

# Appendix S1. Updating of latent encounter history frequencies in **multimark**

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1 The feasible set of latent encounter histories ( $\mathbf{Y}$ ) is explored in **multimark** using  
2 an extension of the MCMC algorithms proposed by Bonner & Holmberg (2013)  
3 and McClintock *et al.* (2013, 2014) that were originally conceived for a different  
4 application by Link *et al.* (2010). The new algorithm conditions on the observed  
5 data (thus reducing the dimension of the problem) and only proposes updates with  
6 non-negative latent encounter history frequencies,  $\mathbf{x} = (x_1, x_2, \dots, x_{5^T})$ . Let  $\mathbf{r}$  denote  
7 the set of  $4^T - 2^{T+1} + 1$  indices for latent encounter histories that spawn  $>1$  observed  
8 history, and let  $x_{j(1)}$  and  $x_{j(2)}$  denote the corresponding frequencies for type 1 and  
9 type 2 histories that arise from latent encounter history  $j \in \mathbf{r}$ . Referring back to  
10 Table 2 with  $T = 2$ ,  $\mathbf{r} = \{4, 8, 9, 12, 14, 16, 17, 18, 19\}$  and one potential update would  
11 involve frequencies for latent history ‘31’ ( $x_{17}$ ) and its progeny ‘11’ ( $x_{17(1)} = x_7$ ) and  
12 ‘20’ ( $x_{17(2)} = x_{11}$ ).

13 When conditioning on the observed encounter histories, the size of the prob-  
 14 lem is typically greatly reduced because many of the potential latent histories and  
 15 corresponding moves are not permissable. For example, with  $T = 2$ , if encounter  
 16 history ‘11’ was never observed, then  $x_7 = x_9 = x_{17} = x_{19} = 0$  can be ignored and  
 17  $j \in \{9, 17, 19\}$  can be removed from  $\mathbf{r}$  for subsequent computations.

18 Starting from a permissible  $\mathbf{x}$  conditional on the observed encounter histories, the  
 19 algorithm proceeds as follows:

- 20 1. Randomly draw a latent encounter history index  $r \in \mathbf{r}^*$ , where  $\mathbf{r}^*$  is the subset of  
 21  $\mathbf{r}$  with corresponding frequencies that satisfy  $\min(x_1, x_j) + \min(x_{j(1)}, x_{j(2)}) > 0$   
 22 for  $j \in \mathbf{r}$ .
- 23 2. Randomly draw  $c_r$  from the integer set  $\{-\min(x_1, x_r), \dots, -1, 1, \dots, \min(x_{r(1)}, x_{r(2)})\}$ .
- 24 3. Propose  $x_r^* = x_r + c_r$ ,  $x_{r(1)}^* = x_{r(1)} - c_r$ , and  $x_{r(2)}^* = x_{r(2)} - c_r$ .
- 25 4. Apportion  $\mathbf{x}^*$  to individuals following McClintock *et al.* (2014), and accept  
 26 proposed move based on the Metropolis-Hastings ratio described therein [pp.  
 27 2470-2472, steps 9(b)-9(c)].

28 Any additional constraints, such as those resulting from encounter histories be-  
 29 ing designated as known with certainty using the *known* argument in *processdata()*,  
 30 are accounted for by simple modifications to steps 1-2. In terms of mixing, it can  
 31 sometimes be advantageous to explore more than one move at a time. At each iter-  
 32 ation of the chain, the argument *maxnumbasis* specifies how many times to perform  
 33 steps 1-3 in sequence before evaluating step 4. The default for *multimarkCJS()* and  
 34 *multimarkClosed()* is *maxnumbasis=1*.

35 Note that because `multimark` uses “semi-complete” data likelihoods that con-  
 36 dition on the number of unique individuals encountered at least once ( $n$ ), the di-  
 37 mension of the data-augmented encounter histories ( $M$ ) described in McClintock  
 38 *et al.* (2014) is determined by the number of observed encounter histories (i.e.  $M =$   
 39  $n_1 + n_2 + n_{known}$ ) such that  $x_1 = M - n$ . Letting  $w_i \sim \text{Bernoulli}(\psi)$  be an indicator  
 40 for whether or not individual  $i$  belongs to the  $n$  unique individuals encountered at  
 41 least once (i.e.  $\sum_{i=1}^M w_i = n$ ), then  $w_i = 1$  if  $H_i > 1$  (otherwise  $w_i = 0$ ), where  
 42  $H_i$  is the latent encounter history index for individual  $i$  ( $\sum_{i=1}^M I(H_i = j) = x_j$ ), and  
 43  $\psi \sim \text{Beta}(a_\psi^0, b_\psi^0)$  is the probability that a randomly selected individual from the  $M$   
 44 observed individuals belongs to the  $n$  unique individuals encountered at least once.  
 45 The defaults in `multimarkCJS()` and `multimarkClosed()` are  $a_\psi^0 = b_\psi^0 = 1$ .

## 46 References

- 47 Bonner, S.J. & Holmberg, J. (2013) Mark-recapture with multiple, non-invasive  
 48 marks. *Biometrics*, **69**, 766–775.
- 49 Link, W.A., Yoshizaki, J., Bailey, L.L. & Pollock, K.H. (2010) Uncovering a latent  
 50 multinomial: analysis of mark-recapture data with misidentification. *Biometrics*,  
 51 **66**, 178–185.
- 52 McClintock, B.T., Bailey, L.L., Dreher, B.P. & Link, W.A. (2014) Probit models for  
 53 capture-recapture data subject to imperfect detection, individual heterogeneity  
 54 and misidentification. *The Annals of Applied Statistics*, **8**, 2461–2484.
- 55 McClintock, B.T., Conn, P.B., Alonso, R.S. & Crooks, K.R. (2013) Integrated mod-

56 eling of bilateral photo-identification data in mark-recapture analyses. *Ecology*,  
57 **94**, 1464–1471.