

Infinite-patch metapopulation models: branching, convergence and chaos

Phil. Pollett

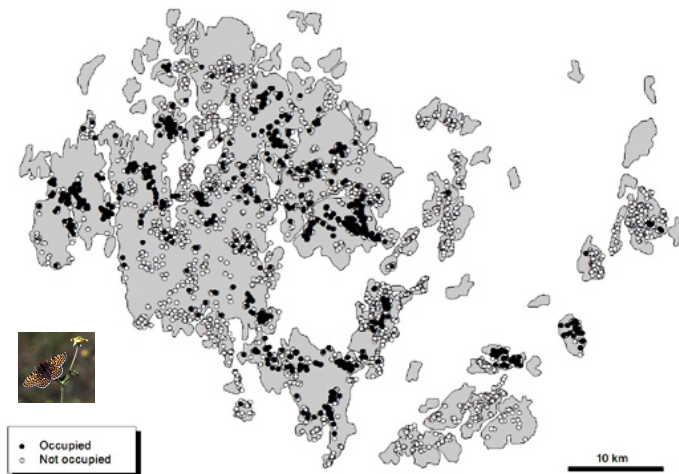
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Metapopulations



Glanville fritillary butterfly (*Melitaea cinxia*) in the Åland Islands in Autumn 2005.

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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)

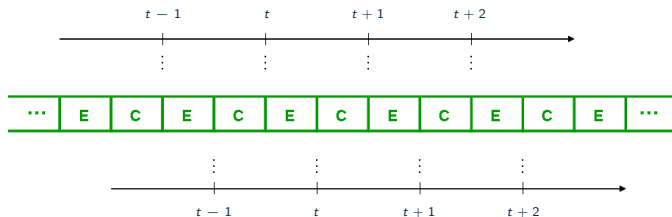


The Jasper Ridge population of Bay checkerspot butterfly (*Euphydryas editha bayensis*), now extinct



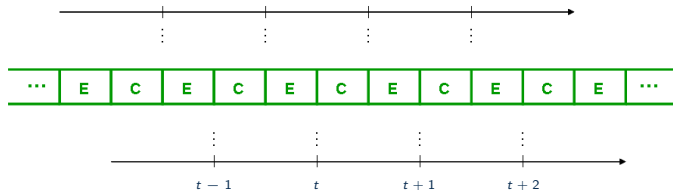
Phase structure

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

Colonization and extinction happen in distinct, successive phases.

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We thus have the following *Chain Binomial* structure¹:

$$n_{t+1} \stackrel{D}{=} \text{Bin}\left(n_t + \text{Bin}\left(N - n_t, c(n_t/N)\right), s\right)$$

[$\text{Bin}(m, p)$ is a binomial random variable with m trials and success probability p .]

¹Buckley, F.M. and Pollett, P.K. (2010) Limit theorems for discrete-time metapopulation models. *Probability Surveys* 7, 53–83.

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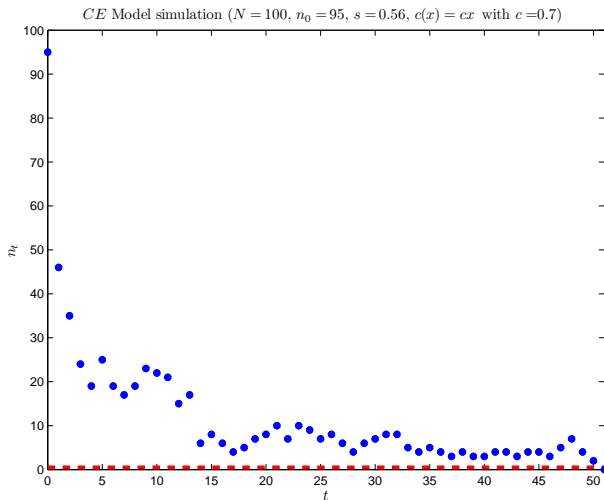
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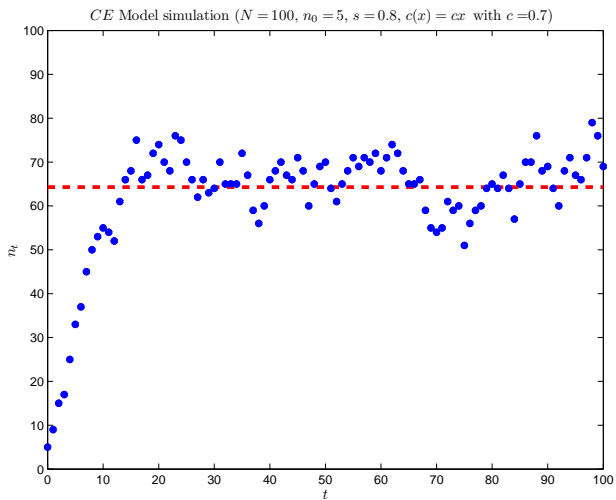
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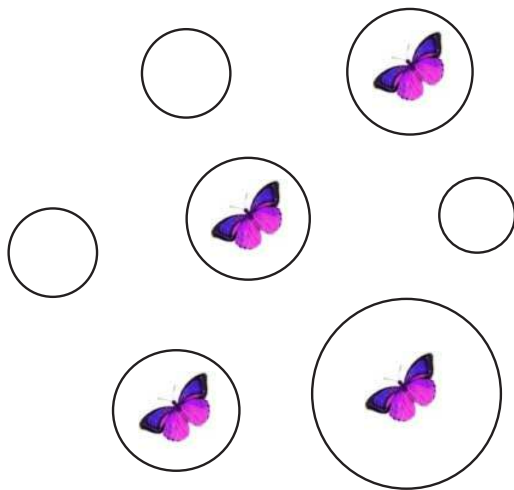
Evanescence: $c'(0) \leq (1 - s)/s$



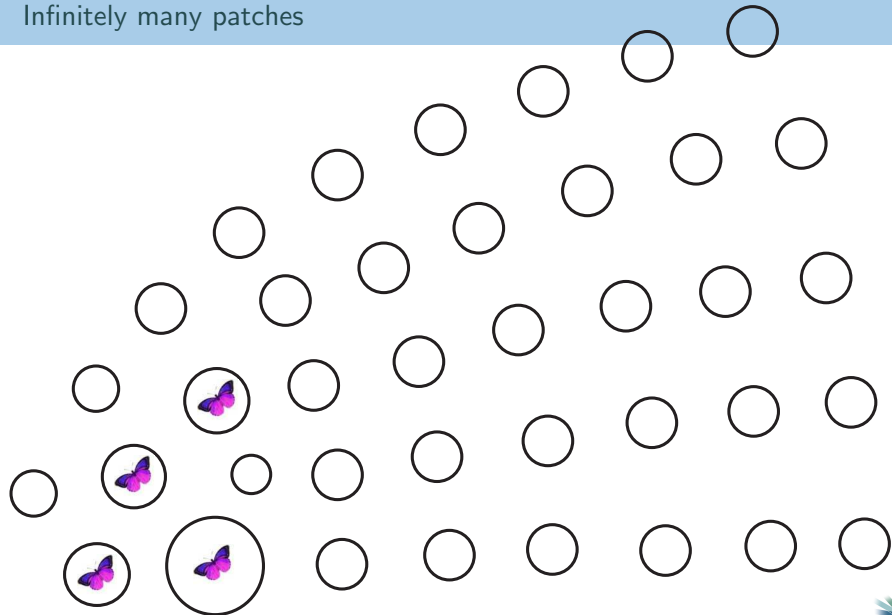
Quasi stationarity: $c'(0) > (1 - s)/s$



N patches



Infinitely many patches



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$$\text{Bin}(N - n, c(n/N)) \xrightarrow{d} \text{Poi}(mn), \quad \text{as } N \rightarrow \infty,$$

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Claim The process $(n_t, t = 0, 1, \dots)$ is a *branching process* (Galton-Watson-Bienaymé process) whose offspring distribution has pgf $G(z) = (1 - s(1 - z))e^{-ms(1-z)}$.

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(Recall the earlier condition for evanescence: $c'(0) \leq (1 - s)/s$)

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For some index N write $m(n) = N\mu(n/N)$, where μ is a continuous function. We may take N to be simply n_0 or, more generally, following Klebaner², we may interpret N as being a 'threshold' with the property that $n_0/N \rightarrow x_0$ as $N \rightarrow \infty$.

²Klebaner, F.C. (1993) Population-dependent branching processes with a threshold. Stochastic Process. Appl. 46, 115–127.

By choosing μ appropriately, we may allow for a degree of regulation in the colonization process.

For example, $\mu(x)$ might be of the form

- $\mu(x) = rx(a - x)$ ($0 \leq x \leq a$) (logistic growth);
- $\mu(x) = xe^{r(1-x)}$ ($x \geq 0$) (Ricker dynamics);
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Theorem 2 If $X_0^N \xrightarrow{P} x_0$ as $N \rightarrow \infty$, then $X_t^N \xrightarrow{P} x_t$ for all $t \geq 1$, where (x_t) is determined by $x_{t+1} = f(x_t)$ ($t \geq 0$) with $f(x) = s(x + \mu(x))$.

Infinite-patch SPOM with regulation

The proof uses the following very useful result.

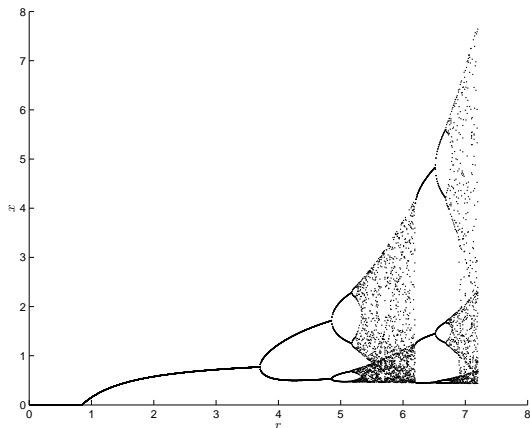
Lemma³ Let U_n , V_n , and u be random variables, where U_n and u are scalar. If $\mathbb{E}(U_n|V_n) \xrightarrow{P} u$ and $\text{Var}(U_n|V_n) \xrightarrow{P} 0$ then $U_n \xrightarrow{P} u$.

³McVinish, R. and Pollett, P.K. (2012) The limiting behaviour of a mainland-island metapopulation. *Journal of Mathematical Biology* 64, 775–801.

Proof: We will use mathematical induction. Suppose $X_t^N \xrightarrow{P} x_t$ for some $t \geq 0$. Since $n_{t+1} \stackrel{D}{=} \text{Bin}(n_t + \text{Poi}(m(n_t)), s)$, a simple calculation gives $\mathbb{E}(n_{t+1}|n_t) = s(n_t + m(n_t))$. But, $m(n) = N\mu(n/N)$. So, dividing by N gives $\mathbb{E}(X_{t+1}^N|X_t^N) = f(X_t^N)$, where $f(x) = s(x + \mu(x))$. Since μ is continuous, so is f , and so $\mathbb{E}(X_{t+1}^N|X_t^N) \xrightarrow{P} f(x_t) = x_{t+1}$. Another simple calculation yields $\text{Var}(n_{t+1}|n_t) = s((1-s)n_t + m(n_t))$, and so $N\text{Var}(X_{t+1}^N|X_t^N) = v(X_t^N)$, where $v(x) = s((1-s)x + \mu(x))$. Since v is continuous, $v(X_t^N) \xrightarrow{P} v(x_t)$, and hence $\text{Var}(X_{t+1}^N|X_t^N) \xrightarrow{P} 0$. Using the technical lemma we arrive at $X_{t+1}^N \xrightarrow{P} x_{t+1}$, and the proof is complete.

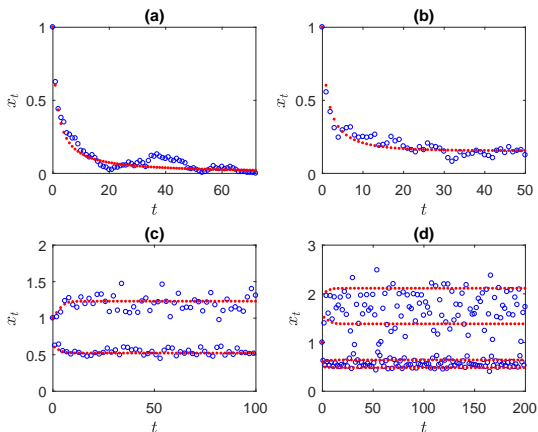


Infinite-patch SPOM with regulation



Bifurcation diagram for the infinite-patch deterministic model with colonization following Ricker growth dynamics: $x_{t+1} = 0.3 x_t (1 + e^{r(1-x_t)})$ (r ranges from 0 to 7.2).

Infinite-patch SPOM with regulation



Simulation (blue circles) of the infinite-patch model with colonization following Ricker growth dynamics, together with the corresponding limiting deterministic trajectories (solid red). Here $s = 0.3$, $N = 200$, and (a) $r = 0.84$, (b) $r = 1$ (c) $r = 4$, (d) $r = 5$.

We can also get a handle on the fluctuations of (X_t^N) about (x_t) . Define Z^N by $Z_t^N = \sqrt{N}(X_t^N - x_t)$ ($t \geq 0$).

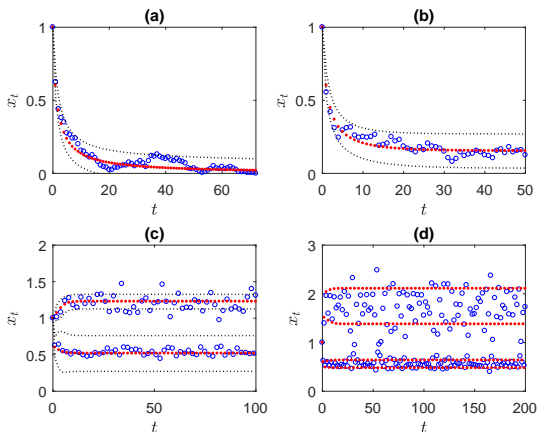
Theorem 3 Suppose that μ is twice continuously differentiable with bounded second derivative, and suppose that $Z_0^N \xrightarrow{d} z_0$. Then, Z^N converges weakly to the Gaussian Markov chain Z defined by $Z_{t+1} \stackrel{D}{=} s(1 + \mu'(x_t))Z_t + E_t$, starting at $(Z_0 =) z_0$, with (E_t) independent and $E_t \sim N(0, v(x_t))$, where $v(x) = s((1-s)x + \mu(x))$.

The proof follows the programme laid out in the proof of Theorem 1 of

Klebaner, F.C. and Nerman, O. (1994) Autoregressive approximation in branching processes with a threshold. *Stochastic Process. Appl.* 51, 1–7,

but note that (n_t) is not a *population-dependent branching processes with threshold*; see last slide.

Infinite-patch SPOM with regulation



Same graphs as earlier, but now in (a), (b) and (c), the black dotted lines indicate ± 2 standard deviations of the Gaussian approximation (in (c) every *second* point is proximate, thus indicating the extent of variation about each of the two limit cycle values).

Recall that $f(x) = s(x + \mu(x))$. Notice that x^* will be a fixed point of f if and only if $\mu(x^*) = \rho x^*$, where $\rho = (1 - s)/s$. Clearly 0 is a fixed point, but there might be others. If there *is* a unique positive fixed point x^* , it will be stable if $\mu'(x^*) < 1$ and unstable if $\mu'(x^*) > 1$ (need to consider higher derivatives when $\mu'(x^*) = 1$).

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Corollary 1 Suppose that f admits a unique positive stable fixed point x^* . Then, if $X_0^N \xrightarrow{P} x^*$, $x_t = x^*$ for all t and, assuming $Z_0^N \rightarrow z_0$, the limit process Z is an AR-1 process of the form $Z_{t+1} \stackrel{D}{=} s(1 + \mu'(x^*))Z_t + E_t$, starting at $(Z_0 =) z_0$, with iid errors $E_t \sim N(0, (1 - s^2)x^*)$.

Corollary 2 Suppose that f admits a stable limit cycle $x_0^*, x_1^*, \dots, x_{d-1}^*$ with $X_0^N \xrightarrow{P} x_0^*$. Then, $x_{nd+j} = x_j^*$ ($n \geq 0, j = 0, \dots, d-1$) and, assuming $Z_0^N \rightarrow z_0$, the limit process Z has the following representation: $(Y_n, n \geq 0)$, where $Y_n = (Z_{nd}, Z_{nd+1}, \dots, Z_{(n+1)d-1})^\top$ with $Z_0 = z_0$, is a d -variate AR-1 process of the form $Y_{n+1} \stackrel{D}{=} AY_n + E_n$, with iid errors $E_n \sim N(\mathbf{0}, \Sigma_d)$; A is the $d \times d$ matrix

$$A = \begin{pmatrix} 0 & 0 & \cdots & a_1 \\ 0 & 0 & \cdots & a_2 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & a_d \end{pmatrix},$$

where $a_j = s^j \prod_{i=0}^{j-1} (1 + \mu'(x_i^*))$, $\Sigma_d = (\sigma_{ij})$ is the $d \times d$ symmetric matrix with entries

$$\sigma_{ij} = a_i a_j \sum_{k=0}^{i-1} v(x_k^*) / a_{k+1}^2 \quad (1 \leq i \leq j \leq d),$$

where $v(x) = s((1-s)x + \mu(x))$, and the random entries, (Z_1, \dots, Z_{d-1}) , of Y_0 have a Gaussian $N(\mathbf{a}z_0, \Sigma_{d-1})$ distribution, where $\mathbf{a} = (a_1, \dots, a_{d-1})$. Furthermore, (Y_n) has a Gaussian $N(\mathbf{0}, V)$ stationary distribution, where $V = (v_{ij})$ has entries $v_{ij} = \sigma_{ij} / (1 - a_d^2)$.

Recall that $n_{t+1} \stackrel{D}{=} \text{Bin}(n_t + \text{Poi}(m(n_t)), s)$. Whilst (n_t) does not exhibit the branching property (required for it to be a *population-dependent branching processes with threshold*), we can say the following.

Theorem $n_{t+1} \stackrel{D}{=} \text{Bin}(n_t, s) + \text{Poi}(sm(n_t))$ (independent RVs).

Proof:

$$\begin{aligned} \mathbb{E}(z^{n_{t+1}} | n_t) &= \mathbb{E}\left(\mathbb{E}\left(z^{n_{t+1}} | \text{Poi}(m(n_t)), n_t\right) \middle| n_t\right) \\ &= \mathbb{E}\left((1 - s + sz)^{n_t + \text{Poi}(m(n_t))} \middle| n_t\right) \\ &= (1 - s + sz)^{n_t} \mathbb{E}\left((1 - s + sz)^{\text{Poi}(m(n_t))} \middle| n_t\right) \\ &= (1 - s(1 - z))^{n_t} e^{-sm(n_t)(1-z)} \end{aligned}$$