# Metapopulations with dynamic extinction probabilities

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#### **SPOM - Phase structure**

For many species the propensity for colonization and local extinction is markedly different in different phases of their life cycle. Examples:

The Vernal pool fairy shrimp (Branchinecta lynchi) and the California linderiella (Linderiella occidentalis), both listed under the Endangered Species Act (USA)

The Jasper Ridge population of Bay checkerspot butterfly (Euphydryas editha bayensis), now extinct





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We will we assume that the population is *observed after successive extinction phases* (CE Model).

*Colonization*: unoccupied patch *i* becomes occupied with probability

$$c\left(\frac{1}{n}\sum_{j=1}^n X_{j,t}^{(n)}d(z_i,z_j)a_j\right),\,$$

where  $d(z, \tilde{z}) \ge 0$  measures the ease of movement between patches located at z and  $\tilde{z}$ ,  $a_j$  is a weight related to the size of the patch j and  $c : [0, \infty) \rightarrow [0, 1]$  (called the *colonisation function*) is increasing and Lipschitz continuous, with c(0) = 0and c'(0) > 0. For simplicity, take  $d \equiv 1$  and  $a \equiv 1$ . So, ...

*Colonization*: unoccupied patch *i* becomes occupied with probability  $c(n^{-1}\sum_{j=1}^{n} X_{j,t}^{(n)})$ , where  $c: [0,1] \rightarrow [0,1]$  (called the *colonisation function*) is increasing and Lipschitz continuous, with c(0) = 0 and c'(0) > 0.

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Proportion of patches occupied

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Then, given the current state  $X_t^{(n)}$  and survival probabilities  $s_t$ , the  $X_{i,t+1}^{(n)}$  (i = 1, ..., n) are independent with transitions

$$\Pr\left(X_{i,t+1}^{(n)}=1 \,|\, X_t^{(n)}, s_t\right) = s_{i,t} X_{i,t}^{(n)} + s_{i,t} \, c \left(n^{-1} \sum_{j=1}^n X_{j,t}^{(n)}\right) \left(1 - X_{i,t}^{(n)}\right).$$

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We will assume that  $(s_{i,t})_{t=0}^{\infty}$  (i = 1, ..., n) are independent Markov chains taking values in [0, 1] with common transition kernel P(s, dr).

This covers the simple but important case where patches are classified as being suitable or unsuitable for occupancy.

## **SPOM - Homogeneous case**

In the *homogeneous case*, where  $s_i = s$  is the same for each *i*, the *number*  $N_t^{(n)}$  of occupied patches at time *t* is Markovian, and, letting the initial number  $N_0^{(n)}$  of occupied patches grow at the same rate as *n* we arrive at:

**Theorem** If  $N_0^{(n)}/n \xrightarrow{p} x_0$  (a constant), then

$$N_t^{(n)}/n \xrightarrow{p} x_t$$
, for all  $t \ge 1$ ,

with  $(x_t)$  determined by  $x_{t+1} = f(x_t)$ , where



#### **CE Model - Evanescence**



#### **CE Model - Quasi stationarity**



 $x_{t+1} = f(x_t)$ , where f(x) = s(x + (1 - x)c(x)).

*Evanescence*:  $1 + c'(0) \le 1/s$ . 0 is the unique fixed point in [0, 1]. It is stable.

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#### **CE Model - Evanescence**



#### **CE Model - Quasi stationarity**



Return now to the general case, where patch survival probabilities evolve in time, and we keep track of which patches are occupied ....

$$\Pr\left(X_{i,t+1}^{(n)}=1 \,|\, X_t^{(n)}, s_t\right) = s_{i,t} X_{i,t}^{(n)} + s_{i,t} \, c \left(n^{-1} \sum_{j=1}^n X_{j,t}^{(n)}\right) \left(1 - X_{i,t}^{(n)}\right).$$

Treat the collection of patch survival probabilities and those of *occupied patches* at time t as point processes on [0, 1].

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 $\sigma_{n,t}(B) = \#\{s_{i,t} \in B\}/n, \qquad B \in \mathcal{B}([0,1]),$ 

 $\mu_{n,t}(B) = \#\{s_{i,t} \in B : X_{i,t}^{(n)} = 1\}/n, \qquad B \in \mathcal{B}([0,1]).$ 

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Think of  $\sigma_0$  as being the initial distribution of survival probabilities.

Equivalently, we may define  $(\sigma_{n,t})$  and  $(\mu_{n,t})$  by

$$\int h(s)\sigma_{n,t}(ds) = \frac{1}{n} \sum_{i=1}^{n} h(s_{i,t})$$
$$\int h(s)\mu_{n,t}(ds) = \frac{1}{n} \sum_{i=1}^{n} X_{i,t}^{(n)} h(s_{i,t}),$$

for *h* in  $C^+([0,1])$ , the class of continuous functions that map [0,1] to  $[0,\infty)$ .

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for *h* in  $C^+([0,1])$ , the class of continuous functions that map [0,1] to  $[0,\infty)$ . For example  $(h \equiv 1)$ ,

$$\int \mu_{n,t}(ds) = \frac{1}{n} \sum_{i=1}^{n} X_{i,t}^{(n)}$$

(proportion occupied).

Suppose that  $\sigma_{n,0} \xrightarrow{d} \sigma_0$  for some non-random (probability) measure  $\sigma_0$ . Although this assumption concerns only the initial variation in the survival probabilities, it implies a similar 'law of large numbers' for them at all times.

**Lemma**  $\sigma_{n,t} \xrightarrow{d} \sigma_t$ , where  $\sigma_t$  is defined by the recursion  $\int h(s)\sigma_{t+1}(ds) = \int h(s) \int P(r, ds)\sigma_t(dr),$ 

for all  $h \in C^+([0,1])$ .

## A measure-valued difference equation

**Theorem** Suppose that  $\mu_{n,0} \xrightarrow{d} \mu_0$  for some non-random measure  $\mu_0$ . Then,  $\mu_{n,t} \xrightarrow{d} \mu_t$  for all t = 1, 2, ..., where  $\mu_t$  is defined by the following recursion: for  $h \in C^+([0,1])$ ,

$$\int h(s)\mu_{t+1}(ds) = c_t \int s \int h(r)P(s,dr)\sigma_t(ds)$$
$$(1-c_t) \int s \int h(r)P(s,dr)\mu_t(ds),$$

where  $c_t = c(\mu_t([0, 1])) = c(\int \mu_t(ds))$ .

#### **Survival probability simulation**



#### **CE Model - Evanescence**



#### **CE Model - Persistence**



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#### **CE Model - Evanescence**



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