#### Peter D Smits

# Committee on Evolutionary Biology, University of Chicago

October 2, 2015

#### Abstract

# 1 Introduction

# 2 Materials and Methods

#### 2.1 Taxon occurrence information

### 2.2 Geographic provinces

#### 2.3 Model specification

Taxon presence was modeled has a hierarchical two-state hidden Markov model (HMM).

 $y_{i,j,t}$  is the observed occurrence of taxon i in province j at time t, where  $i=1,2,\ldots,N,\,j=1,2,\ldots,J,$  and  $t=1,2,\ldots,T.\,y=1$  is occupied while y=0 is unoccupied.  $z_{i,j,t}$  is the "true" occurrence of taxon i in province j at time t, given the estimate of sampling. Just as with y,z=1 is occupied while z=0 is unoccupied.

 $\phi_{j,t}$  is the probability of surviving, in province j, from time t to time t+1  $(Pr(z_{t+1}=1|z_t=1))$ .  $\gamma_{j,t}$  is the probability of newly entering province j at time t+1  $(Pr(z_{t+1}=1|z_t=0))$ .  $p_{j,t}$  is the probability of preservation (Pr(y=1|z=1)) in province j at time t.

 $\psi$  is probability of sit occupancy/probability of occurrence  $(Pr(z_{i,t=1}=1))$ . The first time point is defined in terms of  $\psi$  because there is (assumed) no previous time points.

$$y_{i,t,j} \sim \text{Bern}(p_{t,j}z_{i,t,j})$$
  
 $z_{i,t=1,j} \sim \text{Bern}(\psi_j)$  (1)  
 $z_{i,t,j} \sim \text{Bern}(\phi_{j,t-1}z_{i,t-1,j} + \gamma_{j,t-1}(1 - z_{i,t-1,j}))$ 

The parameters  $\phi$ ,  $\gamma$ , and p are then all defined hierarchically within each province, hierarchical by time bin with the mean of that hierarchical by province.  $\Phi_j$ ,  $\Gamma_j$ , and  $P_j$  are the overall probabilities for province j.  $M_{\phi}$ ,  $M_{\gamma}$ , and  $M_p$  are the overall estimates of survival, origination, and preservation probabilities.

Diversity dependent origination and survival was included as a regression coefficients in the parameterizations of  $\phi_{j,t}$ , and  $\gamma_{j,t}$ . In particular, the diversity of any province can affect the origination and survival probabilities of an province.  $\alpha$  and  $\beta$  are both  $J \times J$  matrices of regression coefficients, where  $\beta_j$  would be a column vector. X is a  $J \times T$  matrix where  $X_t$  is a column vector where each element is defined  $X_{j,t} = \sum_{i=1}^{N} z_{i,j,t}$  (i.e. the sum of the diversity in province j at time t). All regression coefficients (i.e. all elements of the matrices) are given weakly-informative independent normally distributed priors.

And finally, I use independent priors for  $\psi_j$  by province  $j: \psi_j \sim \mathrm{U}(0,1)$ .

$$\log \operatorname{id}(\phi_{j,t}) \sim \operatorname{N}(\Phi_{j} + \beta_{j}^{T} X_{t-1}, \sigma_{\phi,j})$$

$$\Phi_{j} \sim \operatorname{N}(M_{\phi}, \sigma_{\Phi})$$

$$\sigma_{\phi,j} \sim \operatorname{C}^{+}(1)$$

$$M_{\phi} \sim \operatorname{N}(0,1)$$

$$\sigma_{\Phi} \sim \operatorname{C}^{+}(1)$$

$$\beta \sim \operatorname{N}(0,1)$$
(2)

$$\log \operatorname{id}(\gamma_{j,t}) \sim \operatorname{N}(\Gamma_j + \alpha_j^t X_{t-1}, \sigma_{\gamma,j})$$

$$\Gamma_j \sim \operatorname{N}(M_{\gamma}, \sigma_{\Gamma})$$

$$\sigma_{\gamma,j} \sim \operatorname{C}^+(1)$$

$$M_{\gamma} \sim \operatorname{N}(0,1)$$

$$\sigma_{\Gamma} \sim \operatorname{C}^+(1)$$

$$\alpha \sim \operatorname{N}(0,1)$$
(3)

$$\log \operatorname{ic}(p_{j,t}) \sim \operatorname{N}(P_j, \sigma_{p,j})$$

$$P_j \sim \operatorname{N}(M_p, \sigma_P)$$

$$\sigma_{p,j} \sim \operatorname{C}^+(1)$$

$$M_p \sim \operatorname{N}(0, 1)$$

$$\sigma_P \sim \operatorname{C}^+(1)$$
(4)

- 3 Results
- 4 Discussion

Acknowledgements

# References