

# Monte Carlo Likelihood Approximation

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## 1 Monte Carlo Likelihood Approximation

Let  $f_{\theta}(x, y)$  be the complete data density for a missing data model, the missing data being  $x$  and the observed data being  $y$ . Suppose we have observed data  $y_1, \dots, y_n$  which are independent and identically distributed (IID) and simulations  $x_1, \dots, x_m$  which are IID from a known importance sampling distribution with density  $h$ .

The (observed data) log likelihood for this model is

$$l_n(\theta) = \sum_{j=1}^n \log f_{\theta}(y_j) \tag{1}$$

where

$$f_{\theta}(y) = \int f_{\theta}(x, y) dx$$

is the marginal for  $y$ .

The Monte Carlo likelihood approximation for (1) is

$$l_{m,n}(\theta) = \sum_{j=1}^n \log f_{m,\theta}(y_j) \tag{2a}$$

where

$$f_{\theta,m}(y) = \frac{1}{m} \sum_{i=1}^m \frac{f_{\theta}(x_i, y)}{h(x_i)}. \quad (2b)$$

The maximizer  $\hat{\theta}_{m,n}$  of (2a) is the Monte Carlo (approximation to the) MLE (the MCMLLE).

Derivatives of (2a) are, of course,

$$\nabla^k l_{m,n}(\theta) = \sum_{j=1}^n \nabla^k \log f_{m,\theta}(y_j)$$

where  $\nabla$  denotes differentiation with respect to  $\theta$ , and derivatives of (2b) are

$$\nabla f_{\theta,m}(y) = \sum_{i=1}^m \nabla \log f_{\theta}(x_i, y) \cdot v_{\theta}(x_i, y), \quad (3a)$$

where

$$v_{\theta}(x, y) = \frac{\frac{f_{\theta}(x, y)}{h(x)}}{\sum_{i=1}^m \frac{f_{\theta}(x_i, y)}{h(x_i)}}, \quad (3b)$$

and

$$\begin{aligned} \nabla^2 \log f_{\theta,m}(y) &= \sum_{i=1}^m \nabla^2 \log f_{\theta}(x_i, y) \cdot v_{\theta}(x_i, y) \\ &+ \sum_{i=1}^m (\nabla \log f_{\theta}(x_i, y)) (\nabla \log f_{\theta}(x_i, y))^T \cdot v_{\theta}(x_i, y) \\ &- (\nabla \log f_{\theta,m}(y)) (\nabla \log f_{\theta,m}(y))^T. \end{aligned} \quad (3c)$$

These derivative formulas are not obvious but are derived as equations (4.8), (4.9), (4.12), and (4.13) in the first author's thesis.

## 2 Asymptotic Variance

The asymptotic variance of  $\hat{\theta}_{m,n}$ , including both the sampling variation in  $y_1, \dots, y_n$  and the Monte Carlo variation in  $x_1, \dots, x_m$  is

$$J(\theta)^{-1} \left( \frac{V(\theta)}{n} + \frac{W(\theta)}{m} \right) J(\theta)^{-1} \quad (4)$$

where

$$V(\theta) = \text{var}\{\nabla \log f_{\theta}(Y)\} \quad (5a)$$

$$J(\theta) = E\{-\nabla^2 \log f_{\theta}(Y)\} \quad (5b)$$

$$W(\theta) = \text{var} \left\{ E \left[ \frac{\nabla f_{\theta}(X | Y)}{h(X)} \mid X \right] \right\} \quad (5c)$$

where  $X$  and  $Y$  here have the same distribution as  $x_i$  and  $y_j$ , respectively. This is the content of Theorem 3.3.1 in the first author's thesis.

The first two of these quantities have obvious "plug-in" estimators

$$\widehat{V}_{m,n}(\theta) = \frac{1}{n} \sum_{j=1}^n (\nabla \log f_{\theta,m}(y_j)) (\nabla \log f_{\theta,m}(y_j))^T \quad (6a)$$

$$\widehat{J}_{m,n}(\theta) = -\frac{1}{n} \sum_{j=1}^n \nabla^2 \log f_{\theta,m}(y_j) \quad (6b)$$

Thus a natural plug-in estimator is

$$\widehat{W}_{m,n}(\theta) = \frac{1}{m} \sum_{i=1}^m \widehat{S}_{m,n}(\theta, x_i) \widehat{S}_{m,n}(\theta, x_i)^T \quad (6c)$$

where

$$\widehat{S}_{m,n}(\theta, x) = \frac{1}{n} \sum_{j=1}^n (\nabla \log f_{\theta}(x, y_j) - \nabla \log f_{\theta,m}(y_j)) \cdot \frac{f_{\theta}(x, y_j)}{f_{\theta,m}(y_j)h(x)} \quad (6d)$$

See equations (2.7) and (2.9) in the first author's thesis.

Estimation of  $W$  using (6c) and (6d) has the drawback that it either uses  $O(mp)$  memory storing all the  $\log f_{\theta,m}(y_j)$  and their derivatives, where  $p$  is the dimension of the parameter vector  $\theta$  or it uses  $O(mnp)$  time recalculating these quantities. Neither alternative is attractive when  $m$  and  $n$  are large.

Thus we use an alternative method of estimating  $W$  based on the method of batch means, which is usually only used for time series. Let  $n = b \cdot l$ , where  $b$  and  $l$  are positive integers, called the *batch number* and *batch length*, respectively. For  $k = 1, \dots, b$  calculate

$$\widetilde{S}_{m,n,k}(\theta) = \frac{1}{l} \sum_{i=(k-1)l+1}^{kl} \widehat{S}_{m,n}(\theta, x_i) \quad (7a)$$

and use

$$\widetilde{W}_{m,n}(\theta) = \frac{l}{b} \sum_{k=1}^b \widetilde{S}_{m,n,k}(\theta) \widetilde{S}_{m,n,k}(\theta)^T. \quad (7b)$$

The factor  $l$  in (7b) comes from the fact that the batch means (7a) have  $1/l$  times the variance of the individual items (6d).

Using the method of batch means we can estimate  $W$  using  $O(p)$  memory and only  $O(bmp)$  in recalculation. Since the total time is necessarily at least  $O(mnp) + O(bp^2)$ , this recalculation is negligible so long as  $b$  is much smaller than  $n$ .

## 3 Bernoulli Regression with Random Effects

### 3.1 Normal Random Effects

The `bernor` package up through version 0.2 does only normal random effects.

### 3.1.1 Complete Data Density

The complete data density that for Bernoulli regression with normal random effects: the response  $y$  is conditionally Bernoulli given the fixed effect vector  $\beta$  and the random effect vector  $b$ . For this model we change notation, denoting the missing data by  $b$  rather  $x$ , which we used in the general discussion (to avoid confusion with “big  $X$ ” defined presently).

The “other data” for the problem consist of model matrices  $X$  and  $Z$ , both having row dimension equal to the length of  $y$ ,  $X$  having column dimension equal to the length of  $\beta$ , and  $Z$  having column dimension equal to the length of  $b$ . Then the “linear predictor” is

$$\eta = X\beta + Z\Sigma b \tag{8}$$

where  $\Sigma$  is a diagonal matrix that specifies the variance components. In R the linear predictor can be specified by

```
eta <- X %*% beta + Z %*% (sigma[i] * b)
```

where `sigma[i]` is the diagonal of  $\Sigma$ , `sigma` being a vector of scale parameters for the random effects and `i` being an index vector that says which scale parameter goes with which random effect (the lengths of `i` and `b` are equal, and each element of `i` is an integer in `seq(along = sigma)`).

Then

```
p <- 1 / (1 + exp(- eta))
```

is the vector of success probabilities. The complete data log density (or complete data log likelihood) is then

$$\log f_{\theta}(y, b) = \sum [y \log(p) + (1 - y) \log(1 - p)] + \sum \log \phi(b)$$

where the first sum runs over elements of  $y$  and  $p$  (which are the same length), the second sum runs over elements of  $b$ , and  $\phi$  is the density of elements of  $b$ , which are assumed to be IID mean zero normal. The parameter vector  $\theta$  combines  $\beta$  and  $\sigma$ .

### 3.1.2 Gradient

There are two types of elements of the gradient vector (partials with respect to  $\theta$ 's that are  $\beta$ 's and partials with respect to  $\theta$ 's that are  $\sigma$ 's). The first are

$$\nabla_{\beta} \log f_{\theta}(y, b) = (y - p)X. \tag{9a}$$

The second are

$$\frac{\partial}{\partial \sigma_k} \log f_{\theta}(y, b) = \sum_{j=1}^{|y|} (y_j - p_j) \sum_{\substack{m=1 \\ i_m=k}}^{|b|} z_{jm} b_m. \tag{9b}$$

For parallelism, we might as well rewrite (9a) to look more like (9b).

$$\frac{\partial}{\partial \beta_k} \log f_\theta(y, b) = \sum_{j=1}^{|y|} (y_j - p_j) x_{jk}. \quad (9c)$$

### 3.1.3 Hessian

The hessian is fairly simple. First, note that

$$\frac{\partial p_j}{\partial \eta_j} = p_j(1 - p_j).$$

So

$$\frac{\partial^2}{\partial \beta_k \partial \beta_l} \log f_\theta(y, b) = - \sum_{j=1}^{|y|} p_j(1 - p_j) x_{jk} x_{jl} \quad (10a)$$

$$\frac{\partial^2}{\partial \sigma_k \partial \sigma_l} \log f_\theta(y, b) = - \sum_{j=1}^{|y|} p_j(1 - p_j) \sum_{\substack{m=1 \\ i_m=k}}^{|b|} z_{jm} b_m \sum_{\substack{n=1 \\ i_n=l}}^{|b|} z_{jn} b_n \quad (10b)$$

$$\frac{\partial^2}{\partial \beta_k \partial \sigma_l} \log f_\theta(y, b) = - \sum_{j=1}^{|y|} p_j(1 - p_j) x_{jk} \sum_{\substack{n=1 \\ i_n=l}}^{|b|} z_{jn} b_n \quad (10c)$$