$

Here  $\vartheta$  is an unknown parameter of dimension  $d \leq K$ , and the functions  $p_j(\vartheta)$  which share  $\vartheta$  as a parameter are sufficiently smooth. Also denote

$$X_{K+1} = n - X_1 - \dots - X_K = \sum_{j=K+1}^C Y_j$$

For notational convenience, define

$$q_K(\vartheta) = 1 - p_1(\vartheta) - \dots - p_K(\vartheta)$$

In this setting, we express the conditional joint density of  $\mathbf{Y}$  given  $\mathbf{X}$  by

$$f_{\mathbf{Y}|\mathbf{X}}(\mathbf{y}|\mathbf{X}, \vartheta) = \exp\left(\sum_{j=K+1}^C y_j \log\left(\frac{p_j(\vartheta)}{q_K(\vartheta)}\right)\right) \cdot \binom{X_{K+1}}{Y_{K+1}, \dots, Y_C}$$

It follows that the *E-step* of the EM algorithm replaces  $E_{\vartheta_1}(\log f_{\mathbf{Y}|\mathbf{X}}(\mathbf{Y}|\mathbf{X}, \vartheta))$  by

$$X_{K+1} \sum_{j=K+1}^C \frac{p_j(\vartheta_1)}{q_K(\vartheta_1)} \log\left(\frac{p_j(\vartheta)}{q_K(\vartheta)}\right) + \log\left(\binom{X_{K+1}}{Y_{K+1}, \dots, Y_C}\right)$$

or equivalently, replaces  $Y_j$  by  $X_{K+1} \cdot p_j(\vartheta_1)/q_K(\vartheta_1)$  for  $j = K+1, \dots, C$ .

To confirm that the definition of log-likelihood and conditional log-likelihood terms as above, without multinomial coefficients, is legitimate, we observe that the property needed in the proof of log-likelihood improvement for EM iterations holds, that is,

$$E_{\vartheta}(\log L_{\mathbf{Y}|\mathbf{X}}(n_{C+1}, \dots, n_K | \mathbf{X}, \vartheta) - \log L_{\mathbf{Y}|\mathbf{X}}(n_{C+1}, \dots, n_K | \mathbf{X}, \vartheta_1)) \geq 0$$

or equivalently, for all  $\vartheta, \vartheta_1$ ,

$$\sum_{j=C+1}^K \frac{p_j(\vartheta)}{q(\vartheta)} \log \frac{p_j(\vartheta) q(\vartheta_1)}{q(\vartheta) p_j(\vartheta_1)} \geq 0$$

But this is a standard, discrete version of the famous ‘Information Inequality’ proved more generally in the form  $\int f(x) \log(f(x)/g(x)) d\nu(x) \geq 0$  for probability densities with respect to a measure  $\nu$ , using Jensen’s Inequality.

In the multinomial setting, we express the conditional likelihood for  $\mathbf{Y}$  given  $\mathbf{X}$  by

$$L_{\mathbf{Y}|\mathbf{X}}(n_{C+1}, \dots, n_K | \mathbf{X}, \vartheta) = \exp\left(\sum_{j=C+1}^K n_j \log\left(\frac{p_j(\vartheta)}{q(\vartheta)}\right)\right)$$

The *E-step* of the EM algorithm replaces this conditional log-likelihood, when the current parameter-iterate is  $\vartheta_1$ , by

$$E_{\vartheta_1}\left(\log L_{\mathbf{Y}|\mathbf{X}}(\mathbf{Y}|\mathbf{X}, \vartheta) \mid \mathbf{X}\right) = n^* \sum_{j=C+1}^K \frac{p_j(\vartheta_1)}{q(\vartheta_1)} \log\left(\frac{p_j(\vartheta)}{q(\vartheta)}\right)$$

This expression is also equal to  $\log L_{\mathbf{Y}|\mathbf{X}}(n_{C+1}^*, \dots, n_K^* | \mathbf{X}, \vartheta)$ , where  $n_j^* = E_{\vartheta_1}(n_j^* | \mathbf{X}) = n^* \cdot p_j(\vartheta_1)/q(\vartheta_1)$  for  $j = C+1, \dots, K$ .

Thus we have the following comparison between maximization approaches. First, the complete-data likelihood to maximize, if  $\mathbf{Y}$  could also be observed, would be

$$\sum_{j=1}^K X_j \log p_j(\vartheta) + X_{K+1} \log q_K(\vartheta) + \sum_{j=K+1}^C Y_j \log \frac{p_j(\vartheta)}{q_K(\vartheta)}$$

while the crude marginal-observed-data likelihood to maximize is

$$\sum_{j=1}^K X_j \log p_j(\vartheta) + X_{K+1} \log q_K(\vartheta)$$

On the other hand, the *M-step* of the EM algorithm, after replacement of the unobservable  $Y_j$  values in the complete-data likelihood by their *E-step* imputed values, is

$$\sum_{j=1}^K X_j \log p_j(\vartheta) + X_{K+1} \log q_K(\vartheta) + X_{K+1} \sum_{j=K+1}^C \frac{p_j(\vartheta_1)}{q_K(\vartheta_1)} \log \frac{p_j(\vartheta)}{q_K(\vartheta)}$$

$$= \sum_{j=1}^K X_j \log p_j(\vartheta) + X_{K+1} \sum_{j=K+1}^C \frac{p_j(\vartheta_1)}{q_K(\vartheta_1)} \log p_j(\vartheta)$$

Note that the M-step involves a step of maximizing the complete-data likelihood using imputed data for the  $Y_j$ 's, which will be very easy in some problems.

*A key aspect of the usefulness of the EM algorithm in multinomial missing data problems is that no sums of terms  $p_j(\vartheta)$  appear inside the logarithms arising in the maximization-step. Especially in so-called log-linear contingency-table models with some missing cell-counts, where the  $p_j(\vartheta)$  have some multiplicative structure, this is very useful !*

#### SPECIAL EXAMPLE FROM THE ORIGINAL EM PAPER

This example fits into the structure of the general multinomial example, with scalar unknown parameter  $\vartheta = \pi$ ,  $K = 3$ ,  $C = 5$ , and

$$p_1(\pi) = p_2(\pi) = \frac{1 - \pi}{4}, \quad p_3(\pi) = p_4(\pi) = \frac{\pi}{4}, \quad p_5(\pi) = \frac{1}{2}$$

The cell-counts given as data in Dempster, Laird & Rubin (1978) are:

$(X_1, X_2, X_3, X_4) = (18, 20, 34, 125)$ . Appealing to the formulas above, we find that the complete-data M-step involves maximizing  $\sum_{j=1}^3 X_j \log p_j(\pi) + \sum_{j=4}^5 Y_j \log p_j(\pi)$ . In this particular problem, we are equivalently maximizing  $(X_3 + Y_4) \log(\pi/4) + (X_1 + X_2) \log((1 - \pi)/4)$ , which leads to

$$\hat{\pi} = (X_3 + Y_4)/(n - Y_5)$$

Substituting the E-step imputed valued for the  $Y_j$  gives the EM iteration explicitly, starting from initial guess  $\pi_1$ , as:

$$\begin{aligned} \pi_2 &= \left( X_3 + X_4 \cdot \frac{\pi_1/4}{1/2 + \pi_1/4} \right) / \left( n - X_4 \cdot \frac{1/2}{1/2 + \pi_1/4} \right) \\ &= \frac{34 + 125 \cdot \frac{\pi_1}{2 + \pi_1}}{197 - 125 \cdot \frac{2}{2 + \pi_1}} = \frac{68 + 159 \pi_1}{144 + 197 \pi_1} \end{aligned}$$

In this little example, EM iterates the mapping  $h(\pi) \equiv (68 + 159\pi)/(144 + 197\pi)$  to find the fixed-point. (The unique fixed-point  $\pi = 0.6268$  solves  $h(\pi) = \pi$ , which is a quadratic equation.) The Quasi-Newton optimization of the marginal likelihood is messier but, using a modern computer, quicker and more reliable.

```

> optimize(function(x) 38*log(1-x)+34*log(x)+125*log(x+2),
  c(.01,.99), max=T)$max
[1] 0.6268036
> h <- function(x) (159 * x + 68)/(197 * x + 144)
  x<- .5; for (i in 1:6) {x <- h(x)  cat(round(x,5)," \n")}
0.60825
0.62432
0.62649
0.62678
0.62682    ### converged to 5 places after 5 iterations

```

**General Example 2.** Consider ‘mixture’ data  $X_i$  which are *iid* continuously distributed *rv*’s with density

$$f_X(x) = pe^{-x} + \lambda(1-p)e^{-\lambda x}, \quad x > 0$$

where  $\vartheta = (p, \lambda) \in (0, 1) \times [0, \infty)$  is the unknown parameter. These r.v.’s are of *mixture* type because they have the same density as random variables

$$X_i = \epsilon_i U_i + (1 - \epsilon_i) V_i \quad U_i \sim \text{Expon}(1), \quad V_i \sim \text{Expon}(\lambda)$$

where  $\epsilon_i \sim \text{Binom}(1, p)$  is independent of  $(U_i, V_i)$ . The marginal density for the observed variables is  $f_X$ , but the problem would be much simpler to analyze with the ‘complete’ data  $(X_i, \epsilon_i)$ ,  $i = 1, \dots, n$ . Now the *E-step* of the EM algorithm based on observing only  $\mathbf{X} = (X_i, i = 1, \dots, n)$  consists of calculating

$$E_{\vartheta_1}(\epsilon | X) = \frac{p_1 e^{-X}}{p_1 e^{-X} + \lambda_1 (1 - p_1) e^{-\lambda_1 X}} = \epsilon^*(X, \vartheta_1) = \epsilon^*$$

and then substituting to obtain

$$\begin{aligned}
E_{\vartheta_1} \log p_{\epsilon|X}(\epsilon | X, \vartheta) &= \epsilon^* \log \left( \frac{p e^{-X}}{p e^{-X} + \lambda (1 - p) e^{-\lambda X}} \right) \\
&+ (1 - \epsilon^*) \log \left( \frac{\lambda (1 - p) e^{-\lambda X}}{p e^{-X} + \lambda (1 - p) e^{-\lambda X}} \right)
\end{aligned}$$

As a result, starting from initial guess  $\vartheta_1 = (\lambda_1, p_1)$ , the *M-step* of the EM algorithm is to maximize the ‘complete-data log-likelihood’ for the data

$(X_i, \epsilon^*(X_i, \vartheta_1), i = 1, \dots, n)$ , which is given simply in terms of

$$m^* = \sum_{i=1}^n \epsilon^*(X_i, \vartheta_1) \quad , \quad \bar{U} = (m^*)^{-1} \sum_{i=1}^n X_i \epsilon^*(X_i, \vartheta_1)$$

and

$$\bar{V} = (n - m^*)^{-1} \sum_{i=1}^n X_i (1 - \epsilon^*(X_i, \vartheta_1))$$

as

$$m^* (\log p - \bar{U}) + (n - m^*) (\log(\lambda(1 - p)) - \lambda \bar{V})$$

Thus the *M-step* is given in closed form by maximizing the last expression in  $(\lambda, p)$  to obtain

$$p_2 = m^*/n \quad , \quad \lambda_2 = 1/\bar{V}$$

In summary, the entire EM iteration-step in this example, starting from initial guess  $\vartheta_1 = (\lambda_1, p_1)$ , is given in closed form by:

$$p_2 = \frac{1}{n} \sum_{i=1}^n \frac{p_1 e^{-X_i}}{p_1 e^{-X_i} + \lambda_1 (1 - p_1) e^{-\lambda_1 X_i}}$$

$$1/\lambda_2 = \frac{1}{n(1 - p_2)} \sum_{i=1}^n \frac{(1 - p_1) \lambda_1 X_i e^{-\lambda_1 X_i}}{p_1 e^{-X_i} + \lambda_1 (1 - p_1) e^{-\lambda_1 X_i}}$$

We implement this, and evaluate the results in a little simulated dataset, as follows.

```
> EMiter
function(thet, Xvec)
{
## On input, thet is the vector consisting of old values of
##   p, lambda in General Example 2 of Notes, and Xvec is
##   the observed data vector. The output is the new theta.
  frac <- 1/(1 + (1/thet[1] - 1) * thet[2] * exp((
    1 - thet[2]) * Xvec))
  pnew <- mean(frac)
  lamnew <- (1 - pnew)/mean(Xvec * (1 - frac))
}
```

```

list(thet = c(pnew, lamnew), logL = sum(log(pnew * exp(
  - Xvec) + (1 - pnew) * lamnew * exp( - lamnew * Xvec))))
}

> epsv <- rbinom(10000, 1, .6)
  Xv <- rexp(10000)/exp(.3*(1-epsv))
> round(c(mean(epsv), .4/exp(.3)+.6, mean(Xv)),5)
[1] 0.60050 0.89633 0.89459

> theta <- c(.5,1.5)
## Initial log-likelihood
> sum(log(.5 * exp( - Xv) + .5*1.5*exp(-1.5*Xv))) ## = -8908.9
## Log-likelihood at true values:
> sum(log(.6 * exp( - Xv) + .4*exp(.3-exp(.3)*Xv)))## -8883.5

> unlist(EMiter(theta,Xv)) ## values after one EM iteration
  thet1   thet2   logL
0.5069983 1.413503 -8893.04

> for(i in 1:100) {
  tmpitr <- EMiter(theta,Xv)
  theta <<- tmpitr$thet
  if(i %% 5 ==0) cat(round(unlist(tmpitr),5),"\n") }
0.51477 1.30256 -8883.91956
0.51623 1.28736 -8883.74519
0.51689 1.28652 -8883.74114
0.51749 1.28686 -8883.73824
0.51807 1.28728 -8883.73536
0.51866 1.28772 -8883.73248
0.51925 1.28815 -8883.72962
0.51983 1.28859 -8883.72677
0.52042 1.28903 -8883.72393
0.521 1.28947 -8883.72111
0.52158 1.2899 -8883.71829
0.52216 1.29034 -8883.71549
0.52274 1.29078 -8883.7127
0.52331 1.29122 -8883.70992

```

```

0.52389 1.29165 -8883.70716
0.52446 1.29209 -8883.7044
0.52504 1.29253 -8883.70166
0.52561 1.29297 -8883.69893
0.52618 1.29341 -8883.69622
0.52675 1.29385 -8883.69351

> for(i in 1:100) theta <- EMiter(theta,Xv)$thet
  unlist(EMiter(theta,Xv))
    thet1    thet2    logL
0.5379237 1.302709 -8883.642
> for(i in 1:100) theta <- EMiter(theta,Xv)$thet
  unlist(EMiter(theta,Xv))
>   thet1    thet2    logL
0.5483496 1.311421 -8883.596
## Convergence is painfully slow !!!

> nlminb(c(0.5, 1.5), function(x) - sum(log(x[1] * exp(- Xv) +
  (1 - x[1]) * x[2] * exp(- x[2] * Xv))), lower
  = c(0.01, 0.1), upper = c(0.99, 10))[1:4]
$parameters:
[1] 0.6308609 1.3997779

$objective:
[1] 8883.386

$message:
[1] "RELATIVE FUNCTION CONVERGENCE"

$grad.norm:
[1] 0.001797887

```

Note the very slow convergence of the EM algorithm implemented and tested here. The maximized  $\log Lik$  must be larger than  $-8883.5$ , since that is the value at the true parameters ( $p = .6$ ,  $\lambda = e^{-3}$ ), but from the not-too-awful starting values  $p_1 = .5$ ,  $\lambda = 1.5$ , it took more than 300 EM iterations to get there ! As can be seen from the final converged maximization via **nlminb**, the final maximized  $\log Lik$  is  $-8883.39$ .