

# Stochastic models for population networks

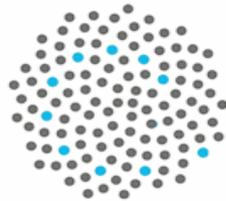
## II: Discrete-time patch occupancy models [Exact results]

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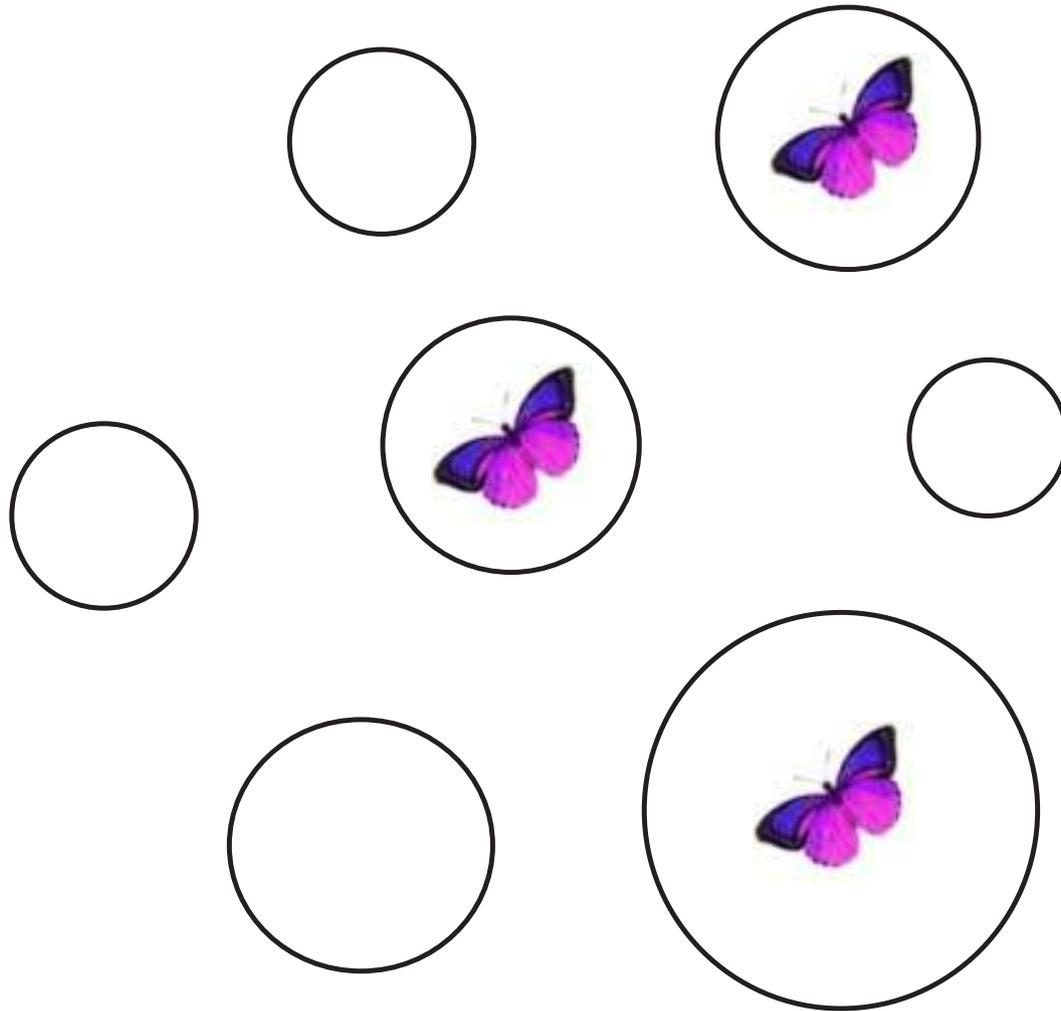


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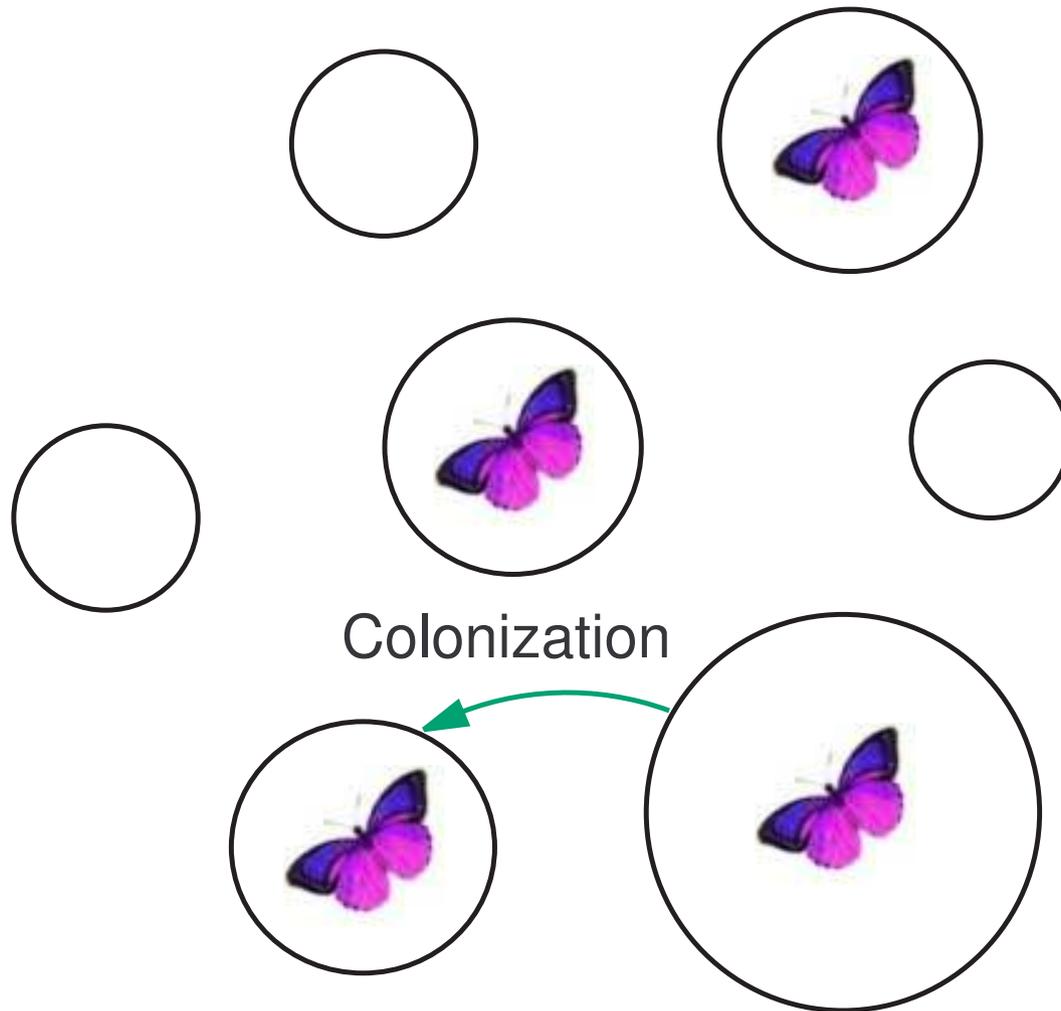
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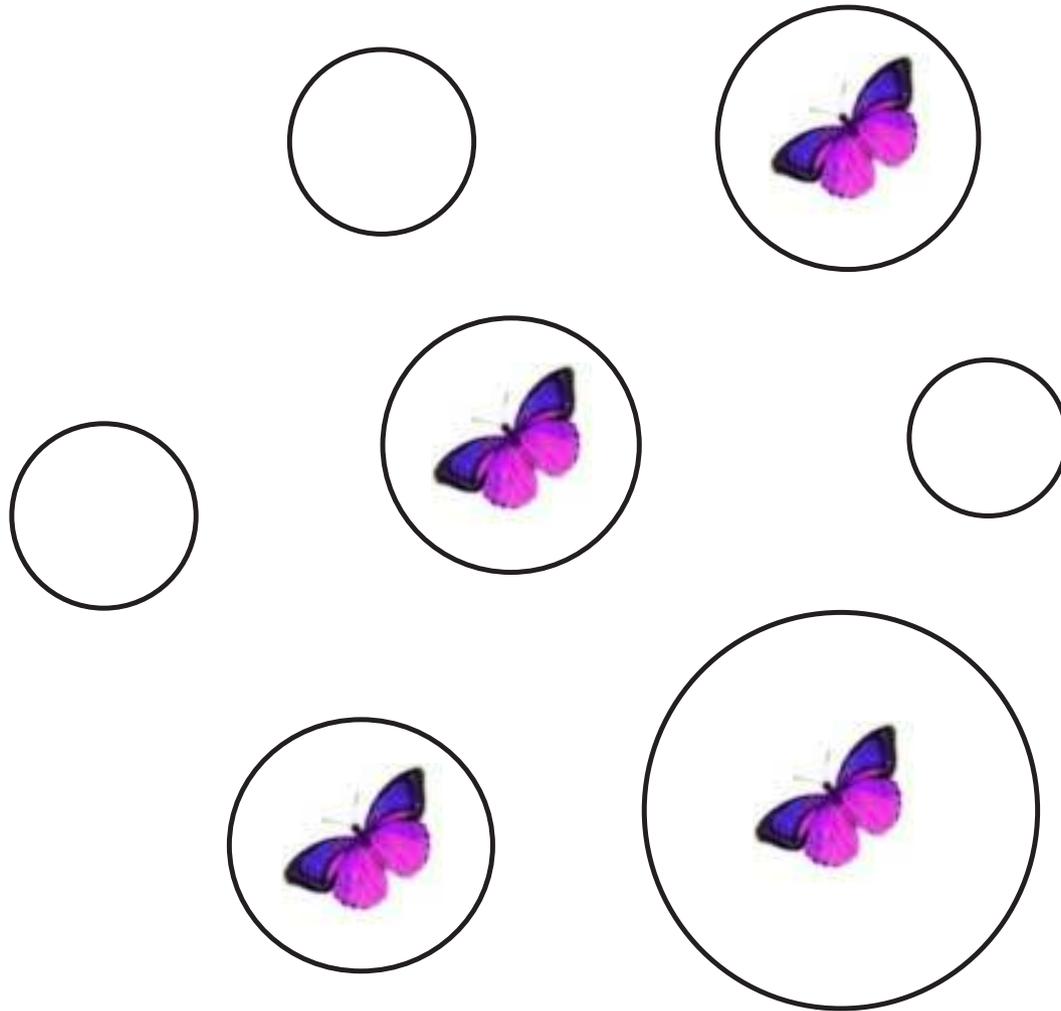
# Metapopulations



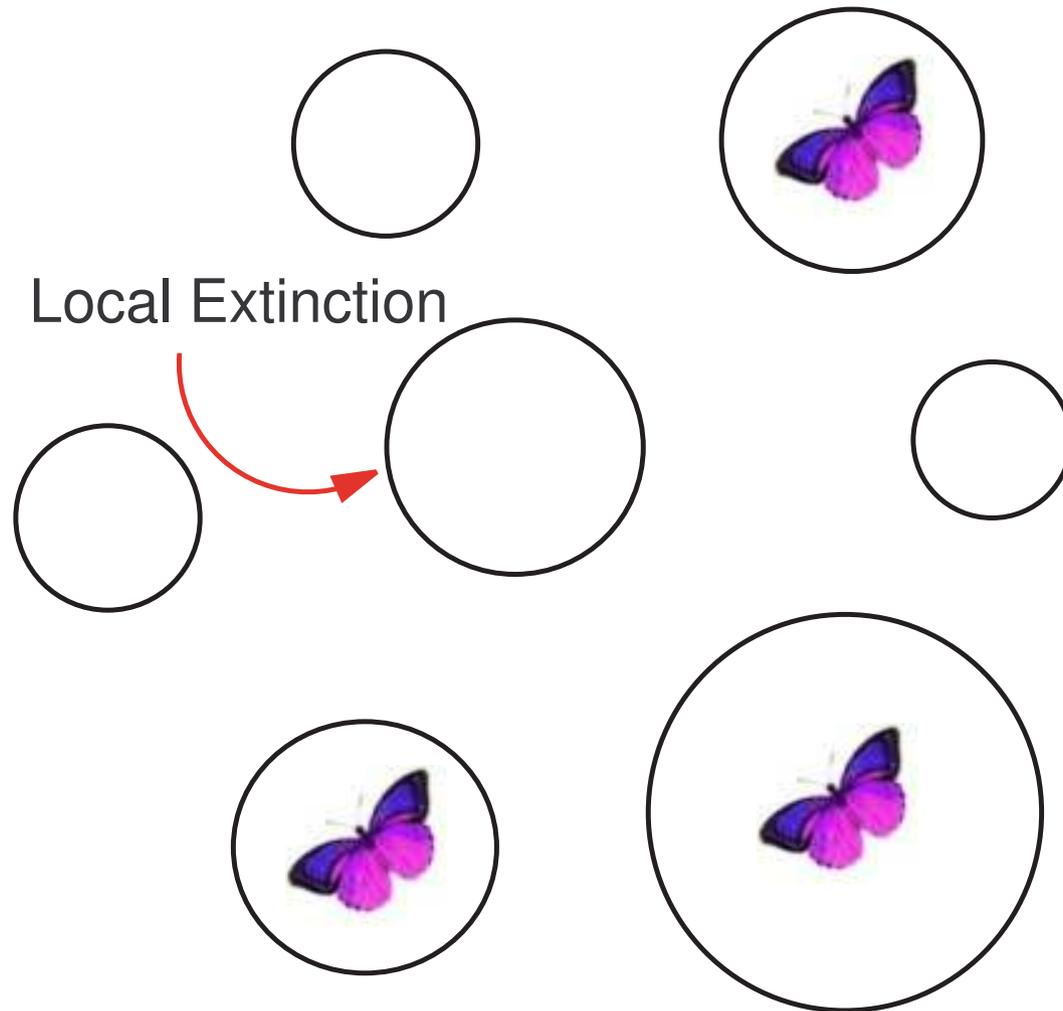
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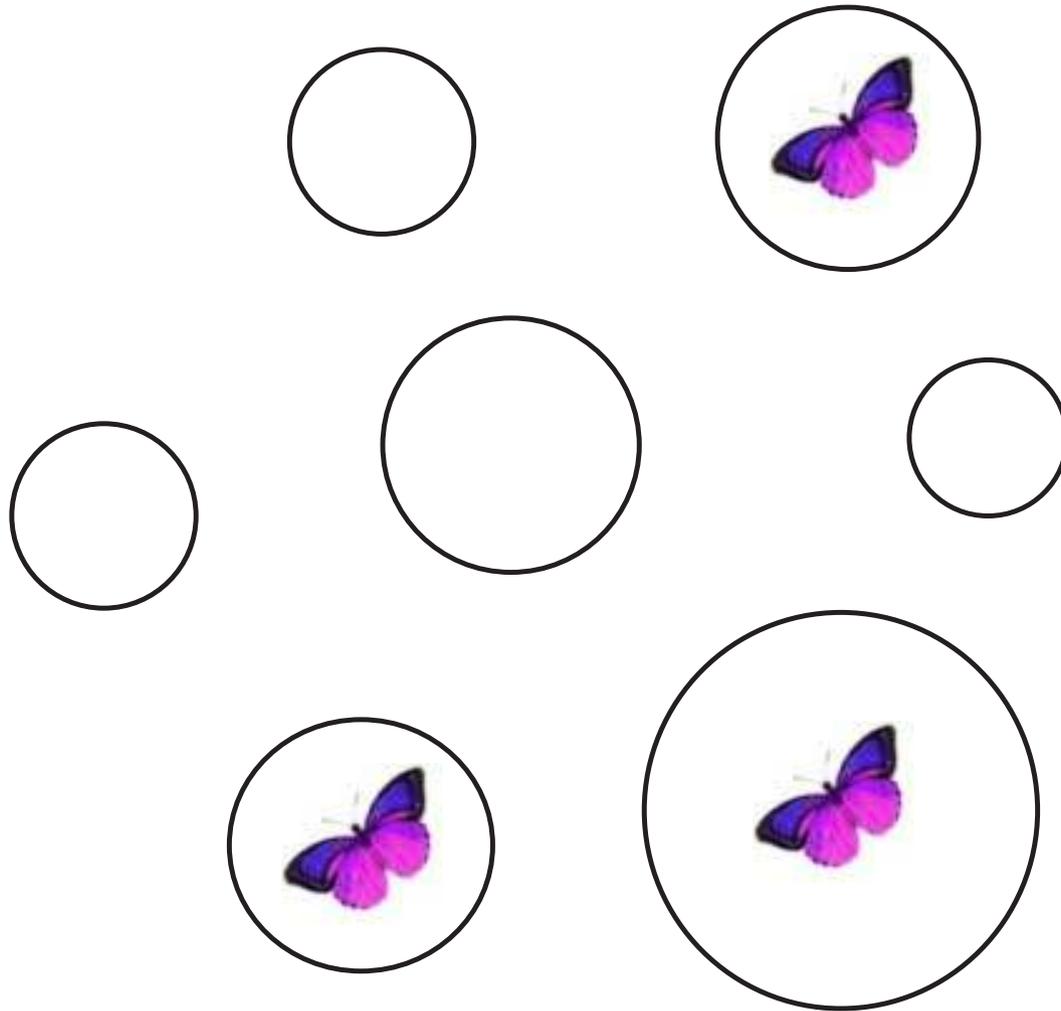
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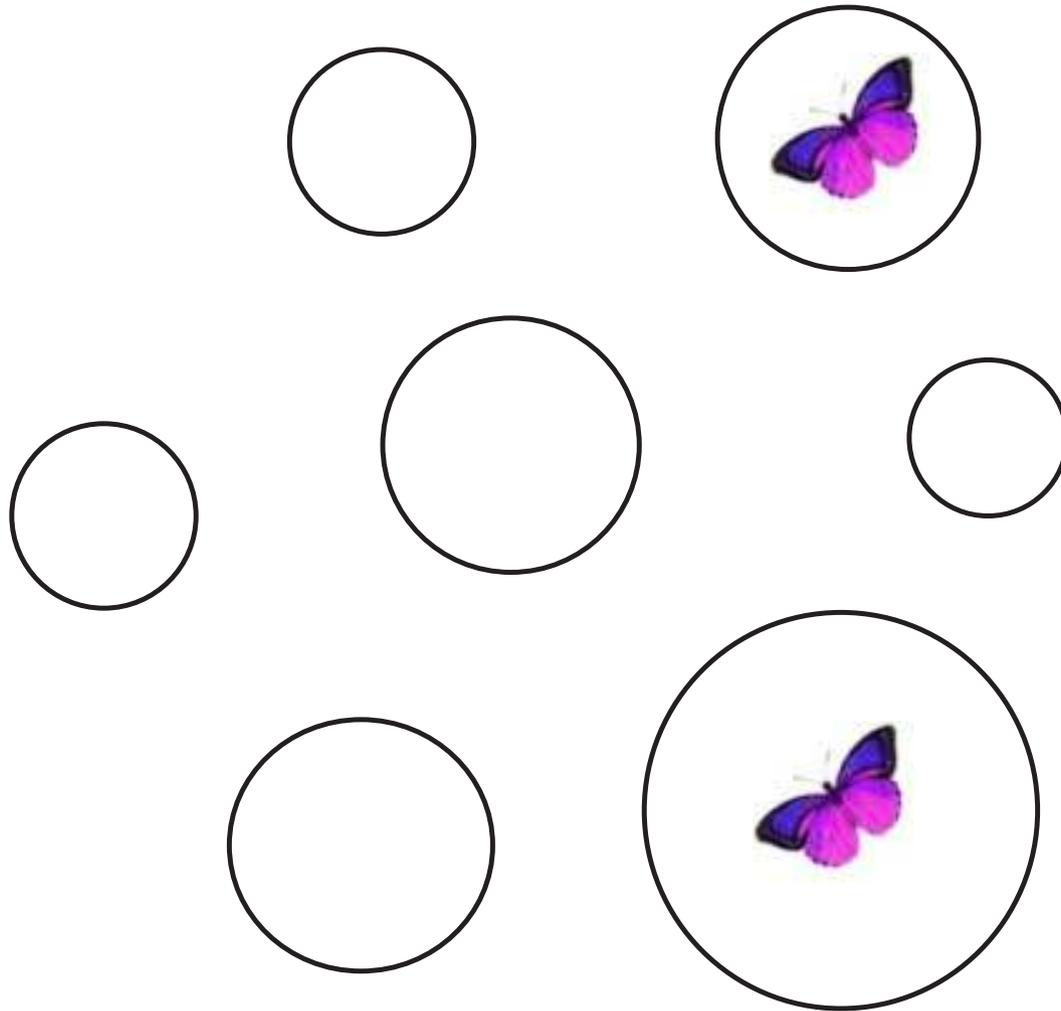
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- Individual patches may suffer local extinction.
- Recolonization can occur through dispersal of individuals from other patches.

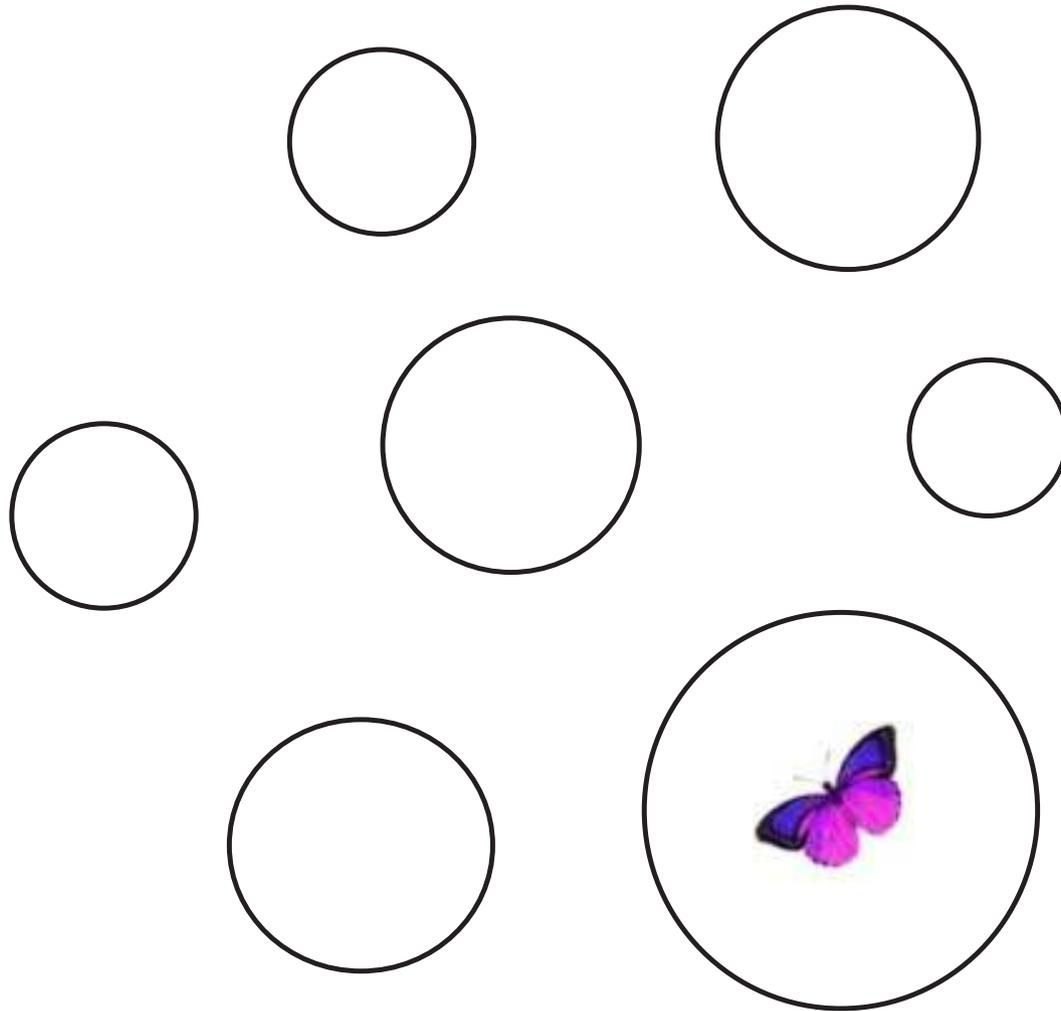
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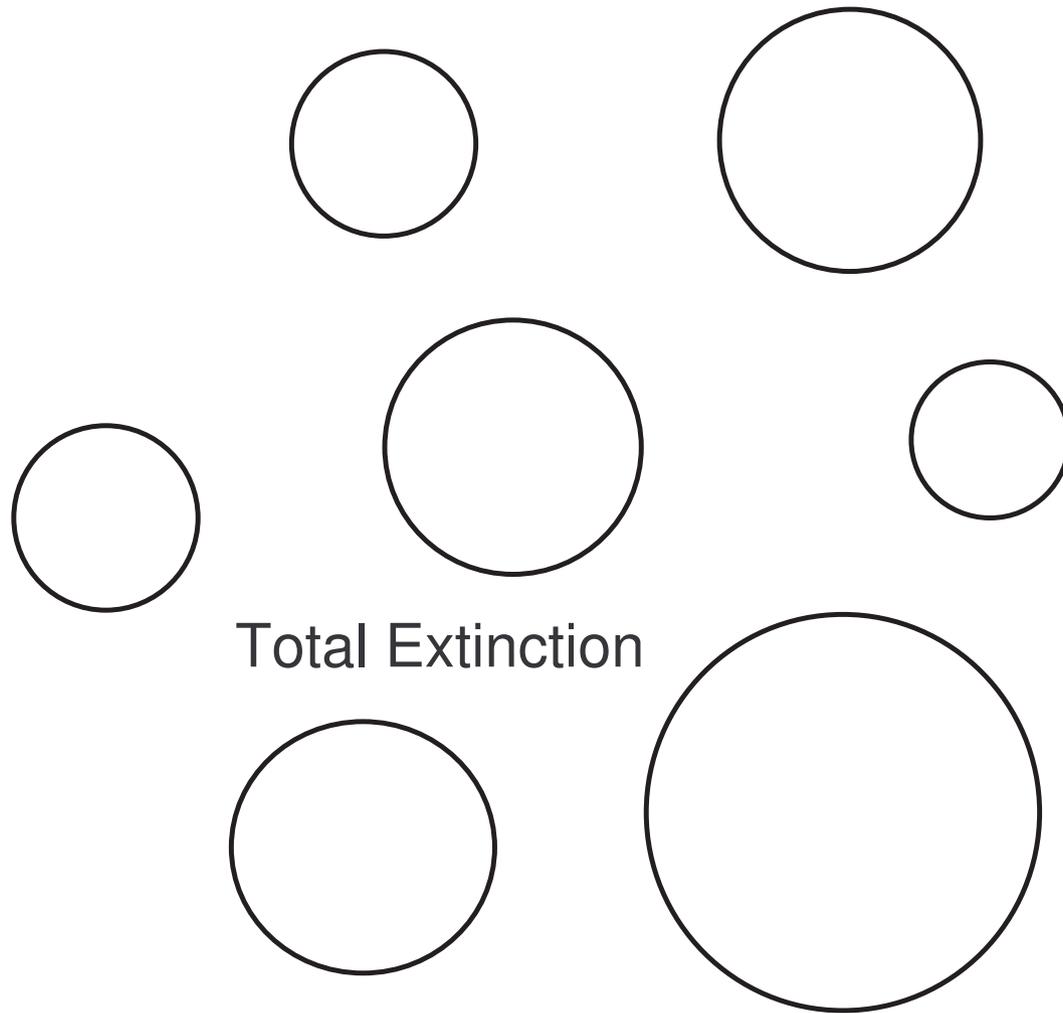
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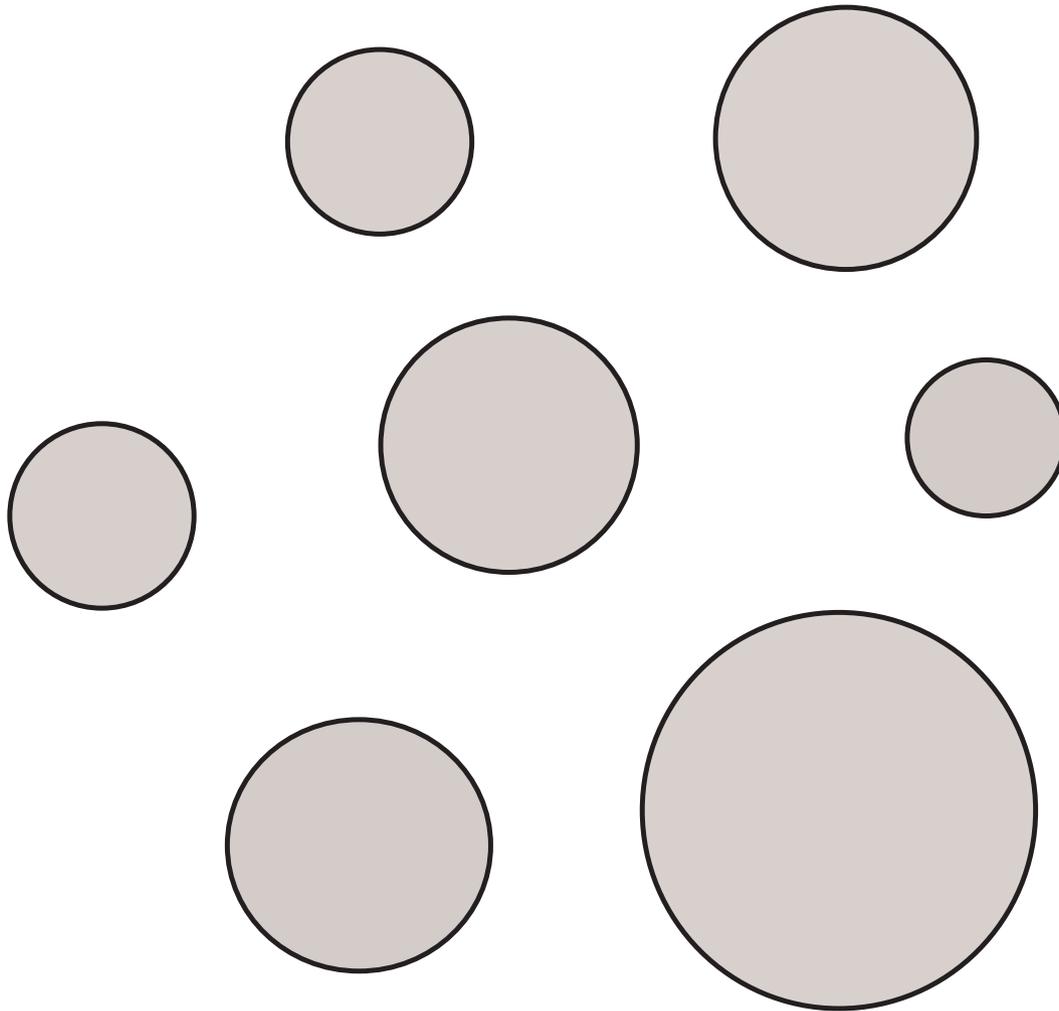


# Metapopulations



Total Extinction

# Metapopulations



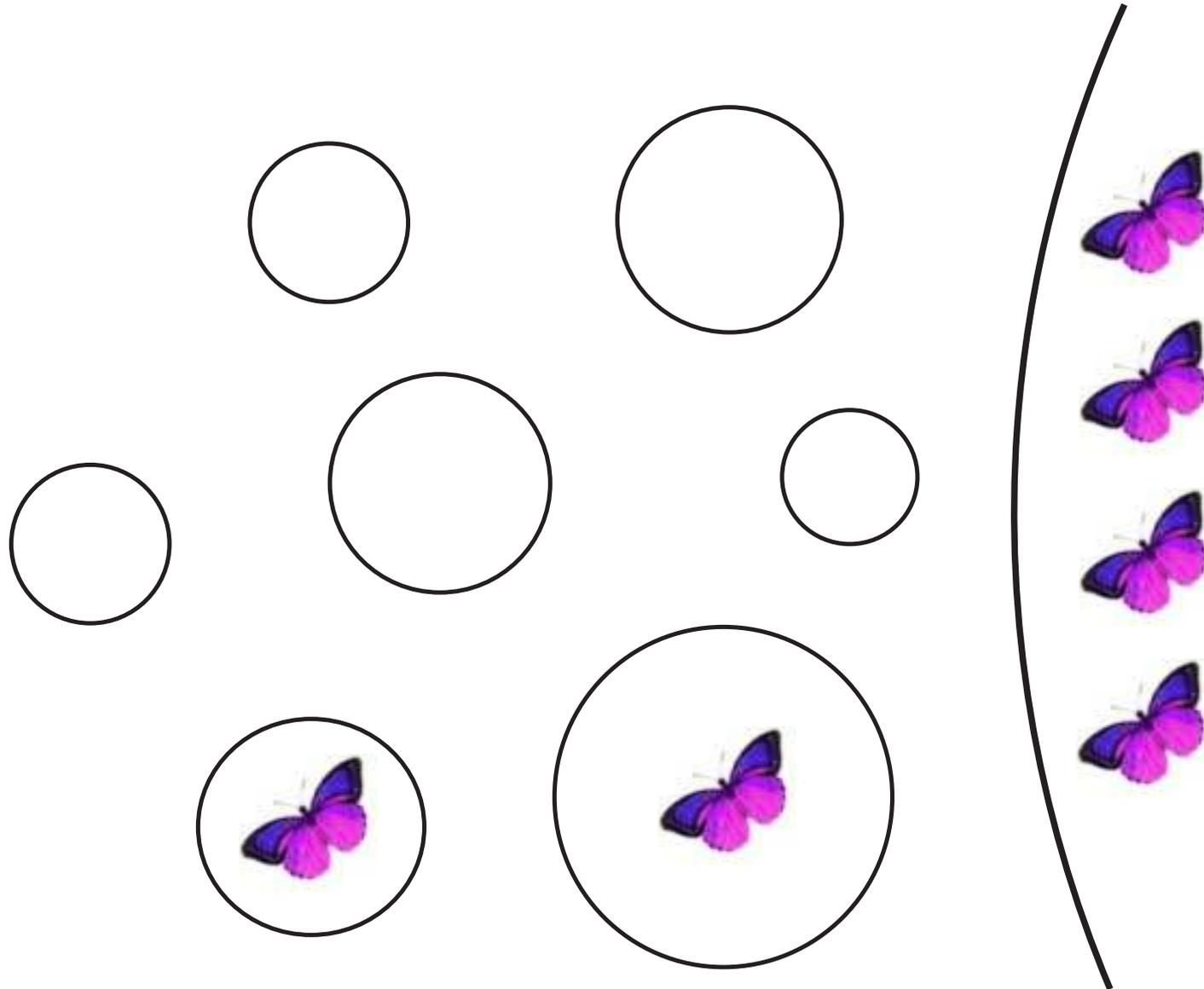
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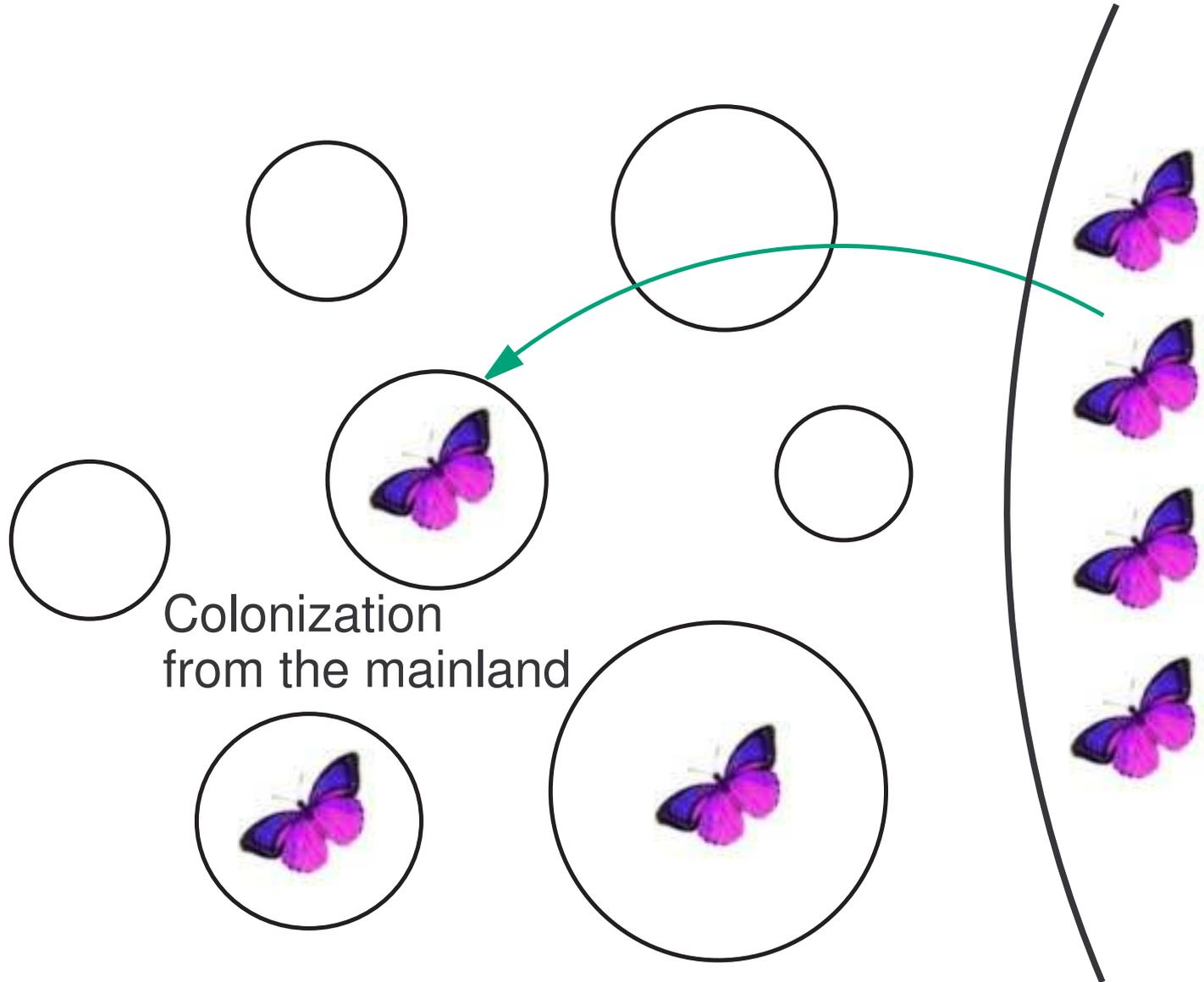
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- In some instances there is an external source of immigration (mainland-island configuration).

# Mainland-island configuration

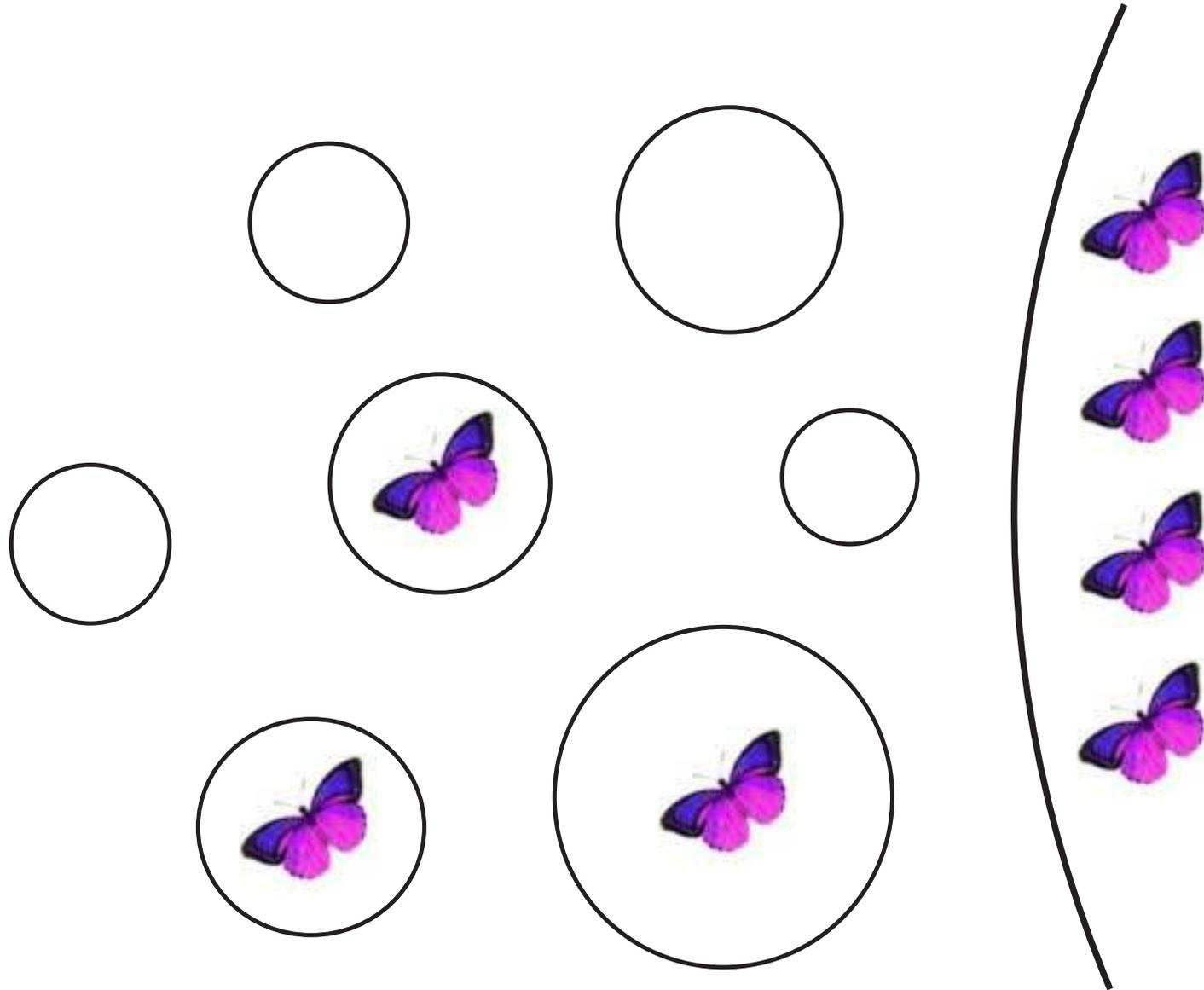


# Mainland-island configuration

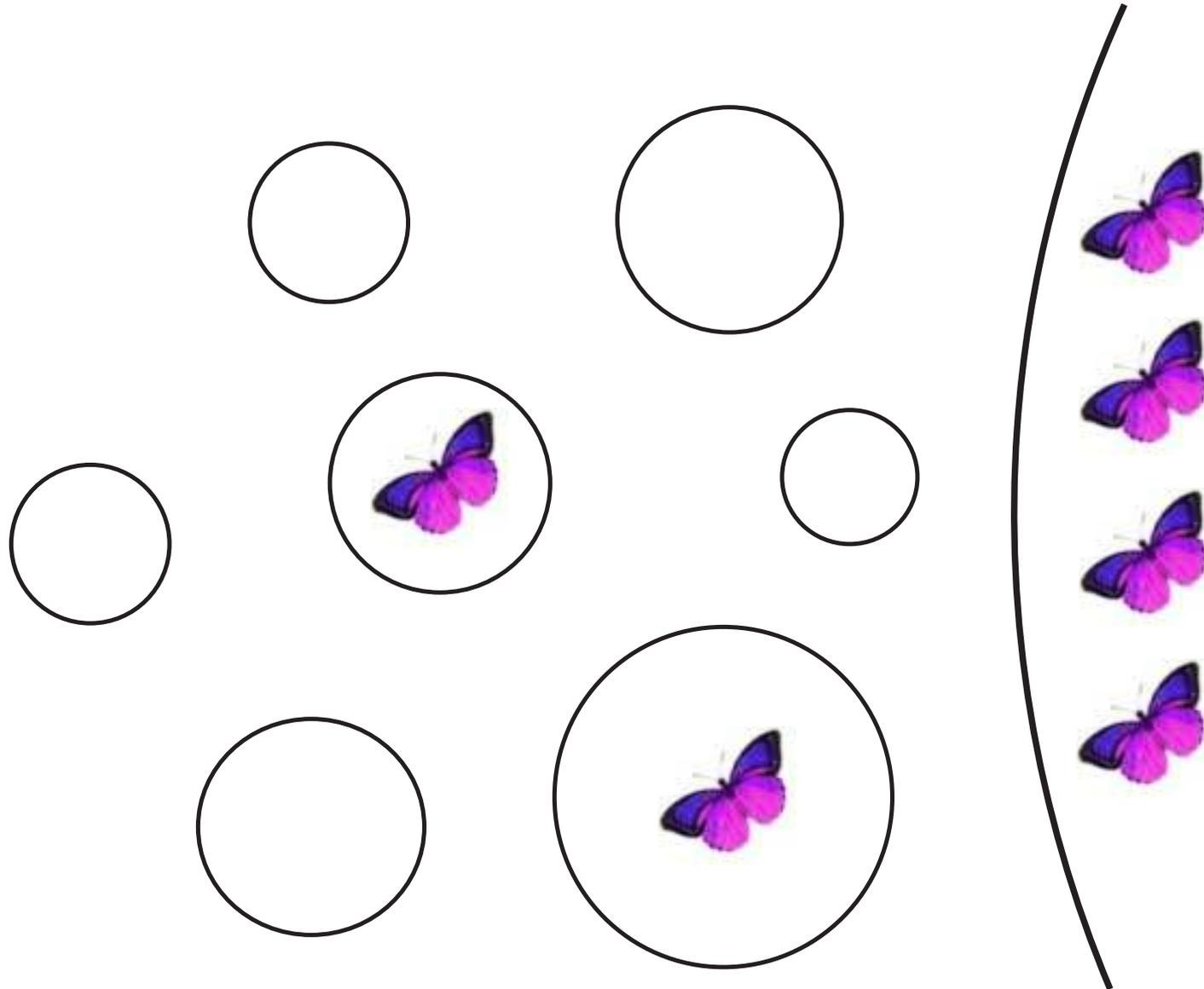


Colonization  
from the mainland

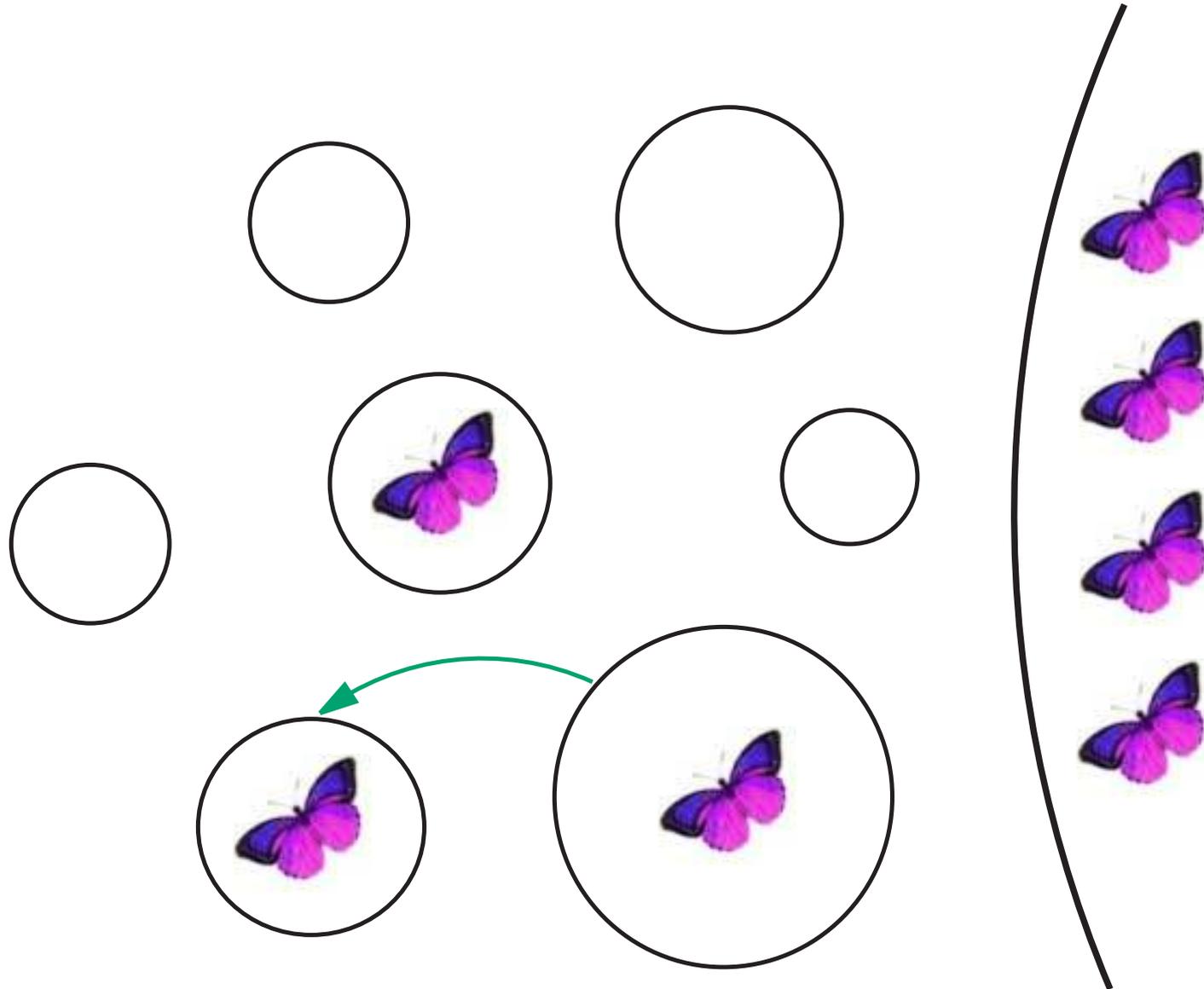
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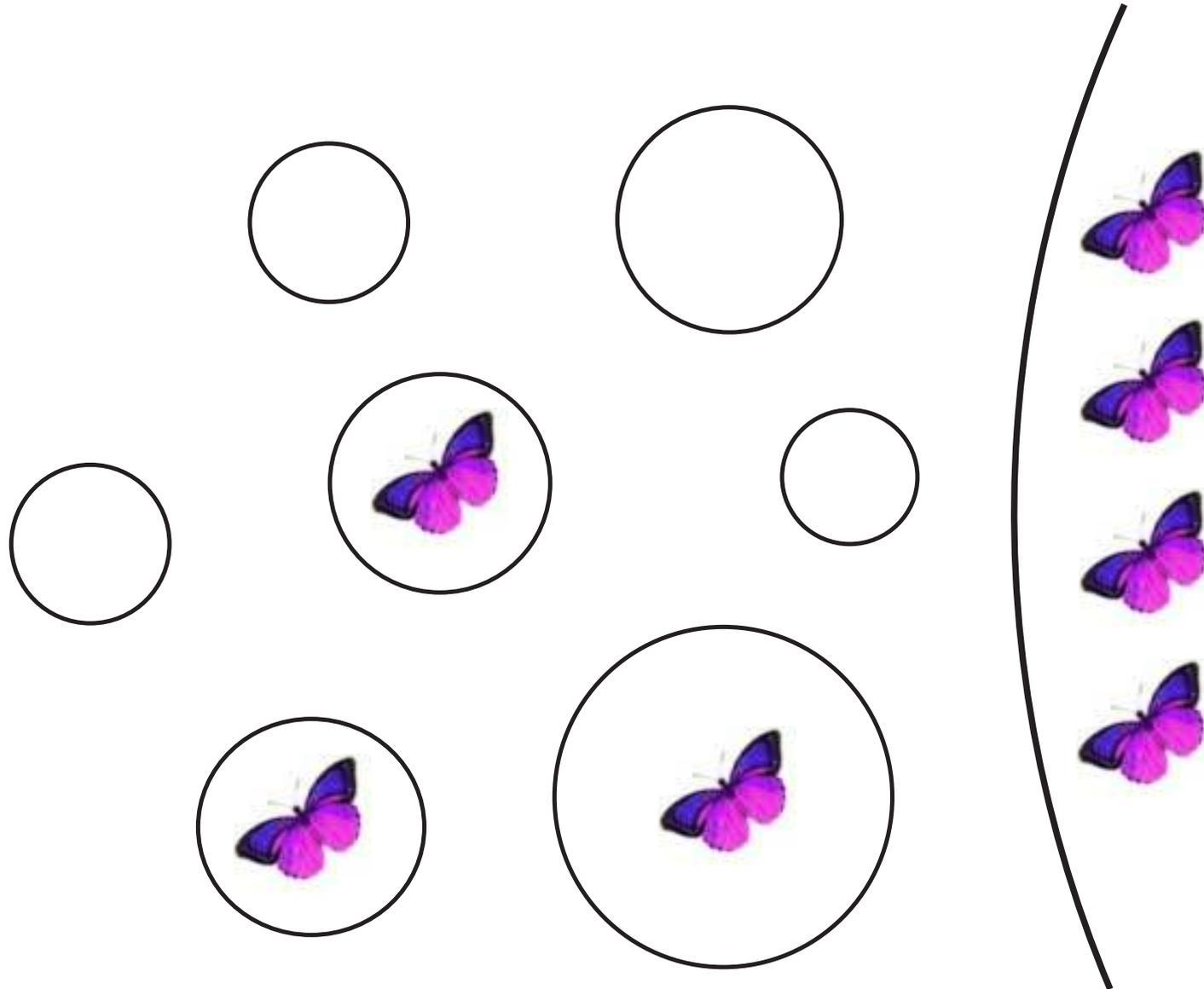
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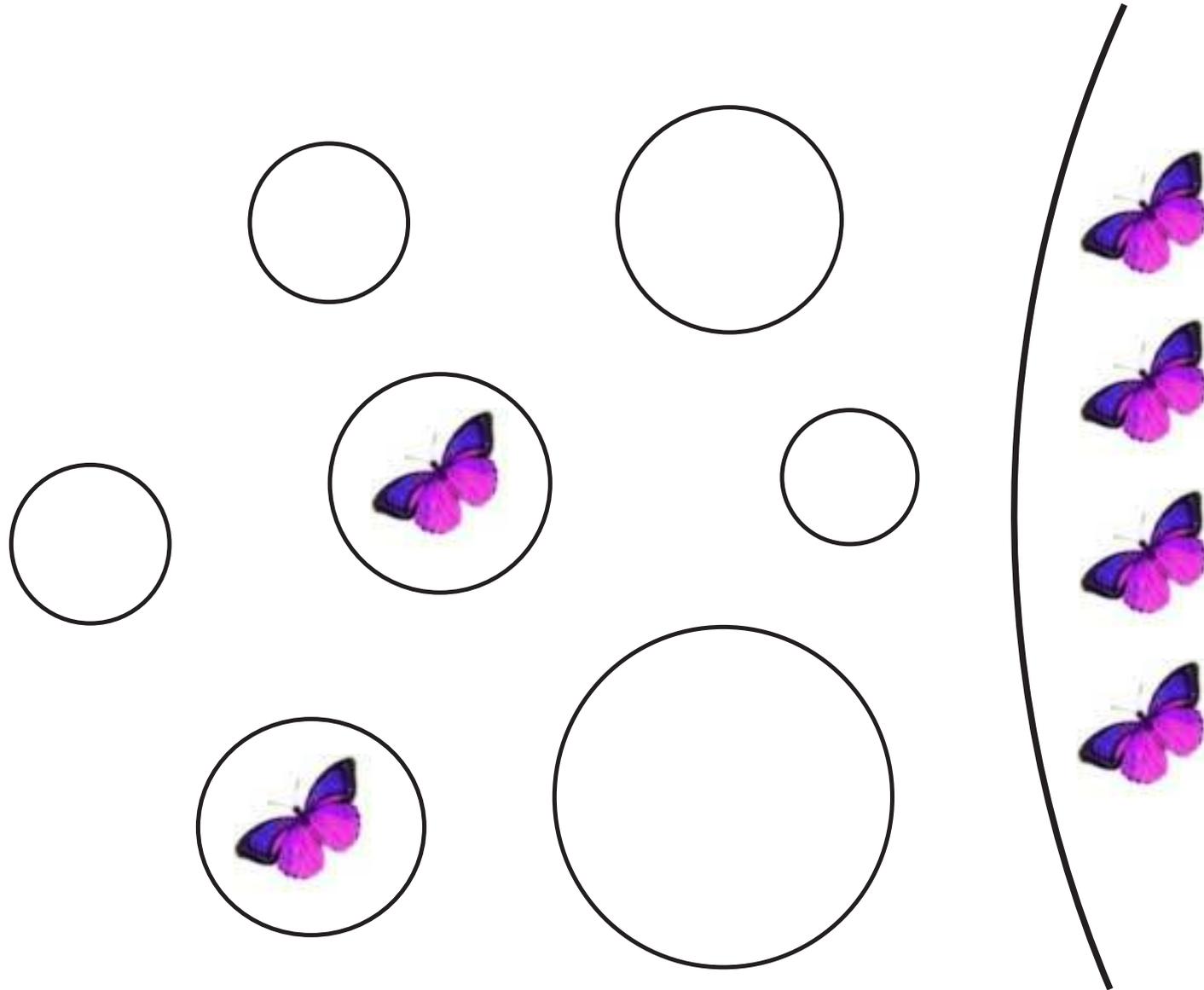
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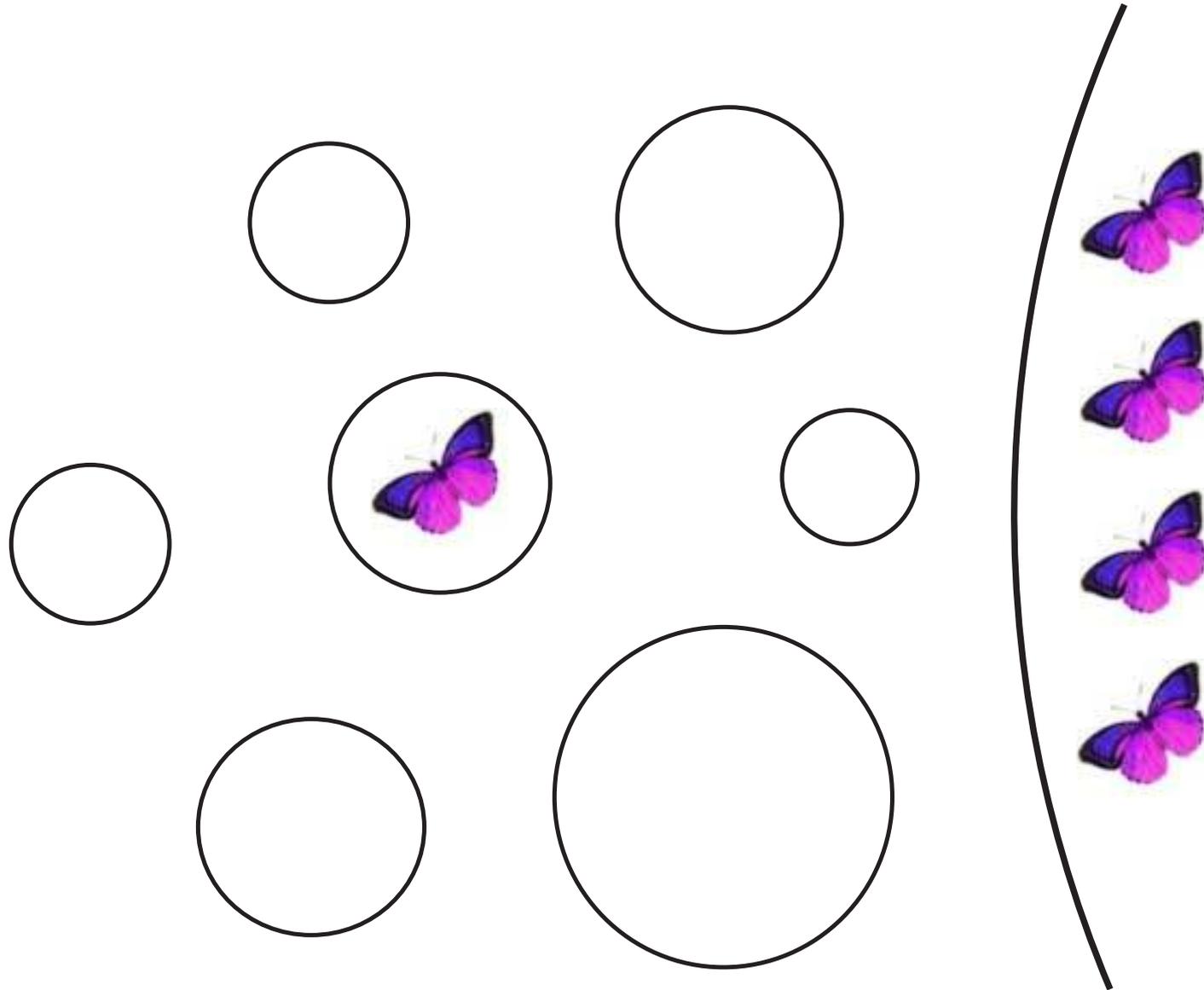
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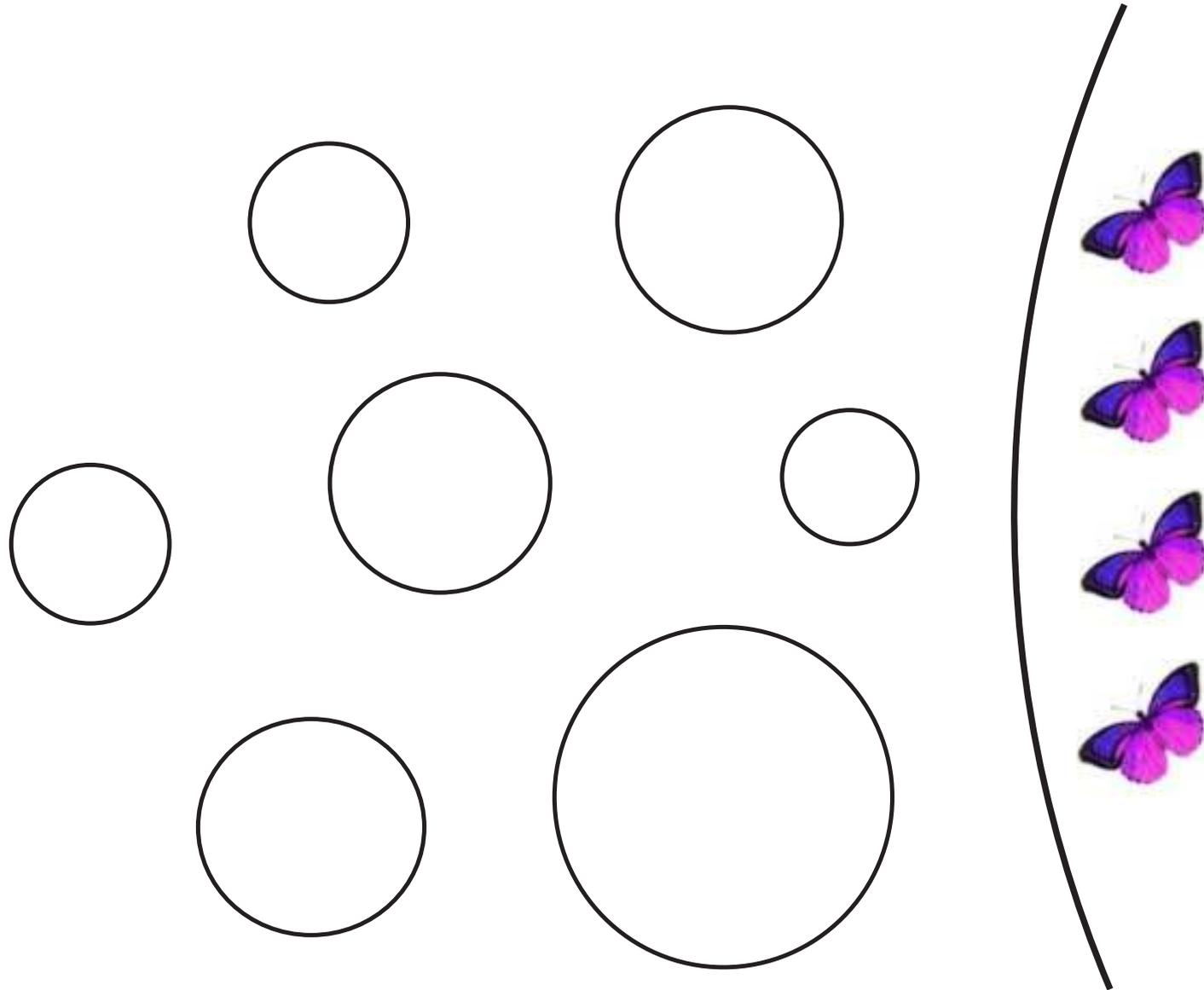
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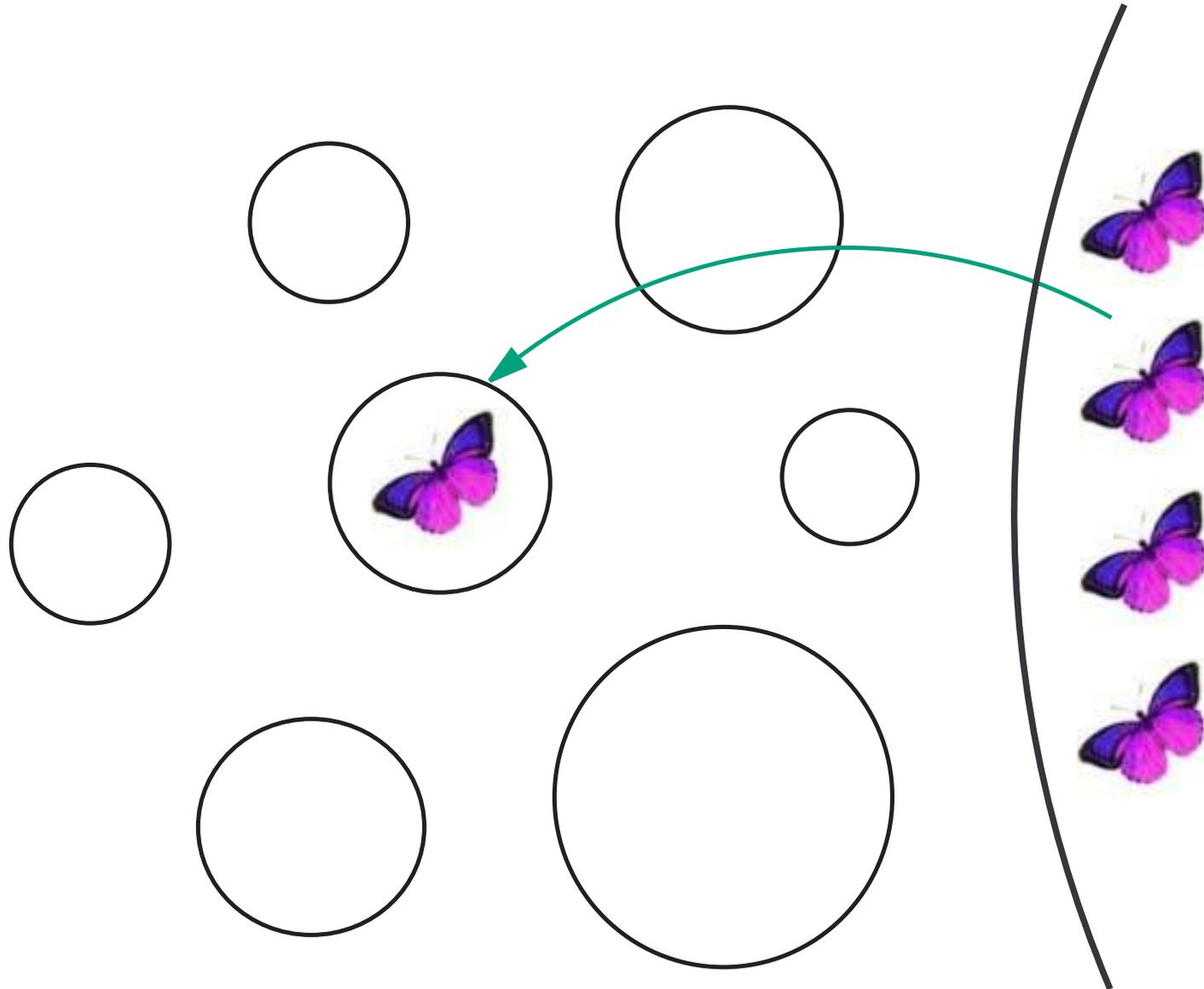
# Mainland-island configuration



# Mainland-island configuration



# Mainland-island configuration



# Typical questions

Given an appropriate model ...

- Assessing population viability:
  - What is the expected time to (total) extinction\* ?
  - What is the probability of extinction by time  $t^*$  ?
- Can we improve population viability ?
- How do we estimate the parameters of the model ?
- Can we determine the stationary/quasi-stationary distributions ?

\*Or *first* total extinction in the mainland-island setup.

# Patch-occupancy models

We record the *number*  $n_t$  of occupied patches at each time  $t$  and suppose that  $(n_t, t \geq 0)$  is a Markov chain in discrete or continuous time.

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In Lecture 1 we looked at the *stochastic logistic (SL) model* of Feller\*.

\*Feller, W. (1939) Die grundlagen der volterraschen theorie des kampfes ums dasein in wahrscheinlichkeitsteoretischer behandlung. Acta Biotheoretica 5, 11–40.

# A continuous-time model

There are  $J$  patches. Each occupied patch becomes empty at rate  $e$  and colonization of empty patches occurs at rate  $c/J$  for each occupied-unoccupied pair.

The state space of the Markov chain  $(n_t, t \geq 0)$  is  $S = \{0, 1, \dots, J\}$  and the transitions are:

$$\begin{array}{lll} n \rightarrow n + 1 & \text{at rate} & \frac{c}{J}n(J - n) \\ n \rightarrow n - 1 & \text{at rate} & en \end{array}$$

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Mainland-island version ( $v$  is the immigration rate):

$$n \rightarrow n + 1 \quad \text{at rate} \quad v(J - n) + \frac{c}{J}n(J - n)$$

$$n \rightarrow n - 1 \quad \text{at rate} \quad en$$

# The SL model

We identified an approximating deterministic model for the *proportion*,  $X_t^{(J)} = n_t/J$ , of occupied patches at time  $t$ . A *functional law of large numbers* established convergence of the family  $(X_t^{(J)})$  to the unique trajectory  $(x_t)$  satisfying

$$x_t' = cx_t(1 - x_t) - ex_t = cx_t(1 - \rho - x_t),$$

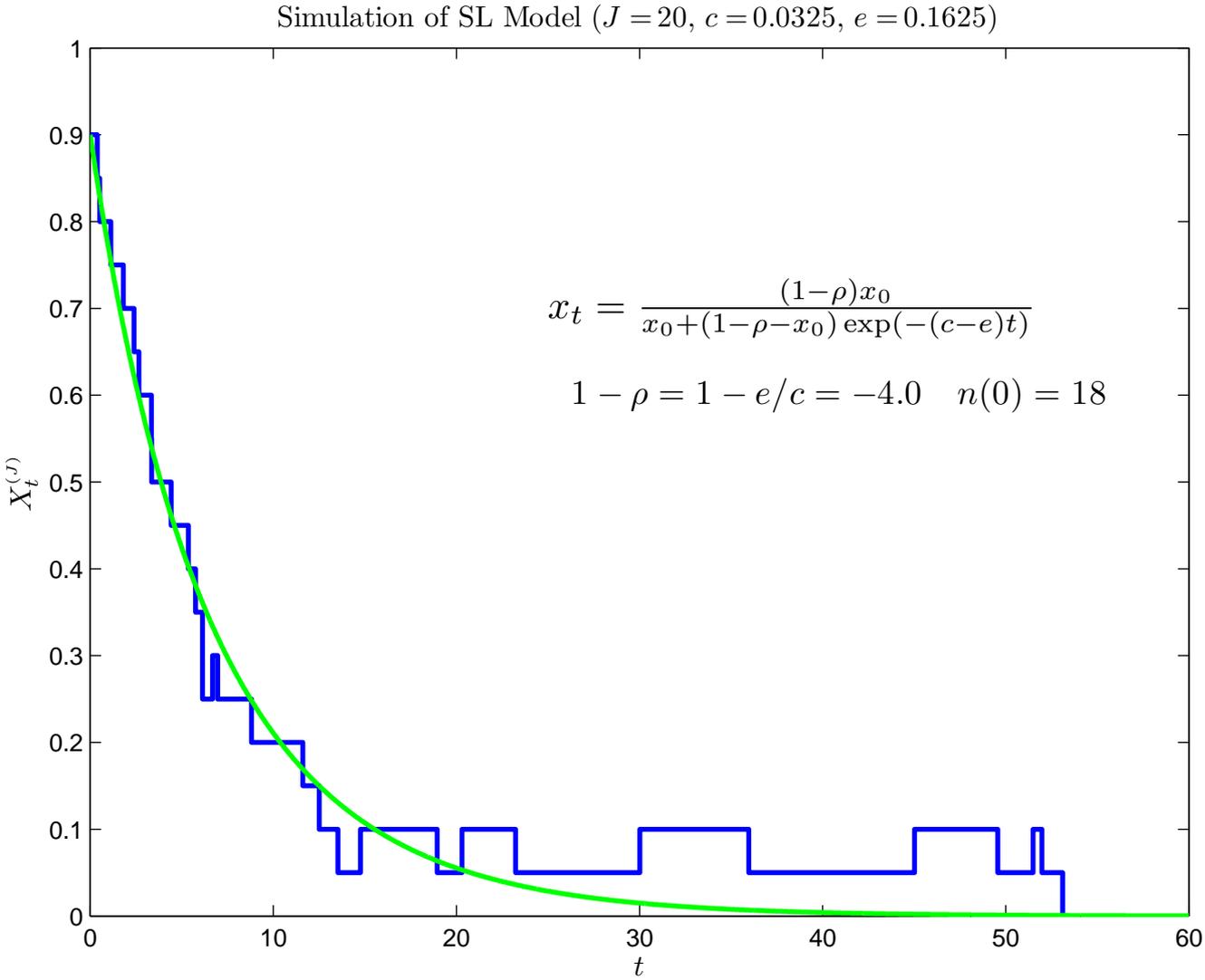
namely

$$x_t = \frac{(1 - \rho)x_0}{x_0 + (1 - \rho - x_0)e^{-(c-e)t}},$$

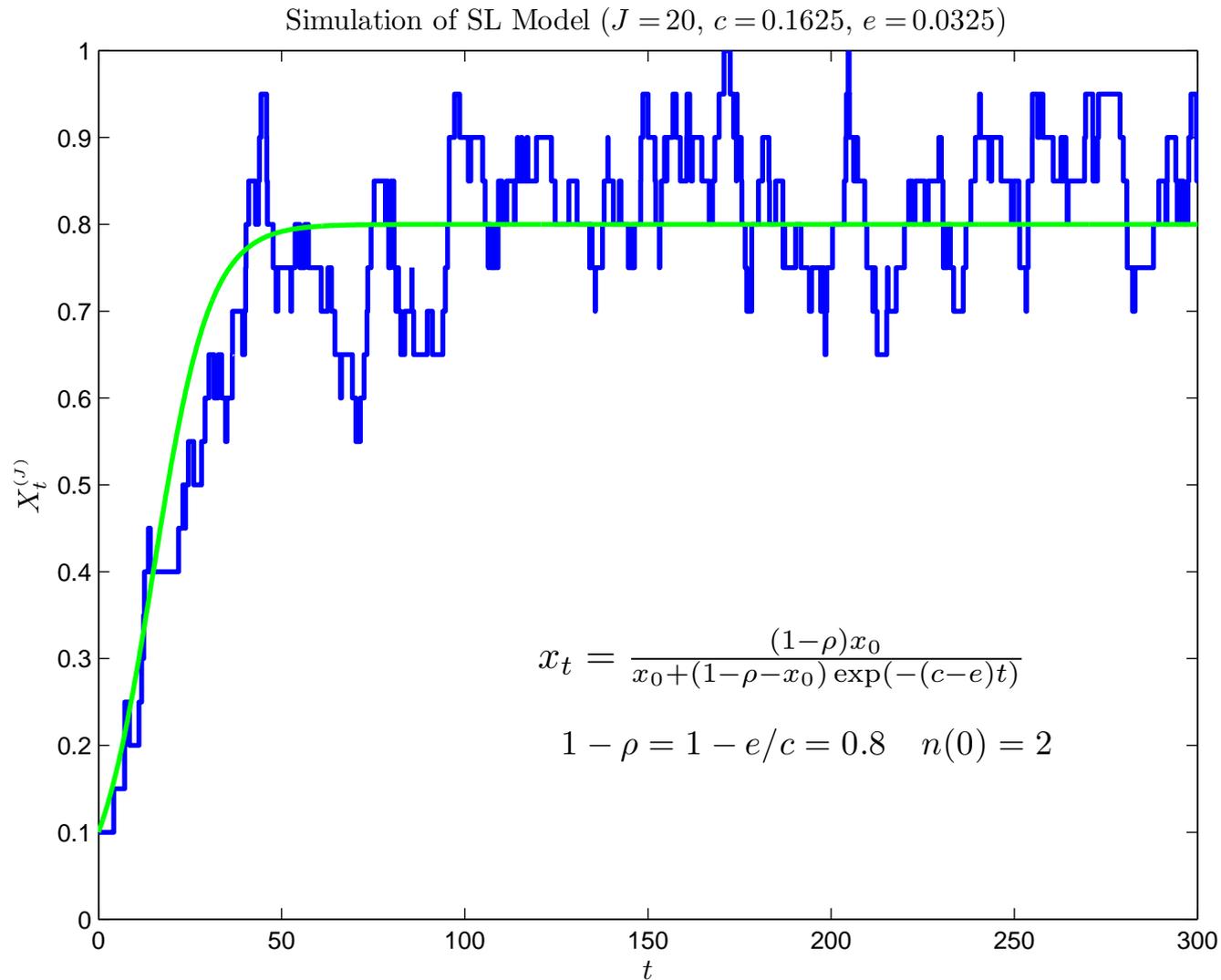
being the classical Verhulst\* model.

\*Verhulst, P.F. (1838) Notice sur la loi que la population suit dans son accroissement. Corr. Math. et Phys. X, 113–121.

# The SL model ( $c < e$ ) $x = 0$ stable



# The SL model ( $c > e$ ) $x = 1 - e/c$ stable

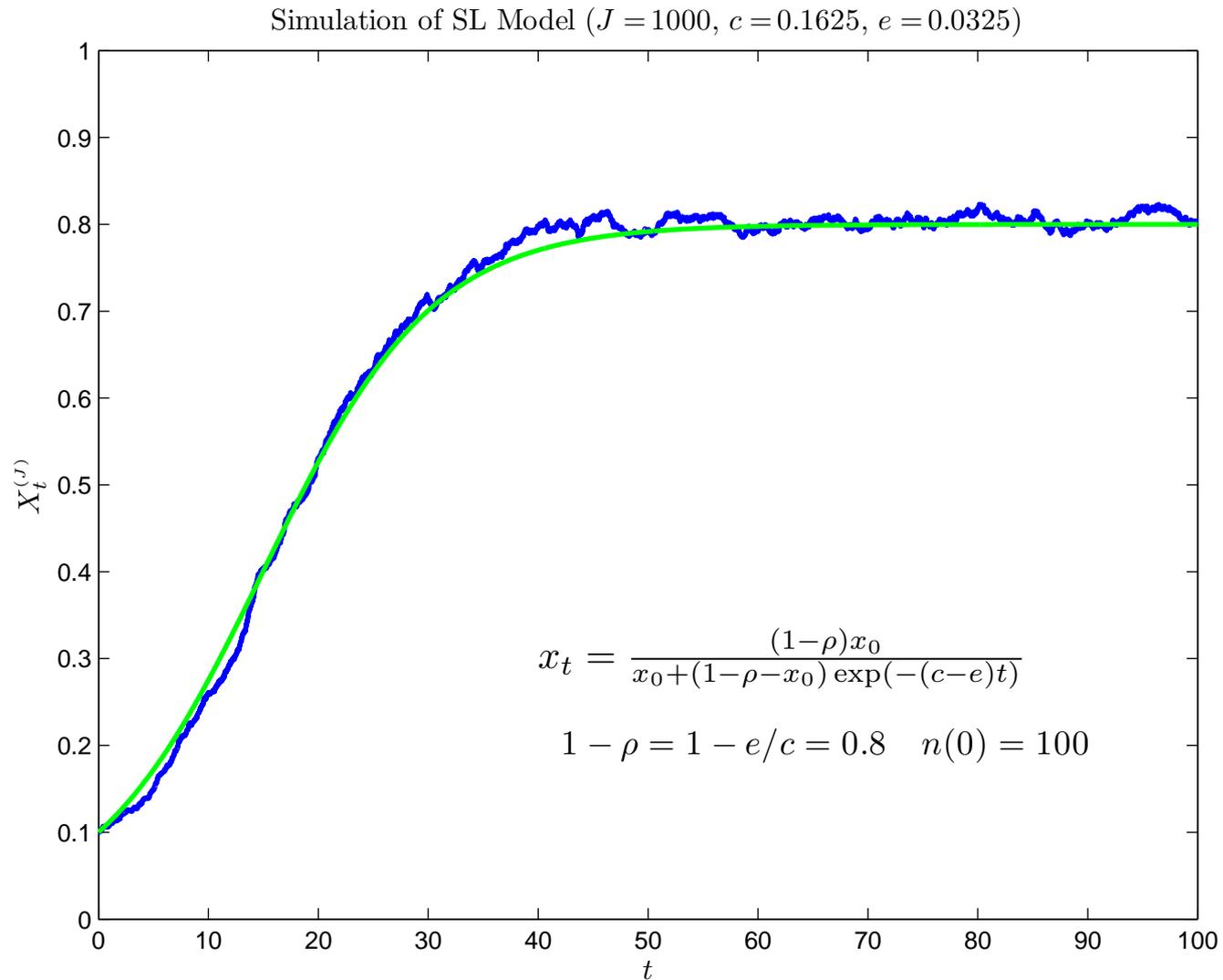


# The SL model

**Theorem** If  $X_0^{(J)} \rightarrow x_0$  as  $J \rightarrow \infty$ , then the family of processes  $(X_t^{(J)})$  converges *uniformly in probability* on *finite time intervals* to the deterministic trajectory  $(x_t)$ : for every  $\epsilon > 0$ ,

$$\lim_{J \rightarrow \infty} \Pr \left( \sup_{s \leq t} |X_s^{(J)} - x_s| > \epsilon \right) = 0.$$

# The SL model ( $c > e$ ) $J \rightarrow \infty$



# Accounting for life cycle

Many species have life cycles (often annual) that consist of distinct phases, and the propensity for colonization and local extinction is different in each phase.

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Many species have life cycles (often annual) that consist of distinct phases, and the propensity for colonization and local extinction is different in each phase. 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



# Butterfly life cycle

Egg  $\simeq$  4 days



Larva (caterpillar)  $\simeq$  14 days



Pupa (chrysalis)  $\simeq$  7 days



Adult (butterfly)  $\simeq$  14 days



# Colonization and extinction phases

Colonization is restricted to the adult phase, and there is a greater propensity for local extinction in the non-adult phases.

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There are several ways to model this:

- A quasi-birth-death process with two phases
- A non-homogeneous continuous-time Markov chain (cycle between two sets of transition rates)
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# A discrete-time Markovian model

Recall that there are  $J$  patches and that  $n_t$  is the number of occupied patches at time  $t$ . We suppose that  $(n_t, t = 0, 1, \dots)$  is a discrete-time Markov chain taking values in  $S = \{0, 1, \dots, J\}$  with a 1-step transition matrix  $P = (p_{ij})$  constructed as follows.

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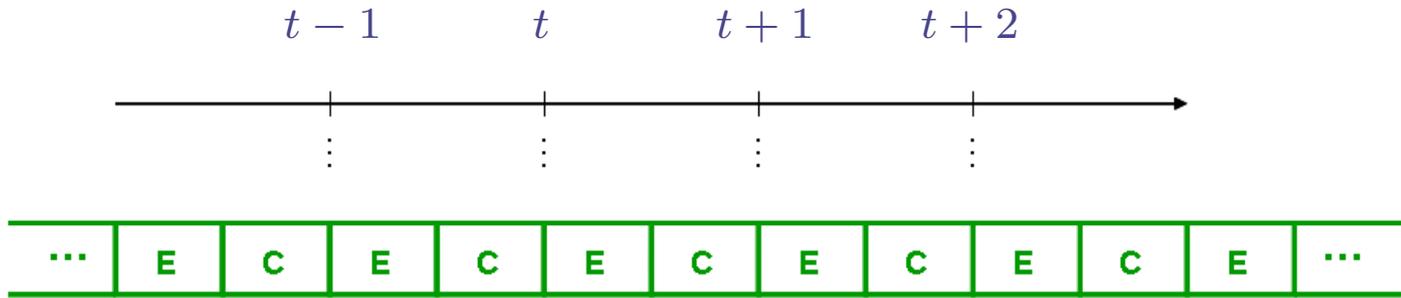
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The extinction and colonization phases are governed by their own transition matrices,  $E = (e_{ij})$  and  $C = (c_{ij})$ .

We let  $P = EC$  if the census is taken after the colonization phase or  $P = CE$  if the census is taken after the extinction phase.

# *EC* versus *CE*

$$P = EC \left\{$$



$$P = CE \left\{$$



# Extinction phase

Suppose that local extinction occurs *at any given patch* with probability  $e$  ( $0 < e < 1$ ), independently of other occupied patches. So, the number of extinctions when there are  $i$  patches occupied has a binomial  $Bin(i, e)$  distribution, and therefore

$$e_{i,i-k} = \binom{i}{k} e^k (1 - e)^{i-k} \quad (k = 0, 1, \dots, i).$$

We also have  $e_{ij} = 0$  if  $j > i$ .

# Colonization phase

Suppose that colonization occurs according to the following mechanism.

If there are  $i$  occupied patches, then each unoccupied patch is colonized with probability  $c_i = (i/J)c$ , where  $c \in (0, 1]$  is a *fixed maximum colonization potential*, the (hypothetical) probability that a single unoccupied patch is colonized by the fully occupied network.

So, the unoccupied patches are colonized independently with the same probability, this probability being *proportional to* the number of patches with the potential to colonize.

# Colonization phase

Therefore, the number of colonizations when there are  $i$  patches occupied has a binomial  $Bin(J - i, c_i)$  distribution, and so

$$c_{i,i+k} = \binom{J-i}{k} c_i^k (1 - c_i)^{J-i-k}, \quad (k = 0, 1, \dots, J - i),$$

In particular,  $c_{0j} = \delta_{0j}$ . We also have  $c_{ij} = 0$ , for  $j < i$ .

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Notice that 0 is an absorbing state and  $C = \{1, \dots, J\}$  is a communicating class.

There are other sensible choices for  $c_i$ : for example  $c_i = c(1 - (1 - c_1/c)^i)$  or  $c_i = 1 - \exp(-i\beta/J)$ .

# Evaluation of $P$

We can evaluate  $P$  elementwise as follows.

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$$p_{ij} = \sum_{k=1}^{\min\{i,j\}} \binom{i}{k} (1-e)^k e^{i-k} \binom{J-k}{j-k} c_k^{j-k} (1-c_k)^{J-j}.$$

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In particular, for  $i \geq 1$ ,  $p_{i0} = e^i (1 - c_i (1 - e))^{J-i}$ .

# Equivalent independent phases

For the  $CE$ -model,

$$\mathbf{E}(z^{n_{t+1}} | n_t = i) = (e + (1 - e)z)^i (1 - (1 - e)c_i(1 - z))^{J-i}.$$

Thus, given  $n_t = i$ ,  $n_{t+1}$  has the *same distribution* as  $B_1 + B_2$ , where  $B_1$  and  $B_2$  are two *independent* random variables with  $B_1 \sim \mathbf{Bin}(i, 1 - e)$  and  $B_2 \sim \mathbf{Bin}(J - i, (1 - e)c_i)$ .

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It is as if each of the  $i$  occupied patches remains occupied with probability  $1 - e$  and each of the  $J - i$  unoccupied patches becomes occupied with probability  $(1 - e)c_i$ , all  $J$  patches being affected independently.

# Equivalent independent phases

For the  $EC$ -model, the best we can do is

$$\mathbf{E}(z^{n_{t+1}} | n_t = i) = \mathbf{E} \left\{ z^B (1 - c_B(1 - z))^{J-B} \right\},$$

where  $B \sim \mathit{Bin}(i, 1 - e)$ .

# Large- $J$

However, note the large- $J$  asymptotics when  $c_i = ic/J$ .  
Write  $p_i^{(J)}(z) = \mathbf{E}(z^{n_{t+1}} | n_t = i)$ .

For the  $CE$ -model,

$$\lim_{J \rightarrow \infty} p_i^{(J)}(z) = [e + (1 - e)z \exp(-c(1 - e)(1 - z))]^i.$$

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**Branching!**

# Infinitely many patches

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The effect is ...

**Theorem** Both infinite patch models are Galton-Watson branching processes.

# Infinitely many patches - branching

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For the *EC*-model,  $p_{10} = e$  and

$$p_{1j} = (1 - e) \exp(-c) \frac{c^{j-1}}{(j-1)!} \quad (j \geq 1),$$

the interpretation being that each individual “dies” with probability  $e$  or otherwise is *joined by* a random number of new offspring that follows a *Poisson*( $c$ ) law.

# Infinitely many patches - branching

For the  $CE$ -model,  $p_{10} = e \exp(-c(1 - e))$  and

$$p_{1j} = (1 - e) \exp(-c(1 - e)) \frac{(c(1 - e))^{j-1}}{(j - 1)!} + e \exp(-c(1 - e)) \frac{(c(1 - e))^j}{j!} \quad (j \geq 1).$$

The individual survives with probability  $1 - e$  or dies with probability  $e$ , and there is a random number of *new* offspring that follows a  $Poisson(c(1 - e))$  law.

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*We can now invoke the encyclopaedic theory of branching processes.*

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Notice that  $\sigma_{EC}^2 - \sigma_{CE}^2 = c(2 + c)e(1 - e) > 0$ .

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Notice that  $\sigma_{EC}^2 - \sigma_{CE}^2 = c(2 + c)e(1 - e) > 0$ .

Recall that, given  $n_0 = i$ ,  $\mathbf{E}(n_t) = i\mu^t$  and

$$\text{Var}(n_t) = \begin{cases} i\sigma^2 t & \text{if } \mu = 1 \quad (e = c/(1 + c)) \\ i\sigma^2(\mu^t - 1)\mu^{t-1}/(\mu - 1) & \text{if } \mu \neq 1 \quad (e \neq c/(1 + c)). \end{cases}$$

# Infinitely many patches - total extinction

**Theorem** For both models extinction occurs with probability 1 if and only if  $e \geq c/(1 + c)$ ; otherwise the extinction probability  $\eta$  is the unique solution to  $s = p(s)$  on the interval  $(0, 1)$ , where:

$$EC\text{-model: } p(s) = e + (1 - e)s \exp(-c(1 - s))$$

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And much more ...

- The expected time to extinction.
- Yaglom's theorem on limiting-conditional (quasi-stationary) distributions.

# Back to the $J$ -patch models

Recall that ...

In the extinction phase the number of extinctions when there are  $i$  patches occupied follows a  $Bin(i, e)$  law.

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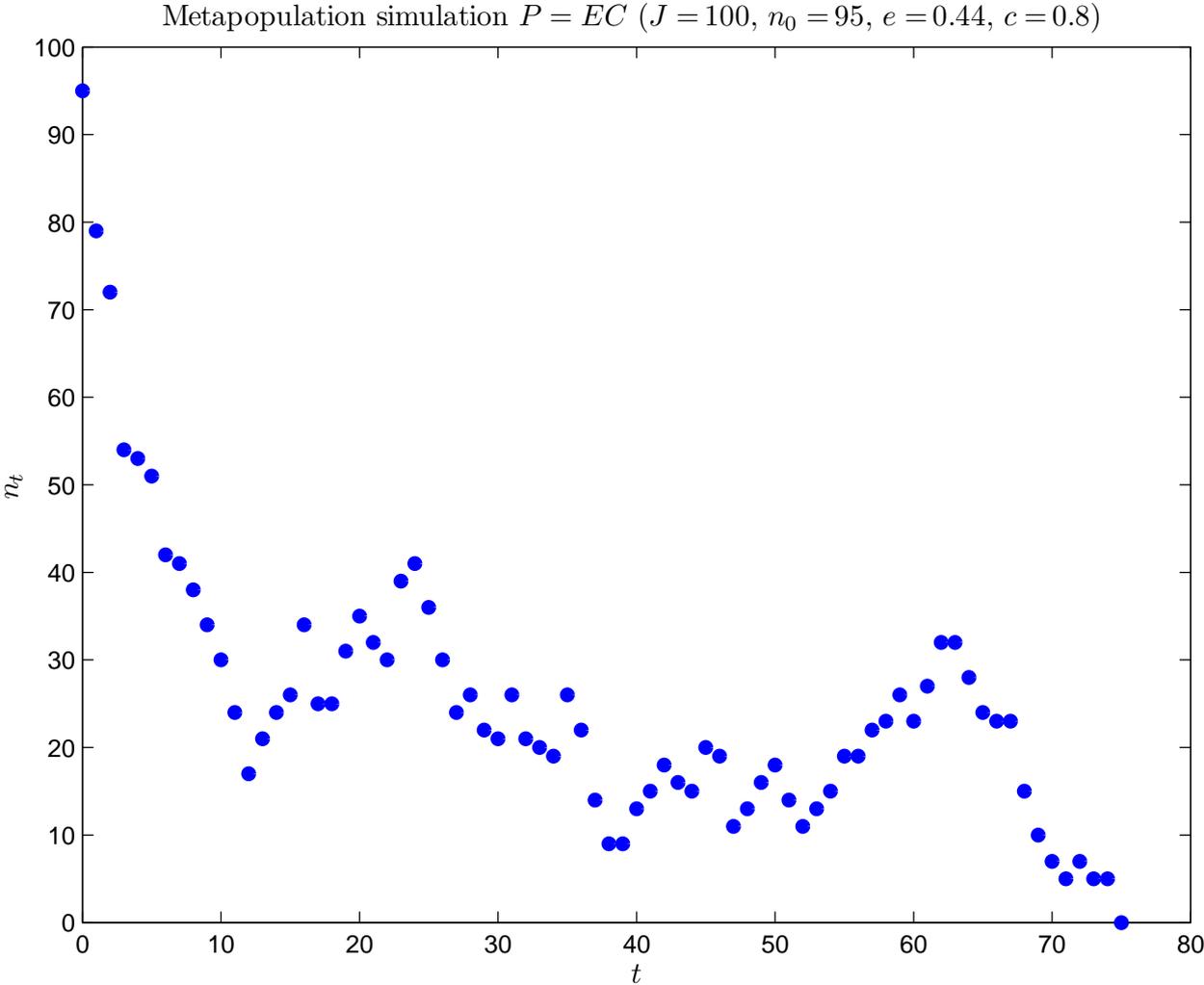
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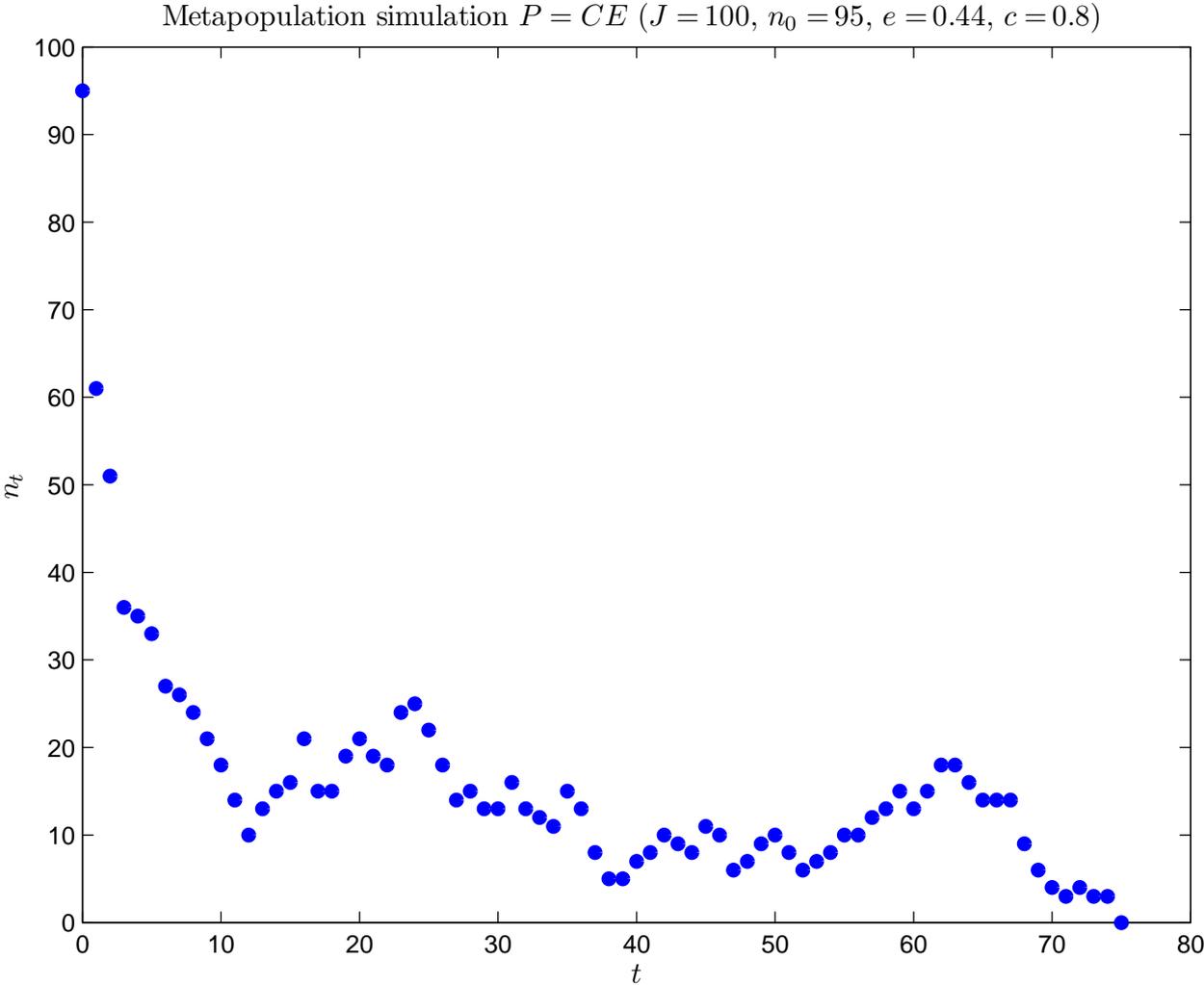
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Numerical procedures are routine.

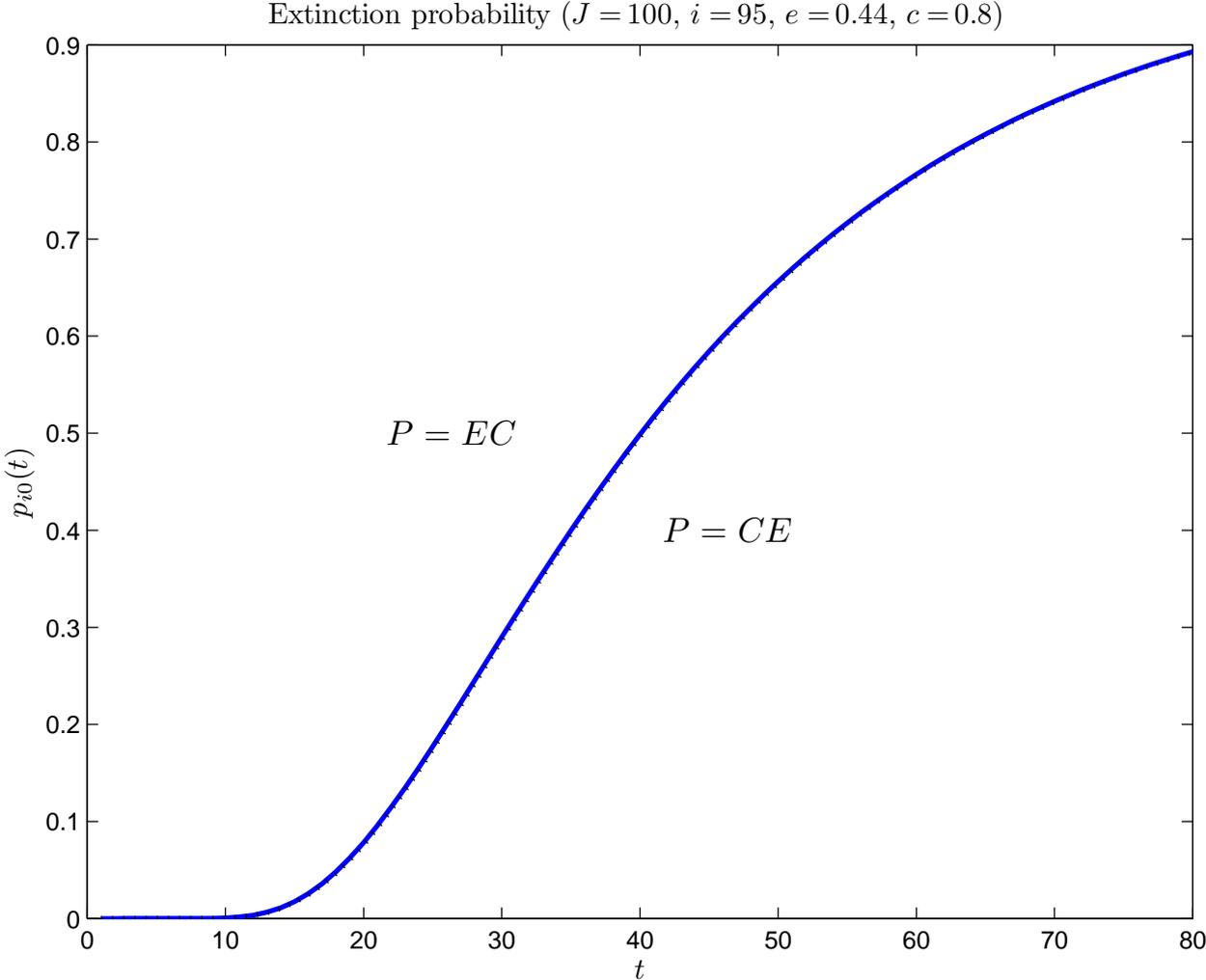
# Simulation: $P = EC$



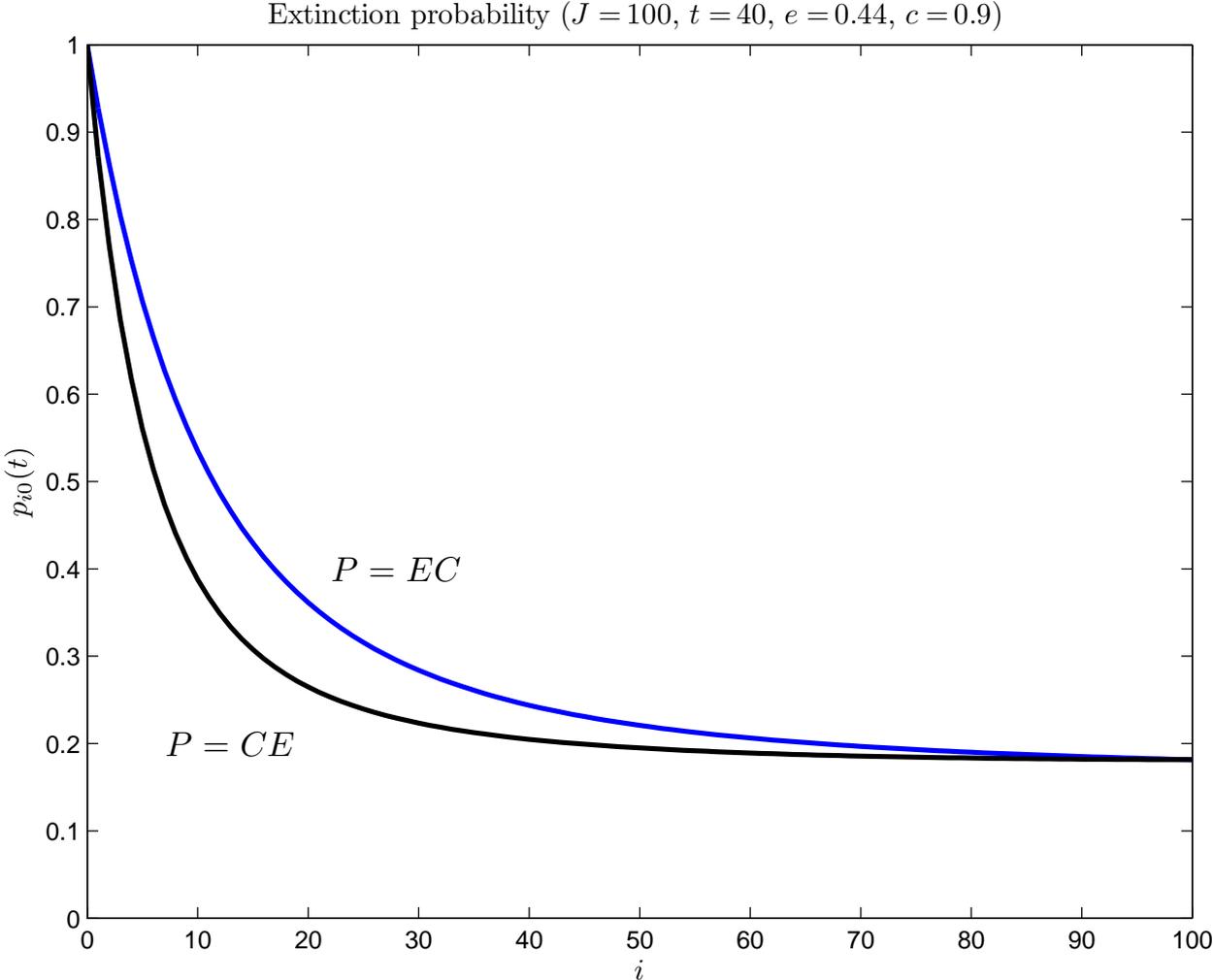
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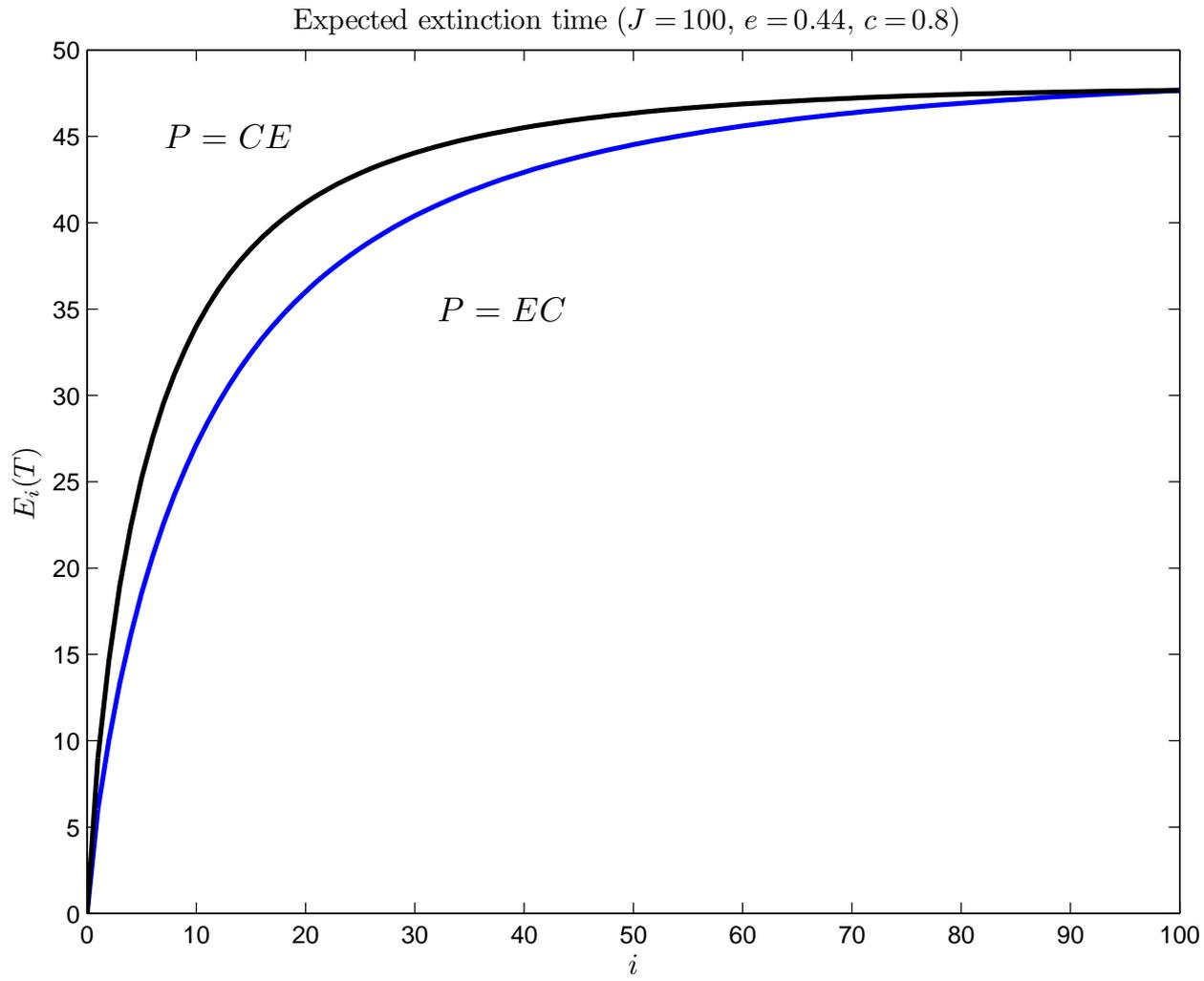
# Extinction probability: vary $t$



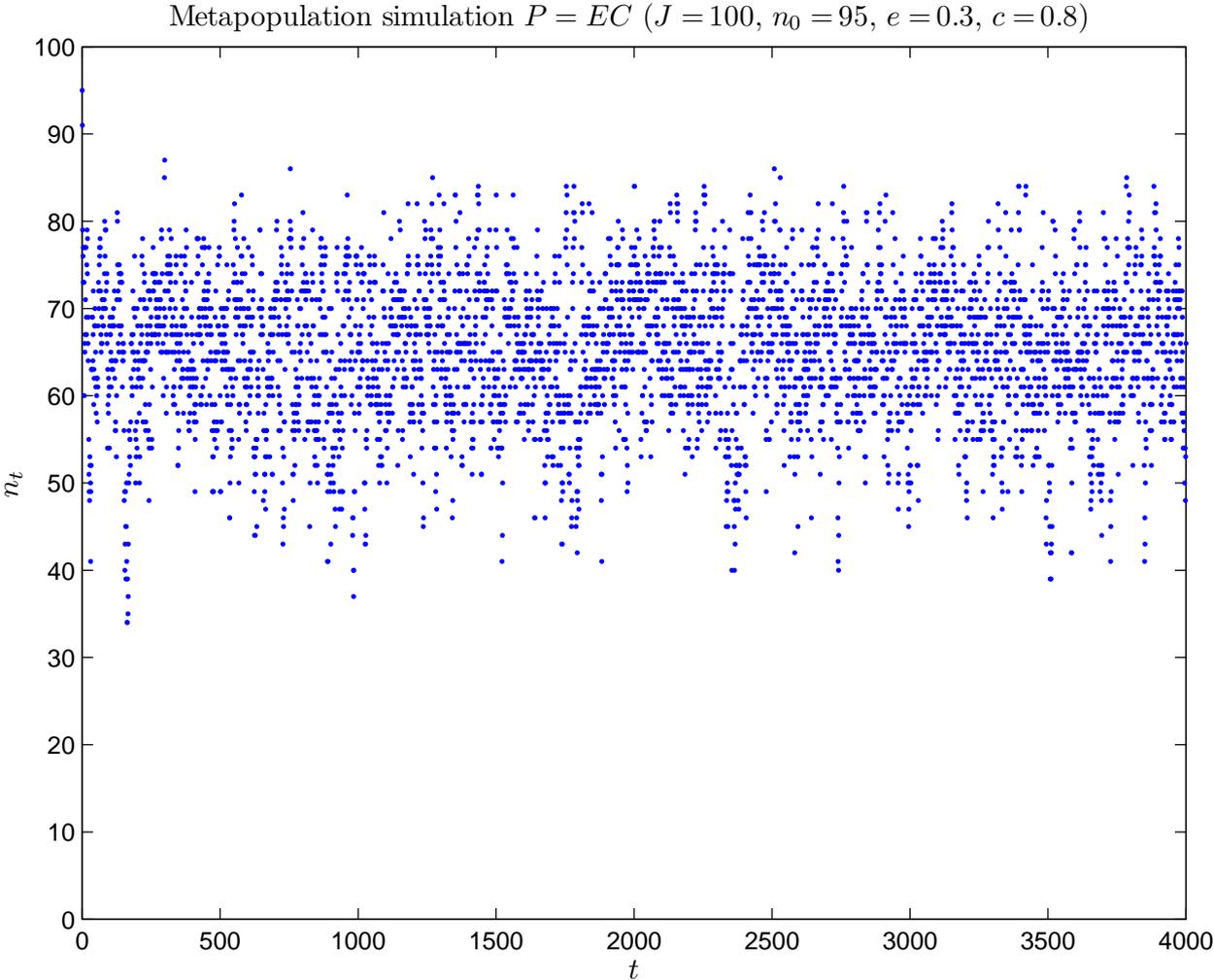
# Extinction probability: vary $n_0$



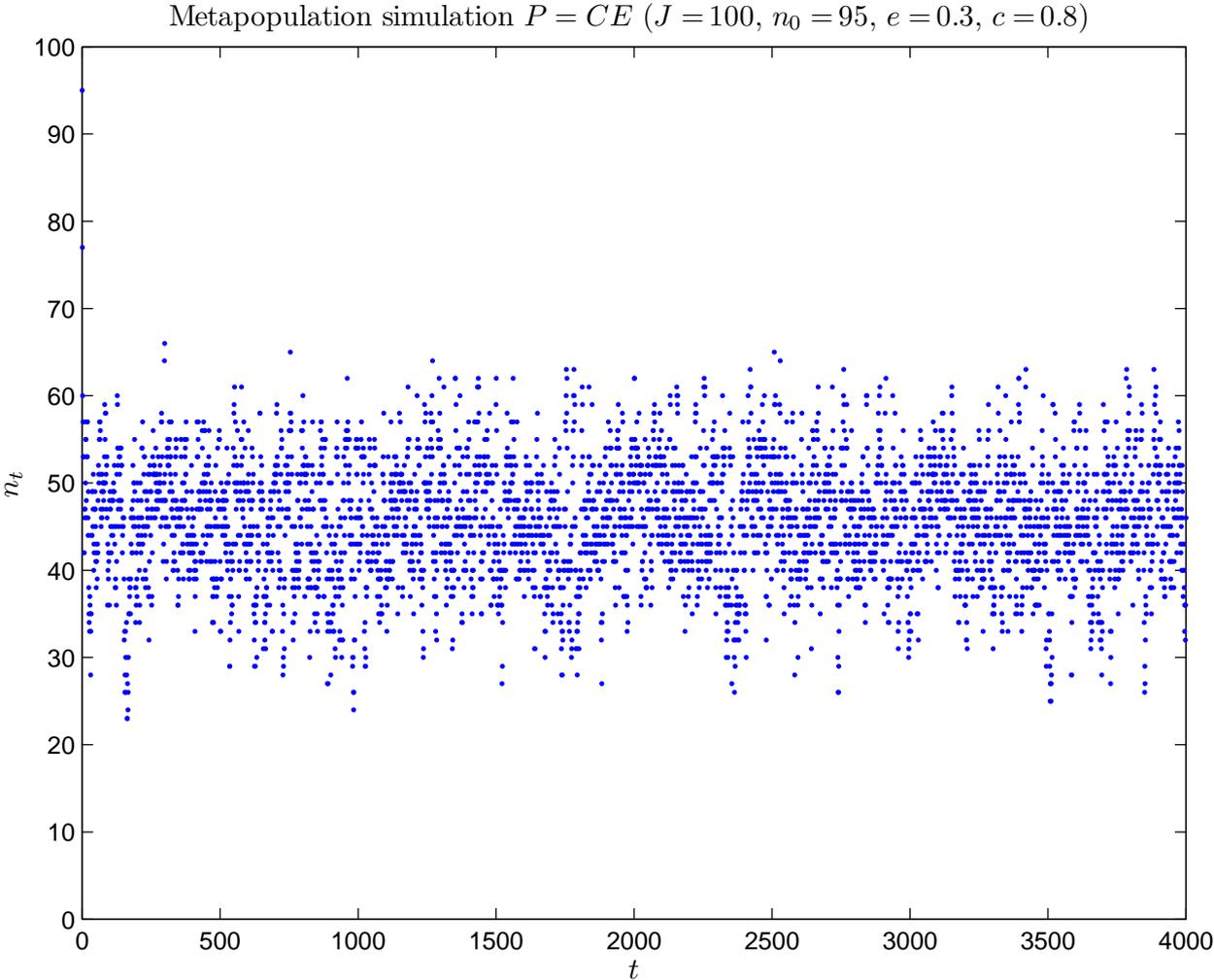
# Expected extinction time: vary $n_0$



# Simulation: $P = EC$



# Simulation: $P = CE$



# Quasi stationarity

We can model this behaviour using a *limiting conditional distribution* (lcd)  $(m_j, j = 1, \dots, J)$ ; often called a *quasi-stationary distribution* (qsd)\*.

lcd:

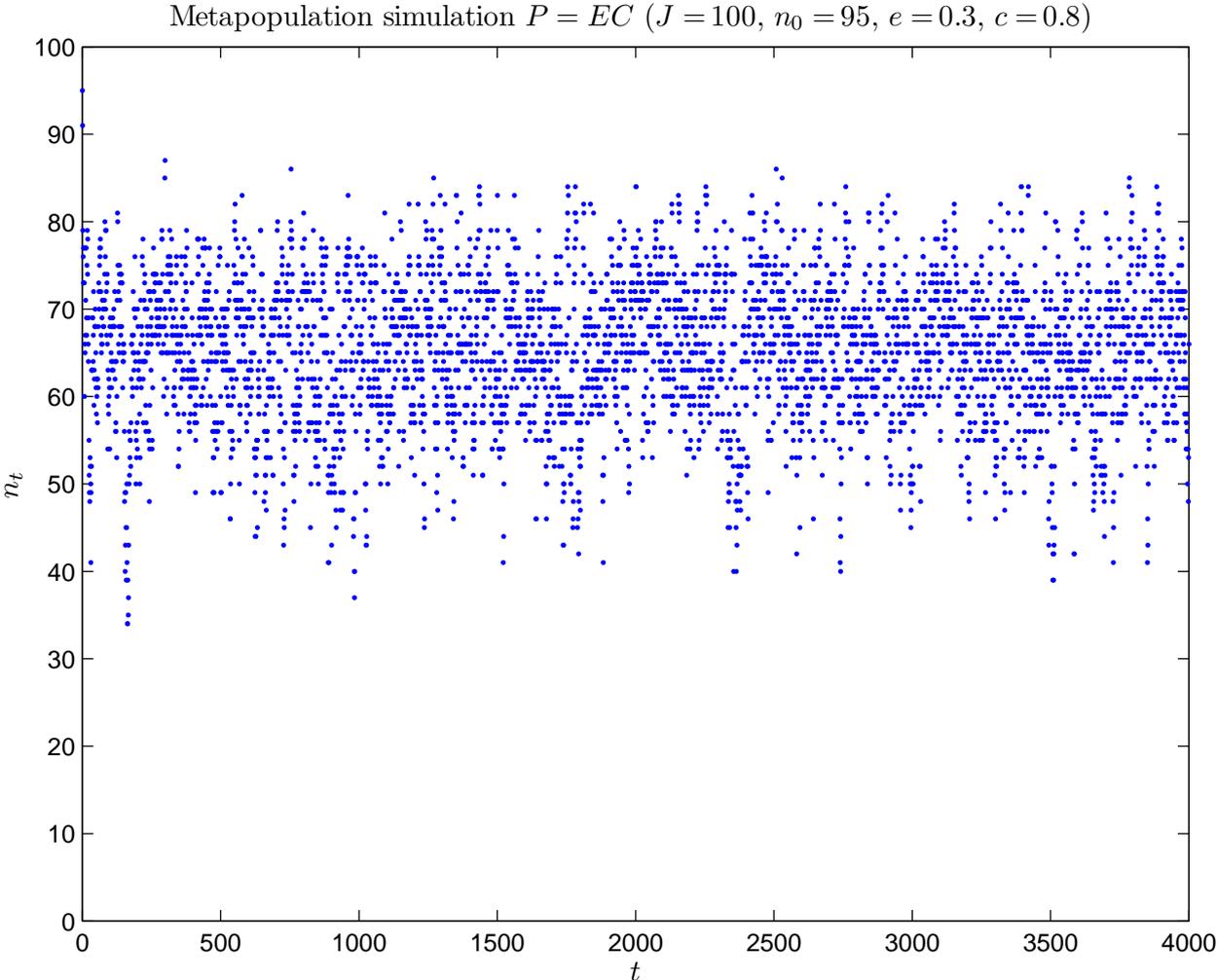
$$\lim_{t \rightarrow \infty} \Pr(n_t = j | n_t \neq 0) = m_j.$$

qsd:

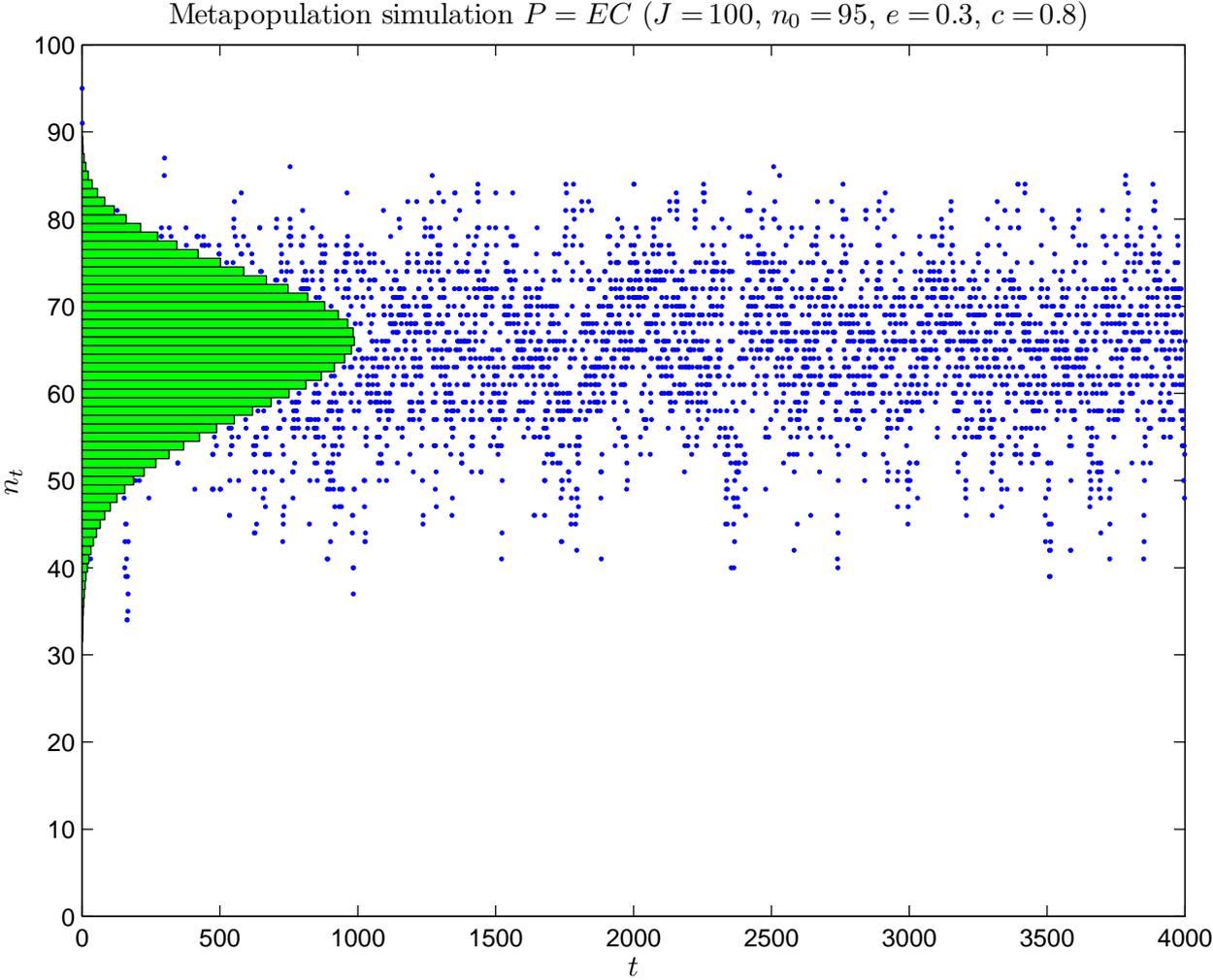
$$\Pr(n_0 = j) = m_j \implies \Pr(n_t = j | n_t \neq 0) = m_j \quad (\forall t > 0).$$

\*In the infinite state space setting, the distinction between lcd and qsd is both subtle and interesting.

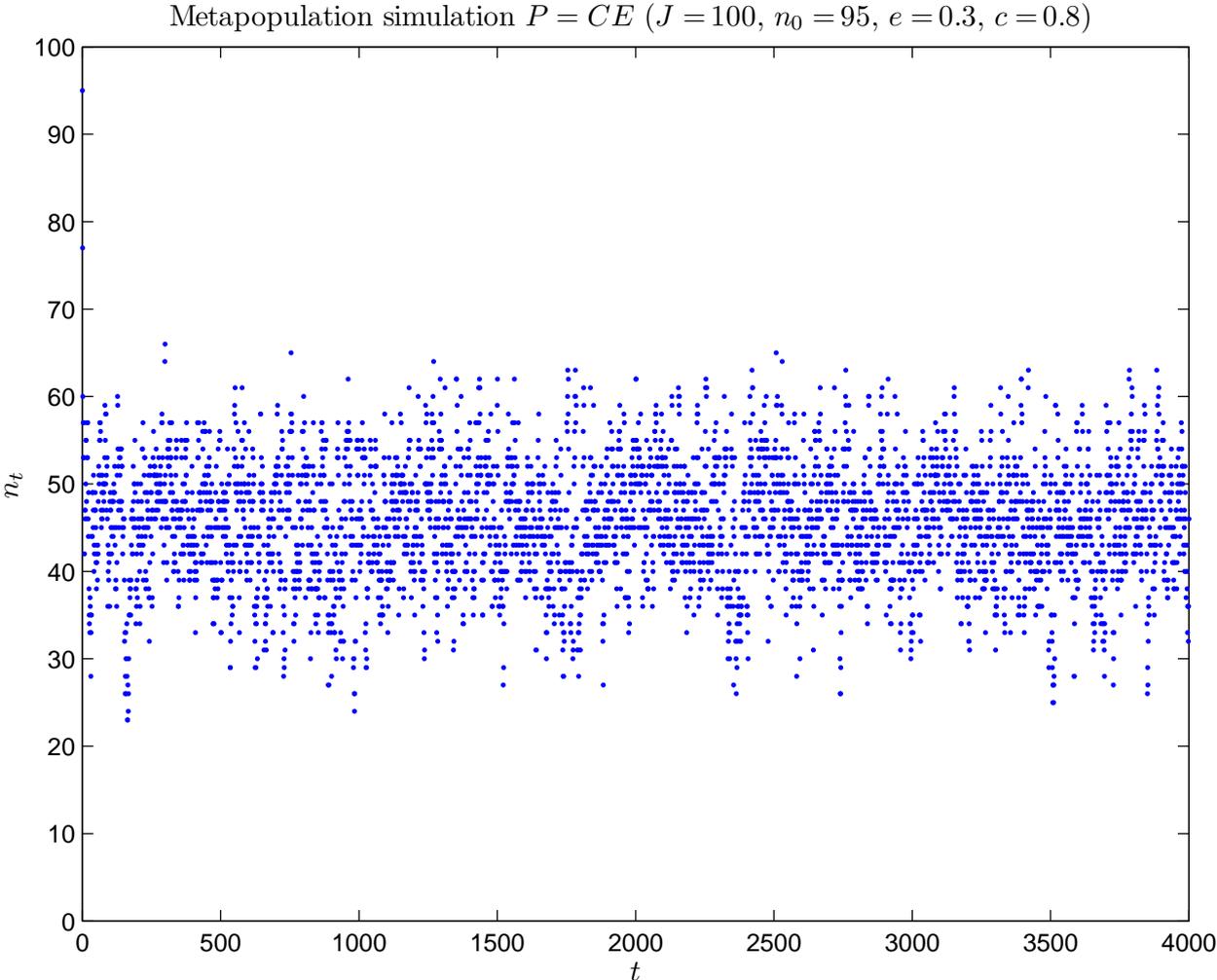
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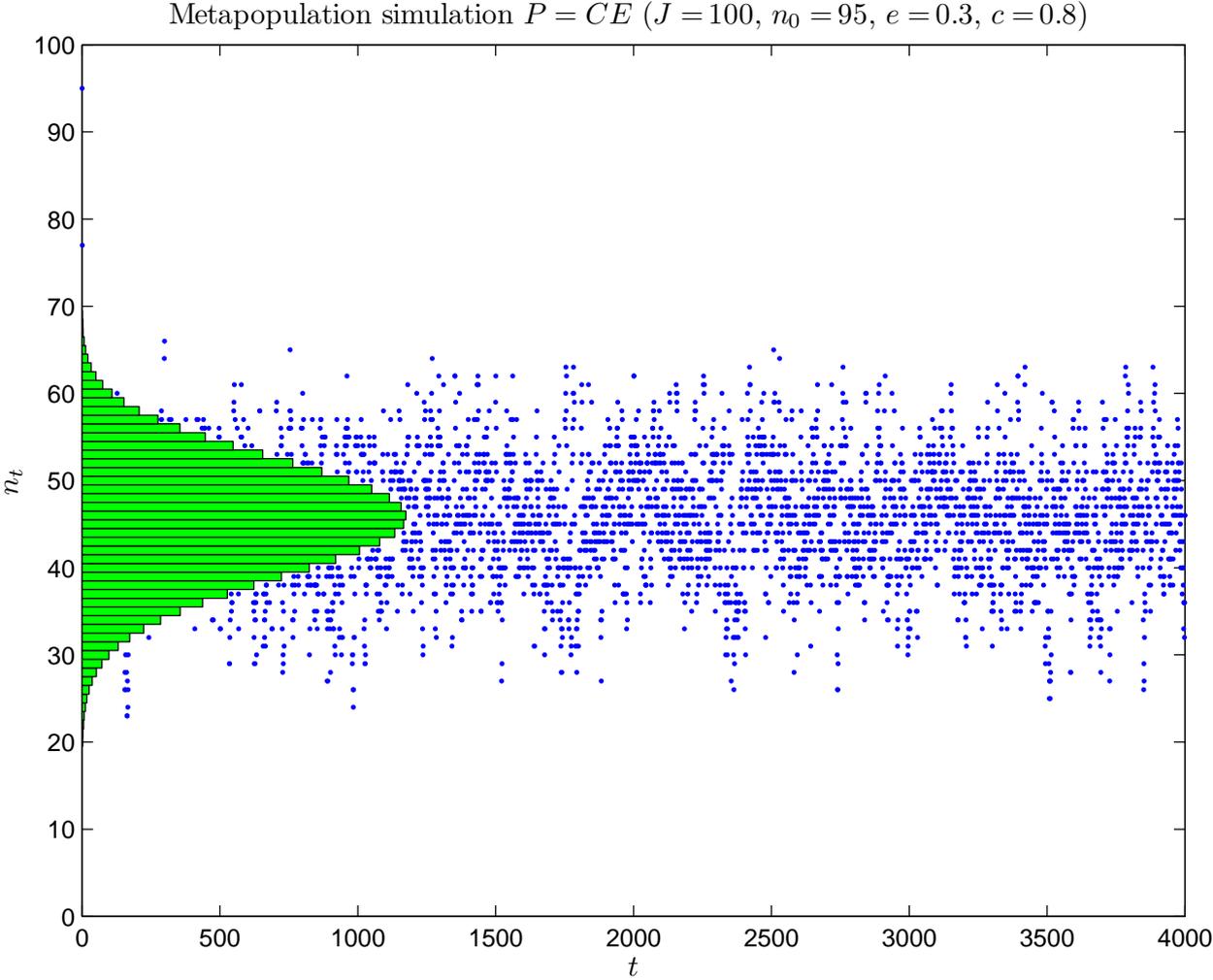
# Simulation and qsd: $P = EC$



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# *J*-patch Mainland-Island models

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This greatly simplifies the analysis!

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The behaviour of both models can be summarized in terms of a single pair of parameters  $(p, q)$ :

$$EC\text{-model: } p = 1 - e(1 - c) \text{ and } q = c$$

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We can improve on this result ...

# *J*-patch Mainland-Island models

Reparameterize by setting  $a = p - q = (1 - e)(1 - c)$ , being the *same* for both models ( $0 < a < 1$ ), and  $q^* = q/(1 - a)$ . Define sequences  $(p_t)$  and  $(q_t)$  by

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# *J*-patch Mainland-Island models

We have in particular that

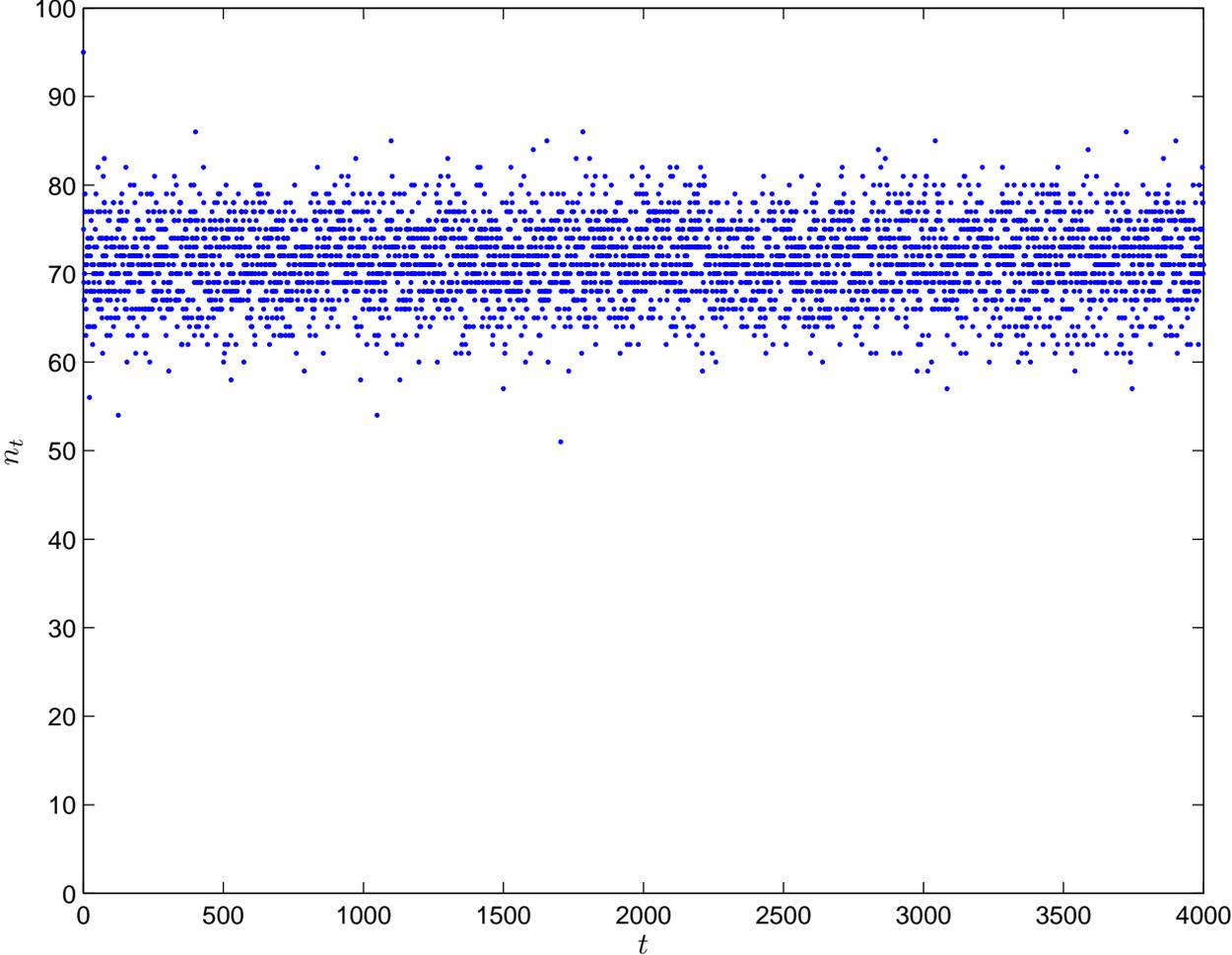
$$\mathbf{E}(n_t | n_0 = i) = ip_t + (J - i)q_t = ia^t + Jq_t$$
$$(\rightarrow Jq^* \text{ as } t \rightarrow \infty)$$

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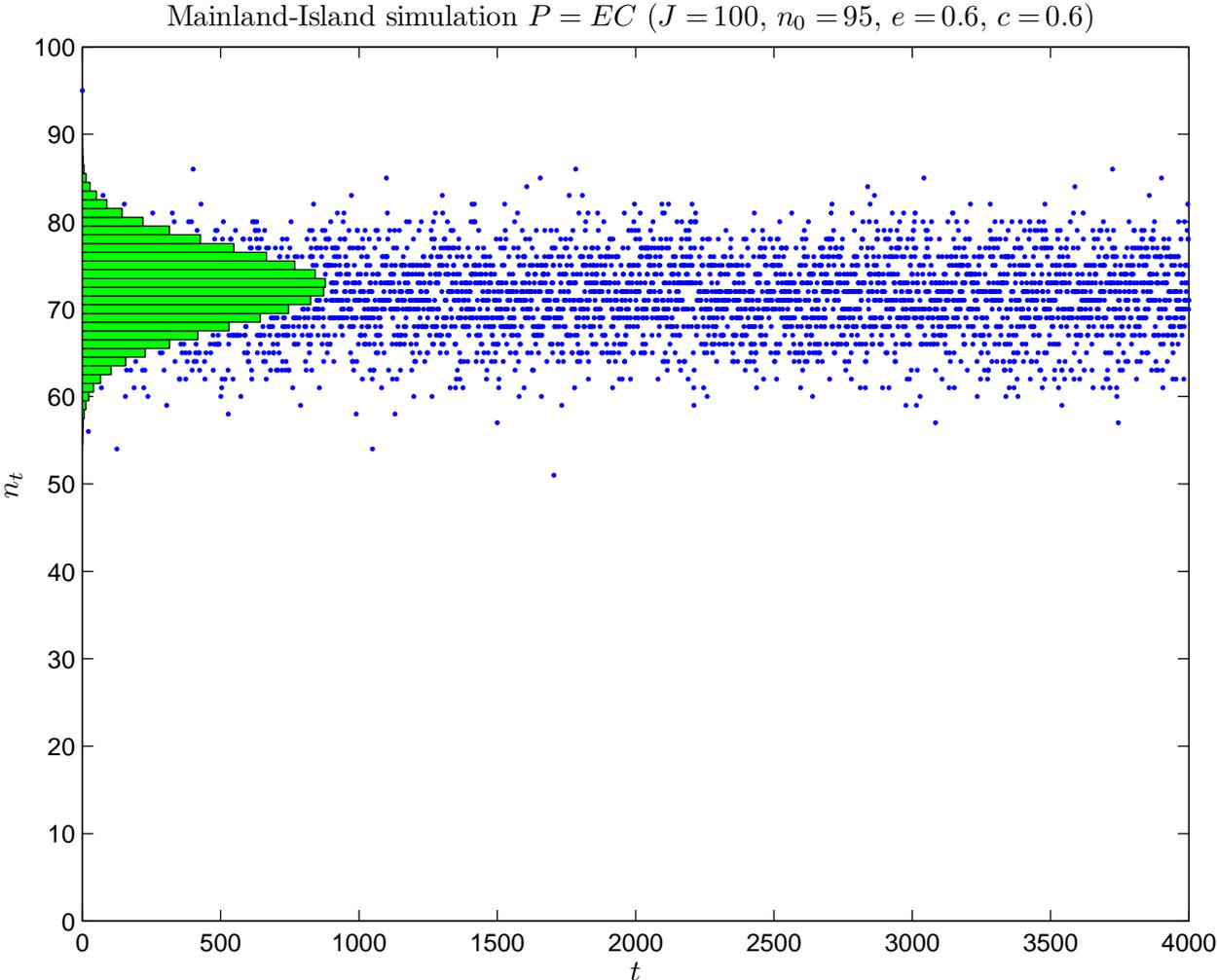
$$\mathbf{Var}(n_t | n_0 = i) = ip_t(1 - p_t) + (J - i)q_t(1 - q_t)$$
$$= ia^t(1 - a^t)(1 - 2q^*) + Jq_t(1 - q_t)$$
$$(\rightarrow Jq^*(1 - q^*) \text{ as } t \rightarrow \infty).$$

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Mainland-Island simulation  $P = EC$  ( $J = 100, n_0 = 95, e = 0.6, c = 0.6$ )

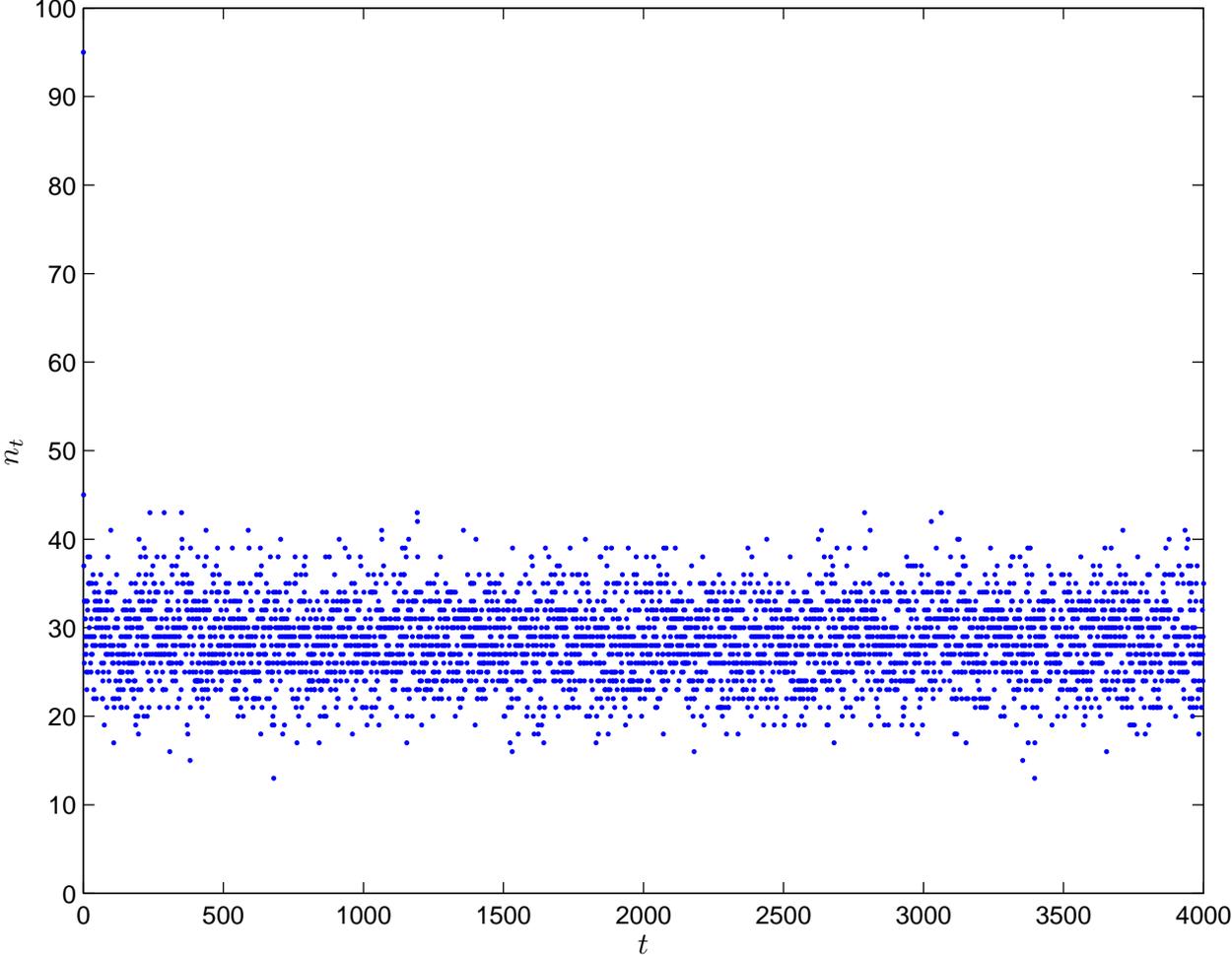


# Simulation and sd: $P = EC$

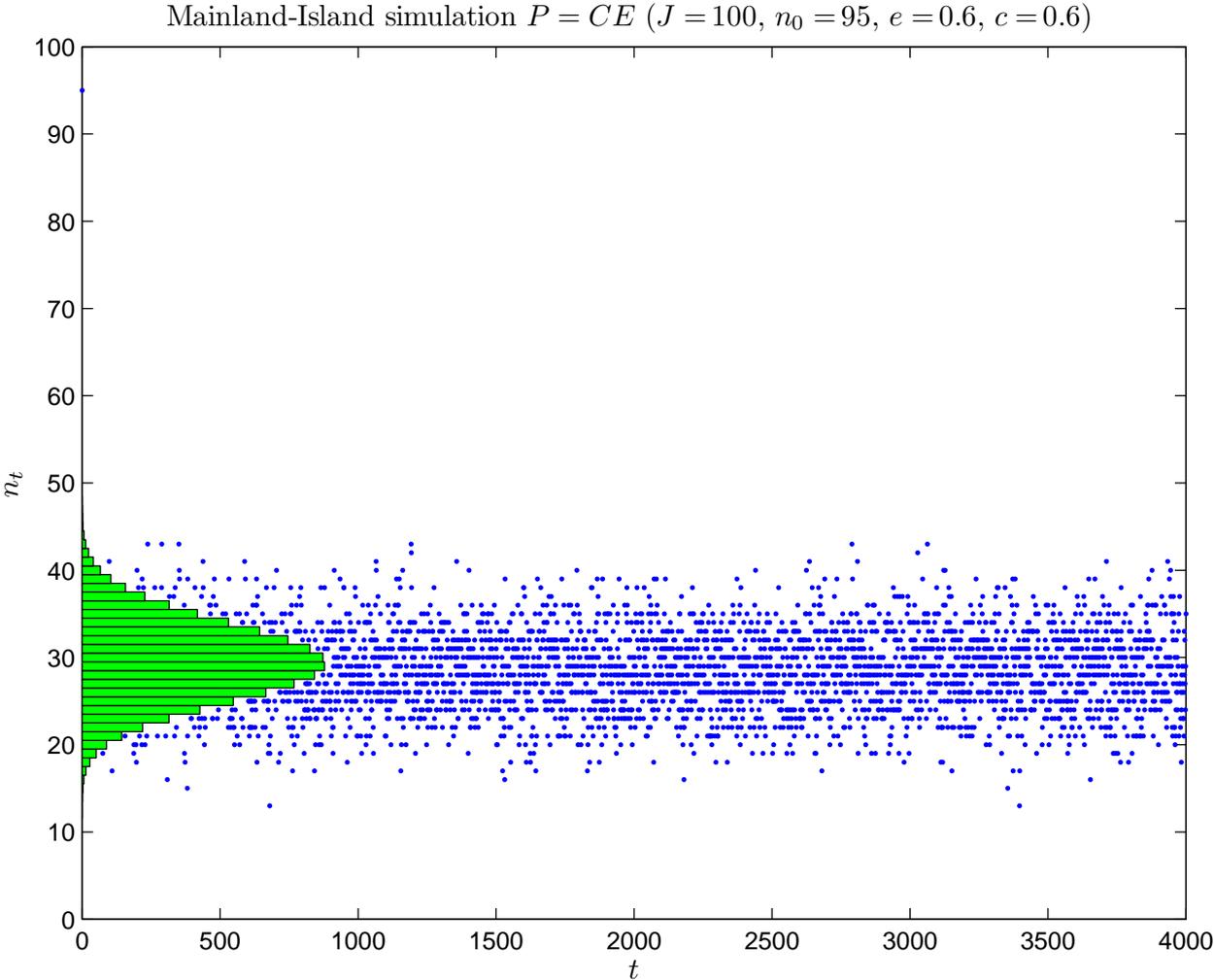


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Note that in contrast with our earlier infinite-state models, state 0 is *no longer absorbing*.

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Indeed we observe that ...

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*Again we can invoke general theory.*

# Infinite-patch Mainland-Island models

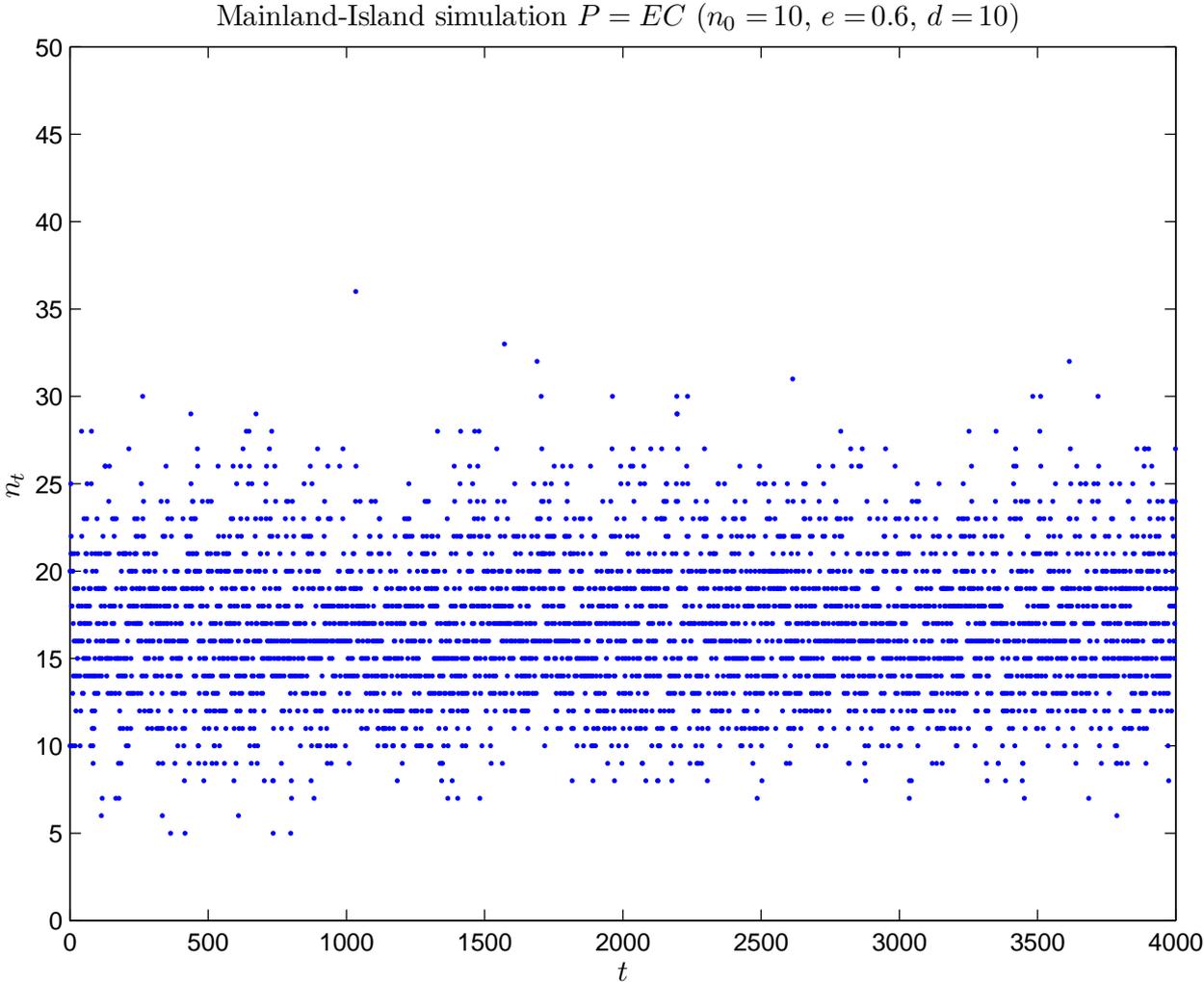
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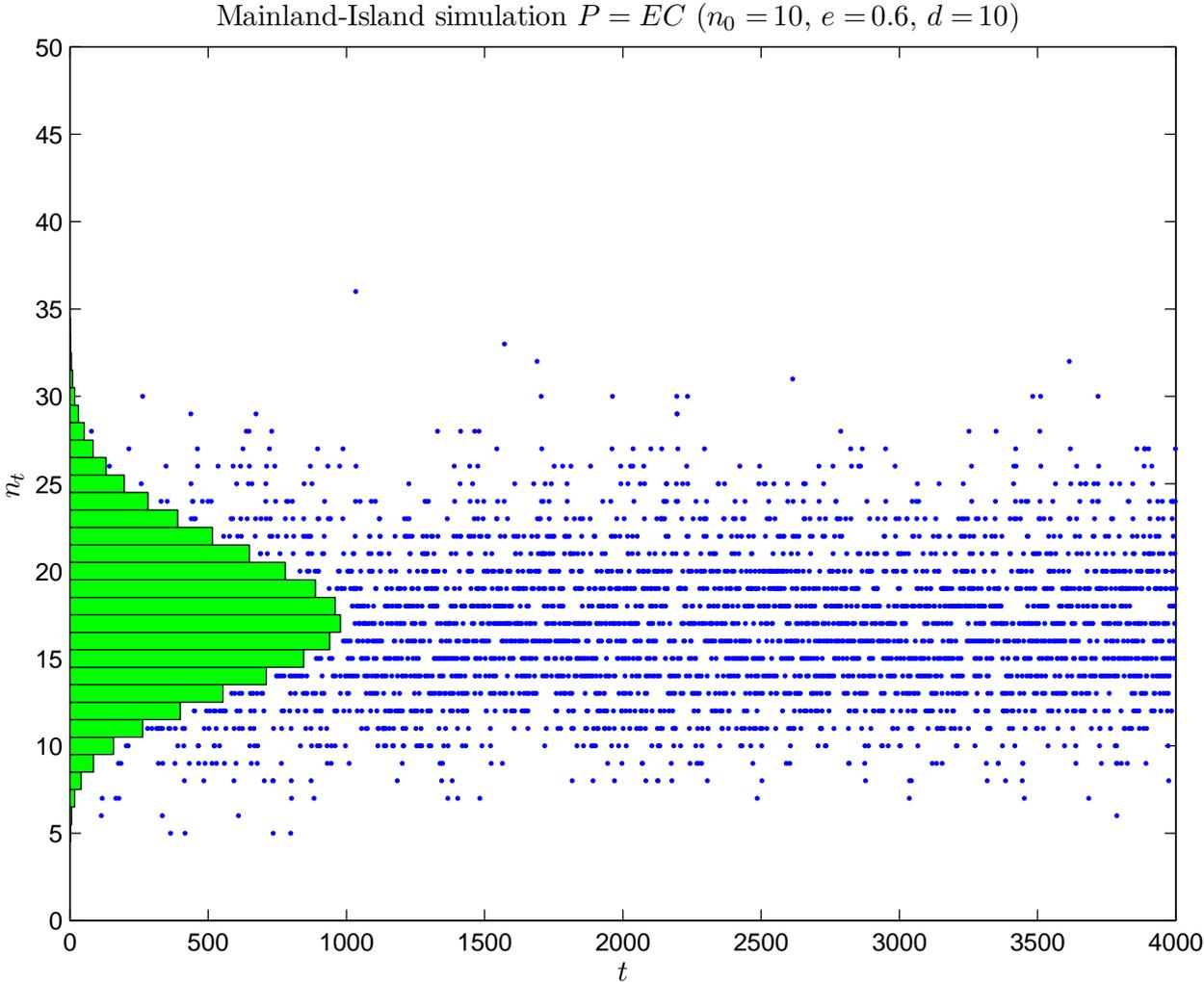
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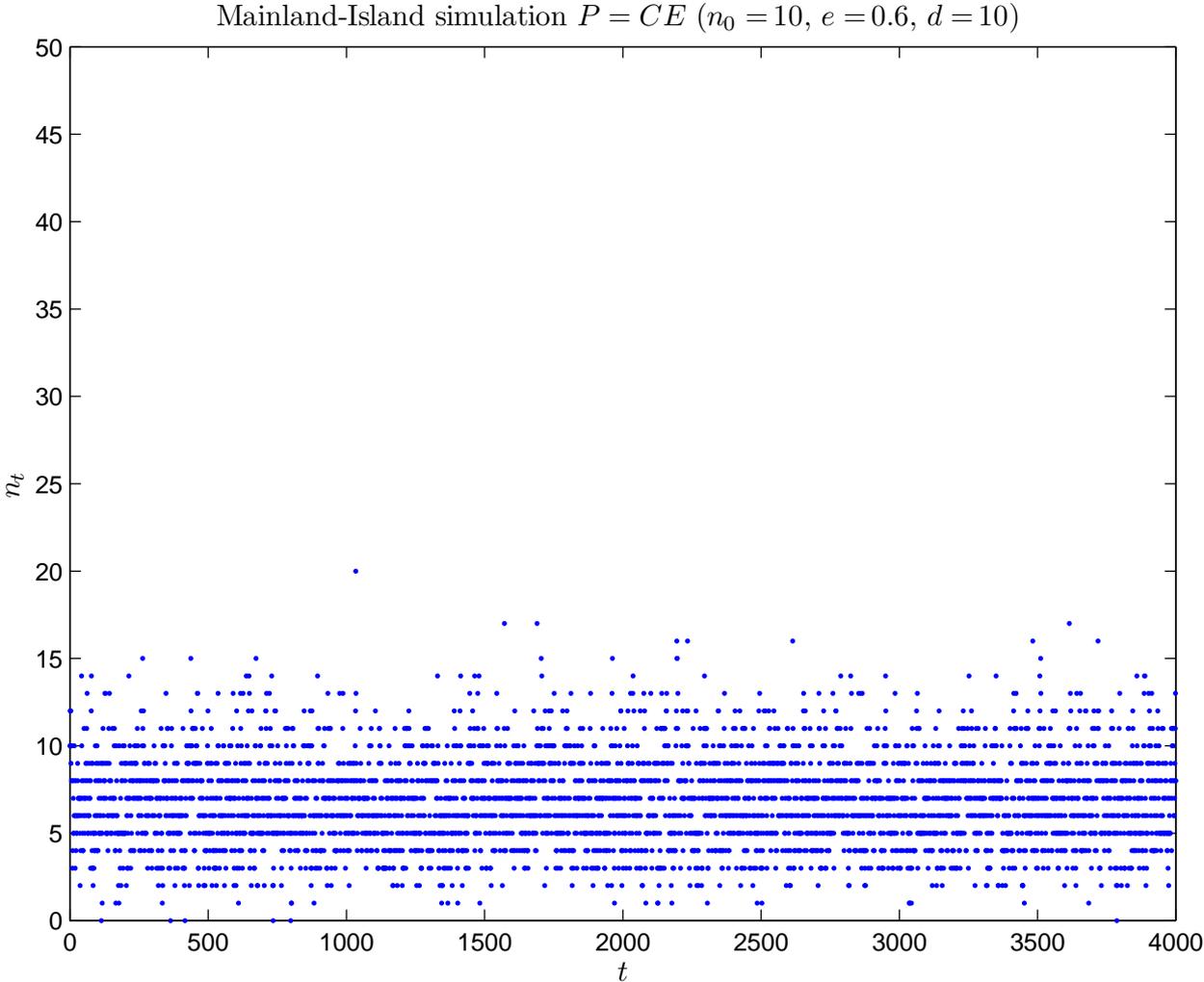
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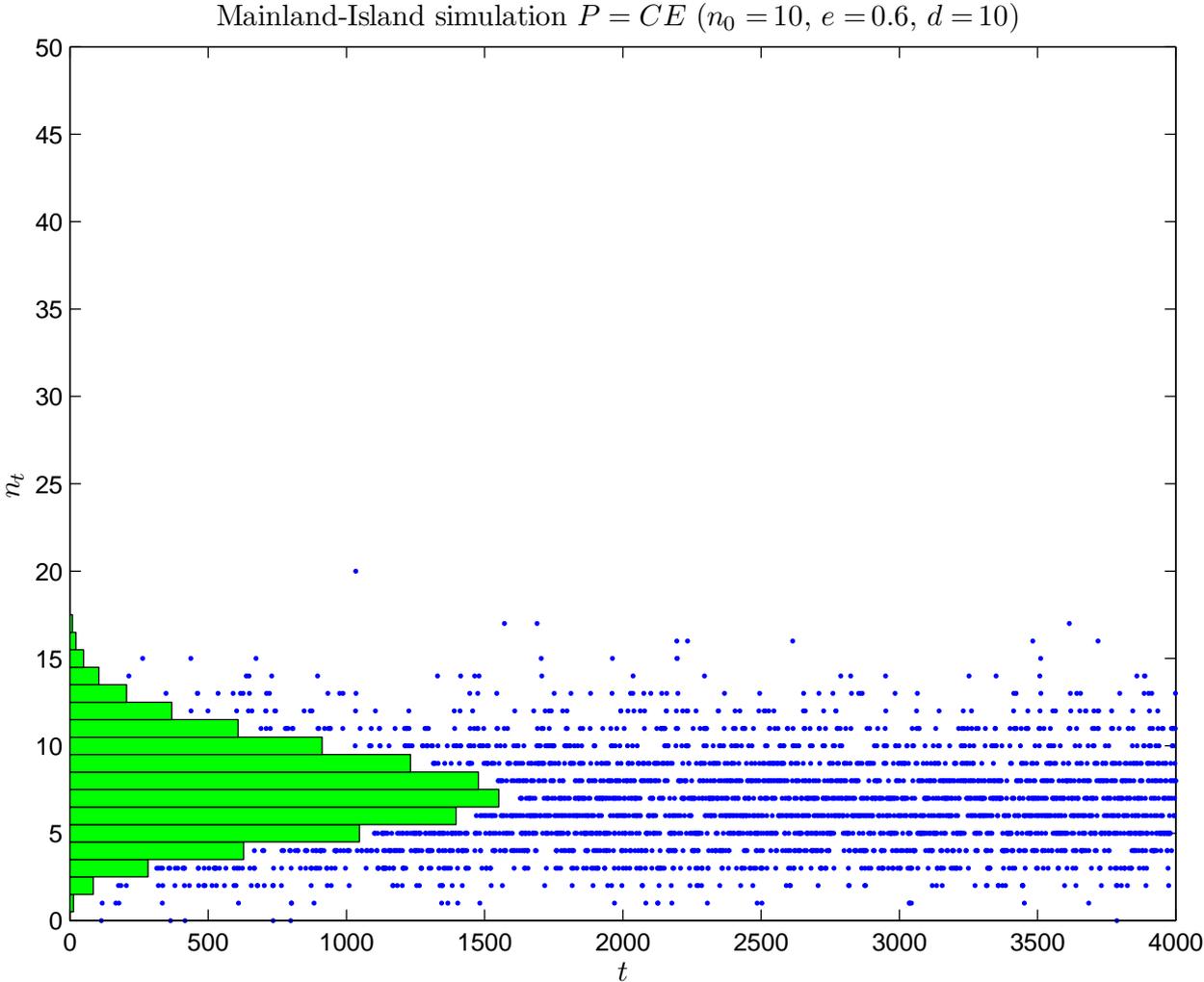
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# First passage times

A measure of persistence for the Mainland-Island models is the expected time to *first* total extinction of the *island network*.

# First passage times

**Theorem** For the  $J$ -patch Mainland-Island model with parameters  $p$  and  $q$ , given  $n_0 = i$  patches occupied initially, the expected time to first enter state 0 is given by

$$\begin{aligned}\mu_{i0} &= \sum_{k=1}^J \binom{J}{k} \frac{b^k}{1 - a^k} - \sum_{j=0}^i \binom{i}{j} (-1)^j \sum_{k=0}^{J-i} \binom{J-i}{k} \frac{b^k (1 - \delta_{j0} \delta_{k0})}{1 - a^{j+k}} \\ &= \sum_{n=0}^{\infty} \left[ (1 + ba^n)^J - (1 - a^n)^i (1 + ba^n)^{J-i} \right],\end{aligned}$$

where  $a = p - q$  and  $b = q/(1 - p)$ .

# First passage times

**Theorem** For the infinite-patch Mainland-Island model with parameters  $e$  and  $m$ , given  $n_0 = i$  patches occupied initially, the expected time to first enter state 0 is always *finite* and is given by

$$\begin{aligned}\mu_{i0} &= \sum_{j=1}^i \binom{i}{j} (-1)^{j+1} \sum_{n=0}^{\infty} (1-e)^{jn} \exp\left(\frac{m}{e}(1-e)^n\right) \\ &= \sum_{n=0}^{\infty} [1 - (1 - (1-e)^n)^i] \exp\left(\frac{m}{e}(1-e)^n\right).\end{aligned}$$