

# Stochastic models for population networks

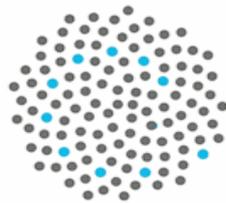
## I: Network models

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AUSTRALIAN RESEARCH COUNCIL  
Centre of Excellence for Mathematics  
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# Collaborators

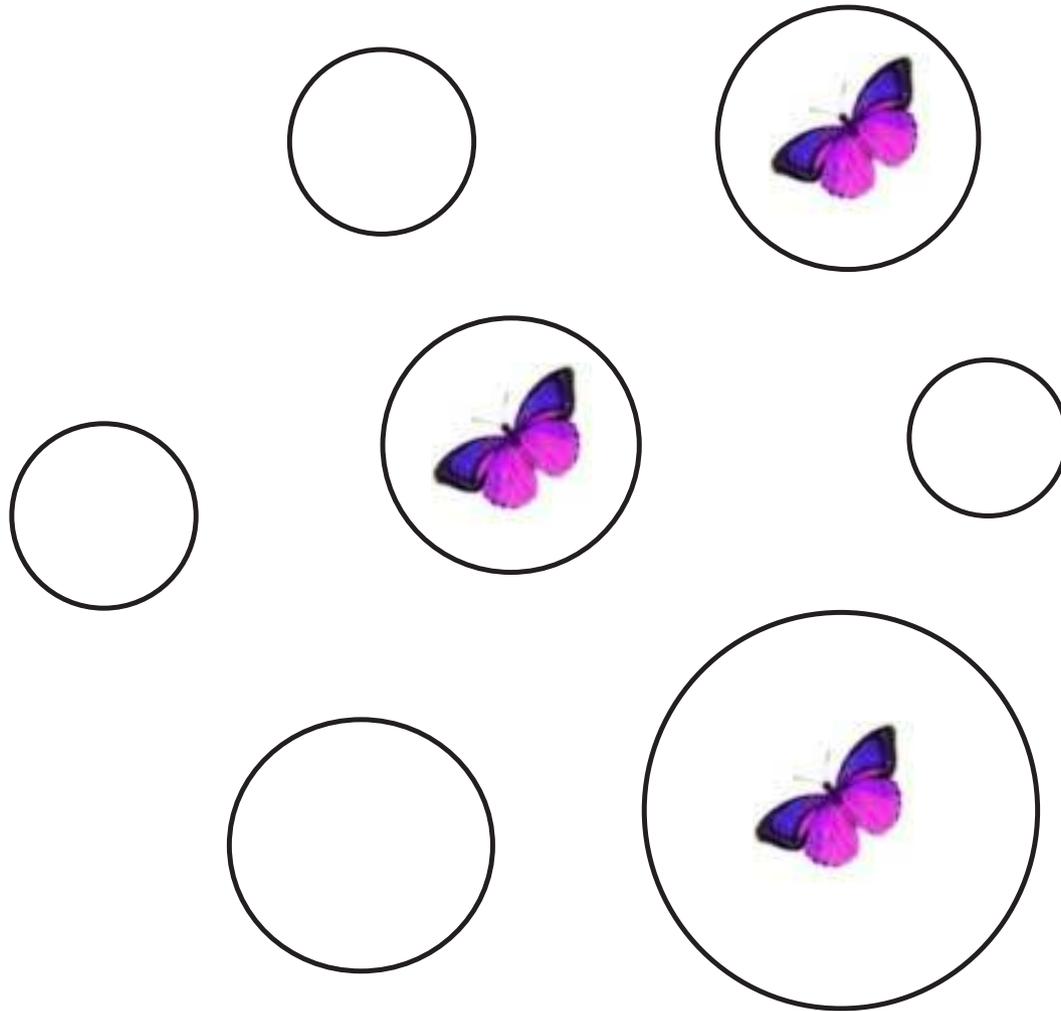
Fionnuala Buckley (MASCOS)  
Department of Mathematics  
The University of Queensland



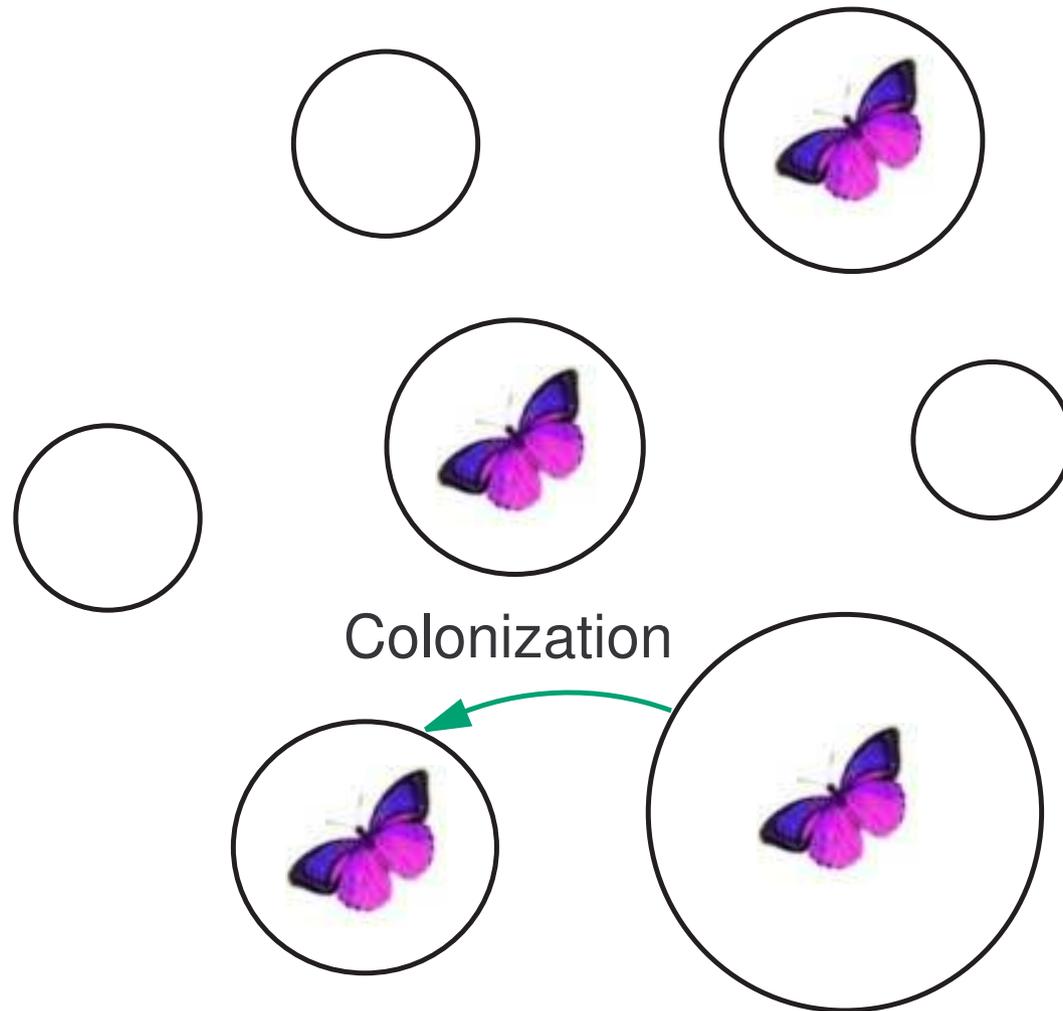
Joshua Ross  
King's College  
University of Cambridge



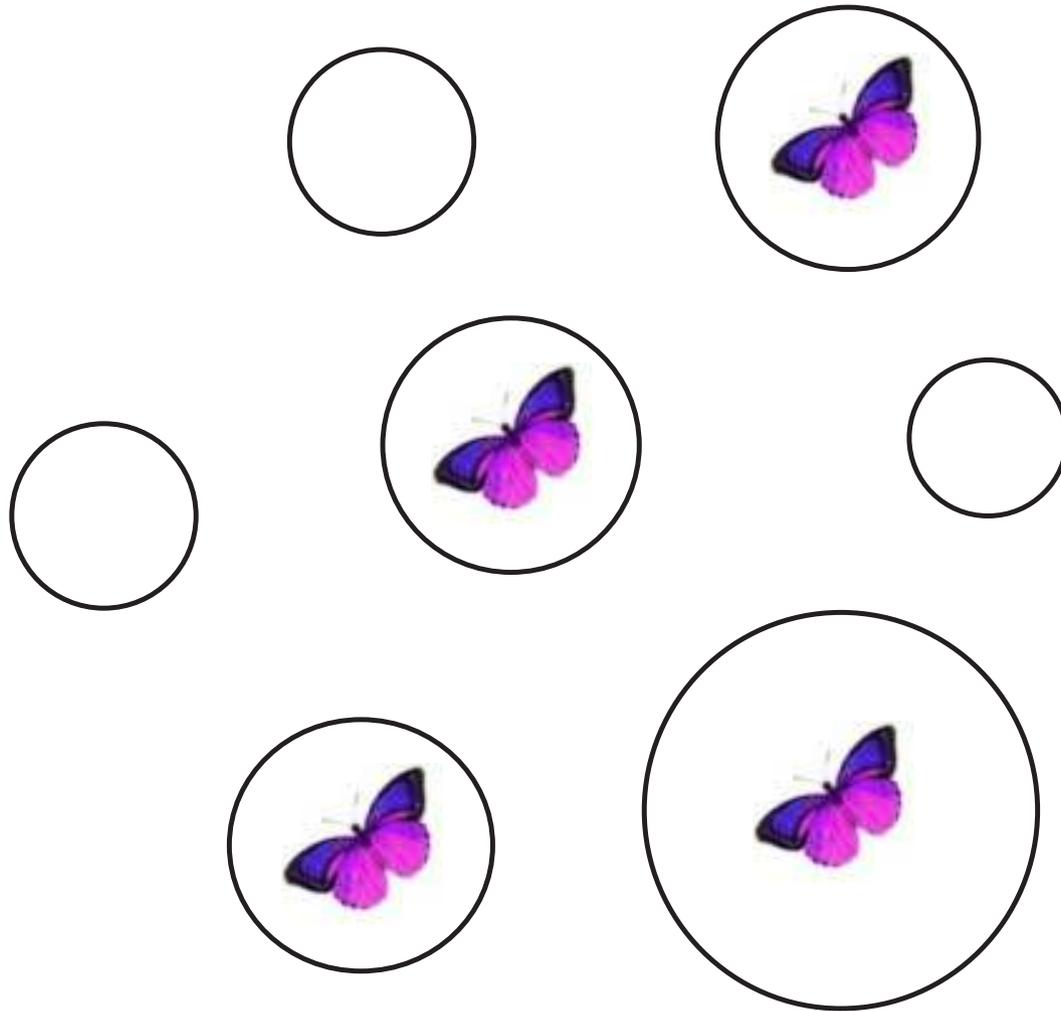
# Metapopulations



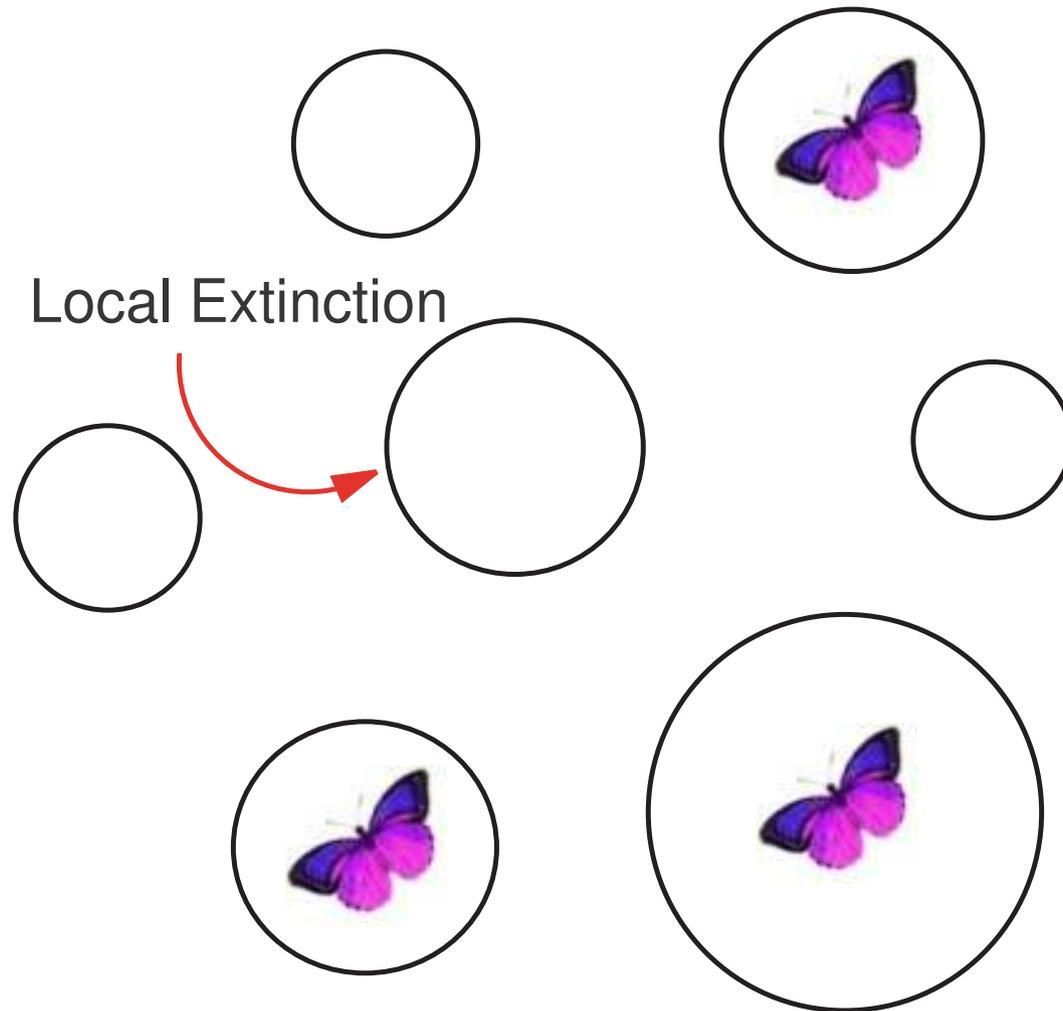
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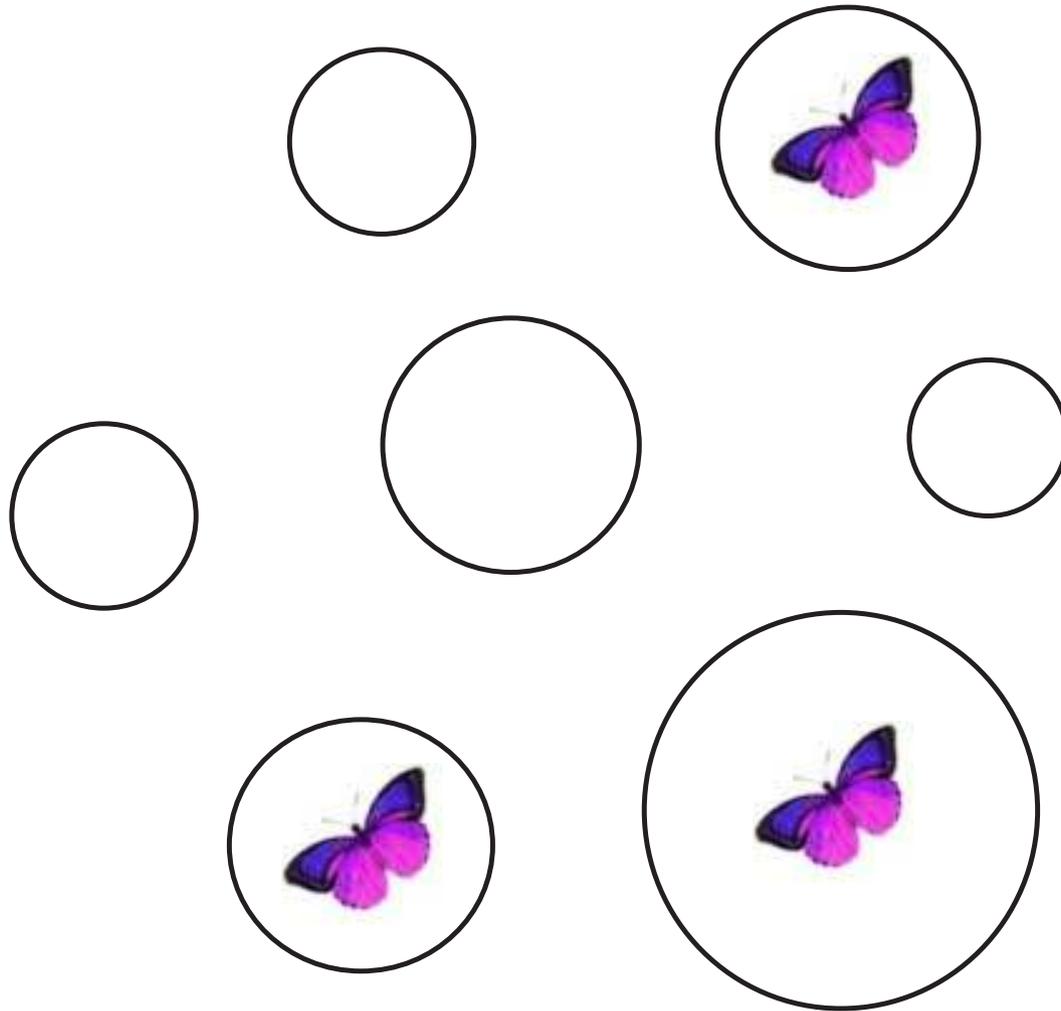
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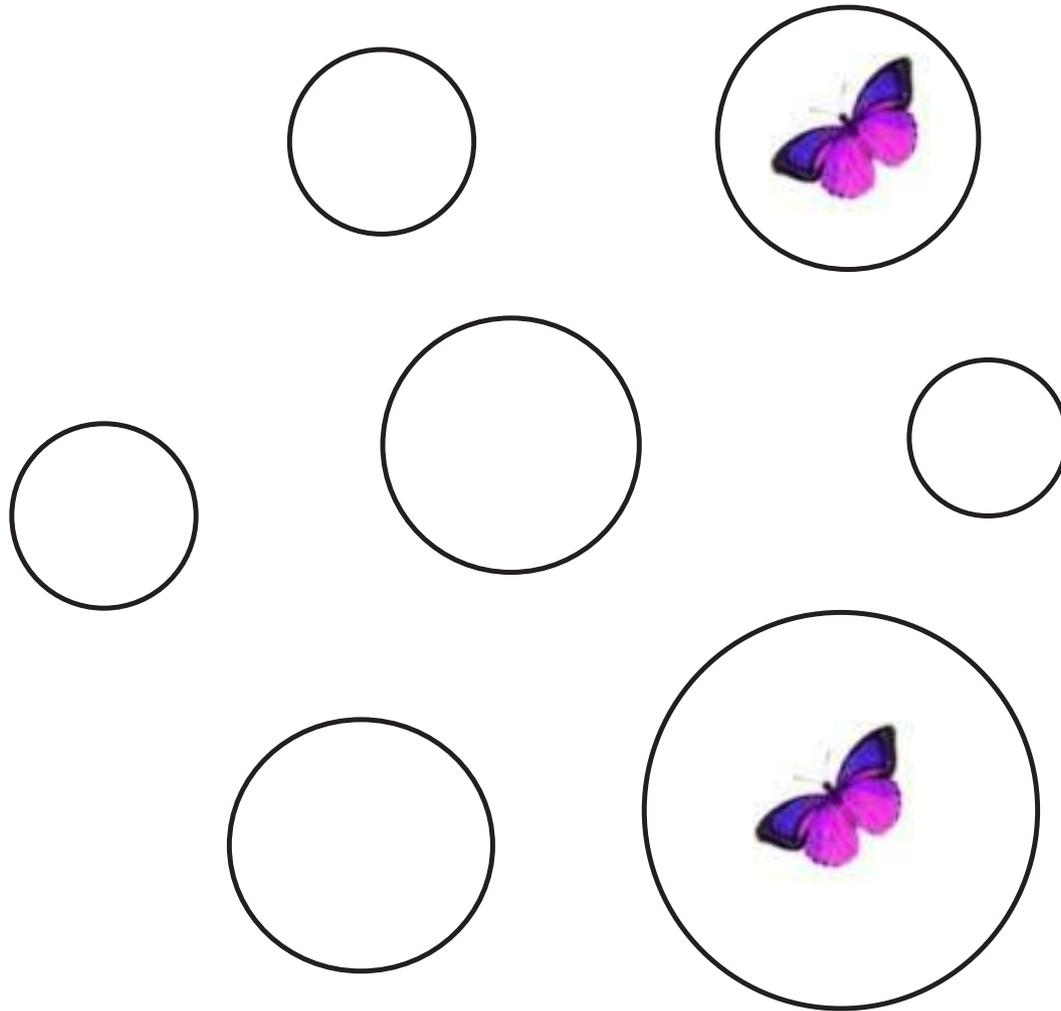
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- A metapopulation is a population that is confined to a network of geographically separated habitat patches (for example a group of islands).
- Individual patches may suffer local extinction.
- Recolonization can occur through dispersal of individuals from other patches.

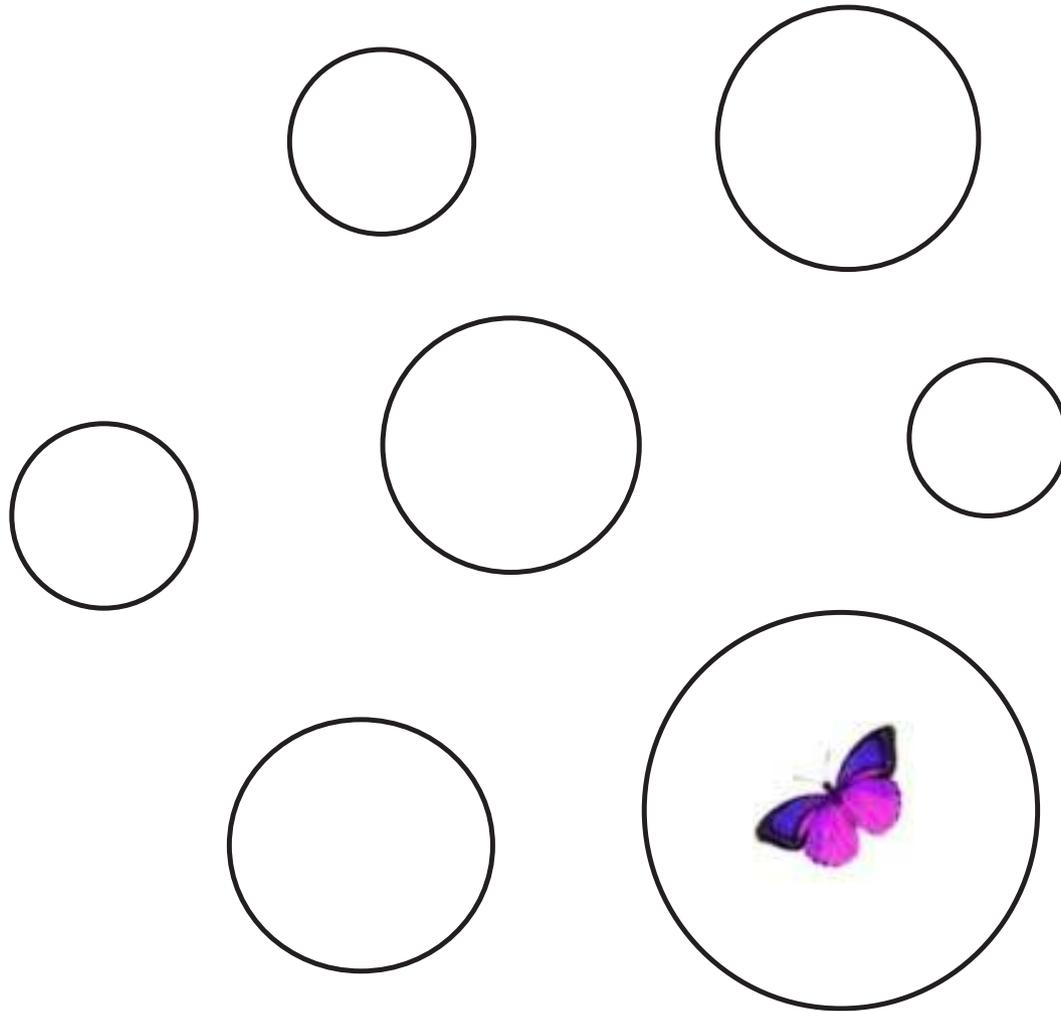
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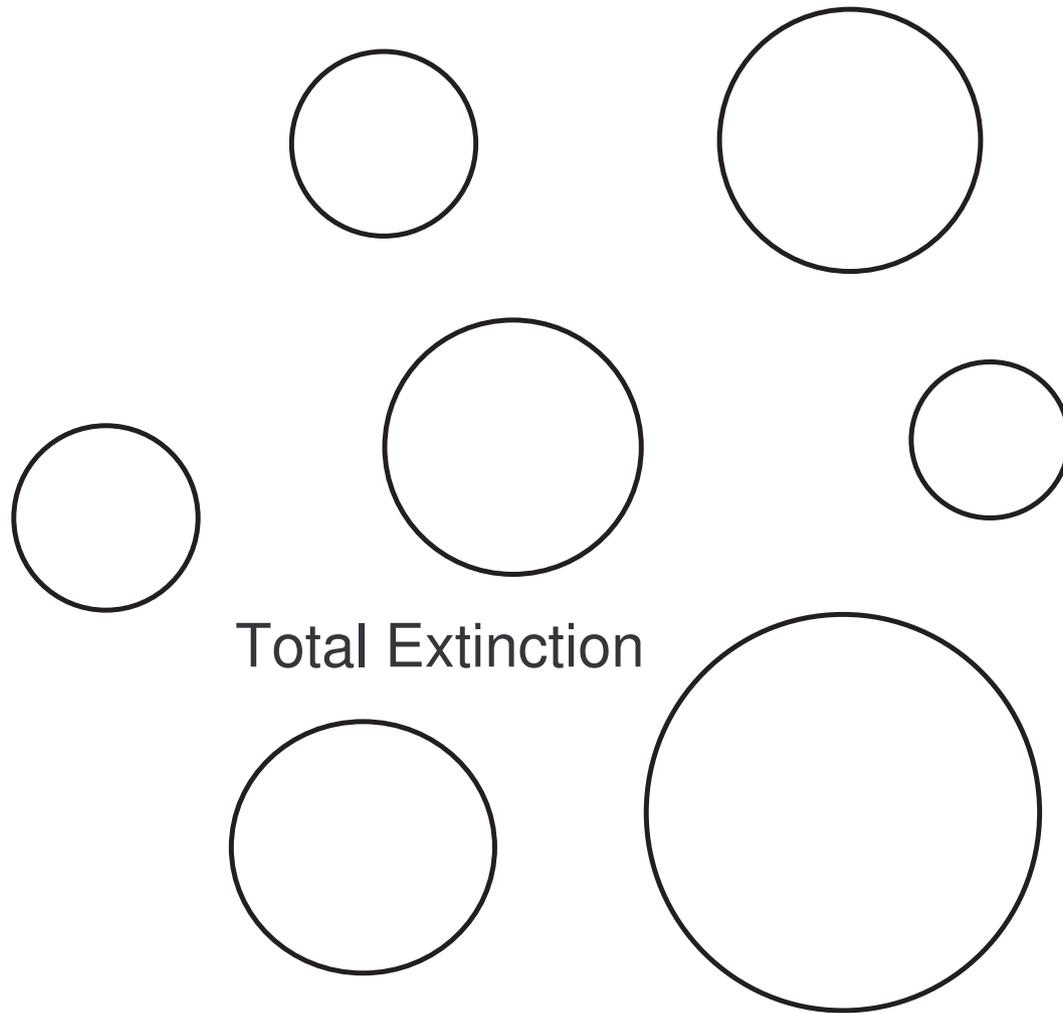
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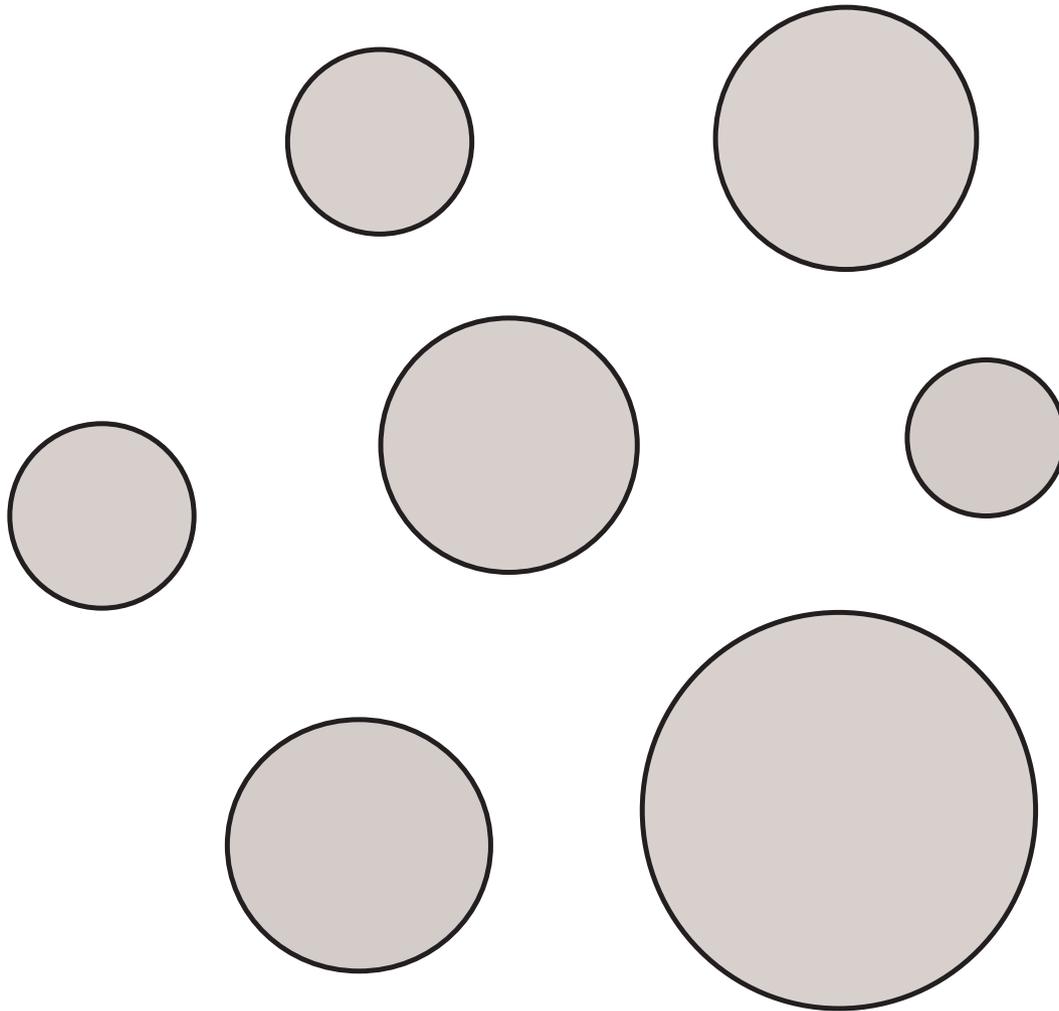


# Metapopulations



Total Extinction

# Metapopulations



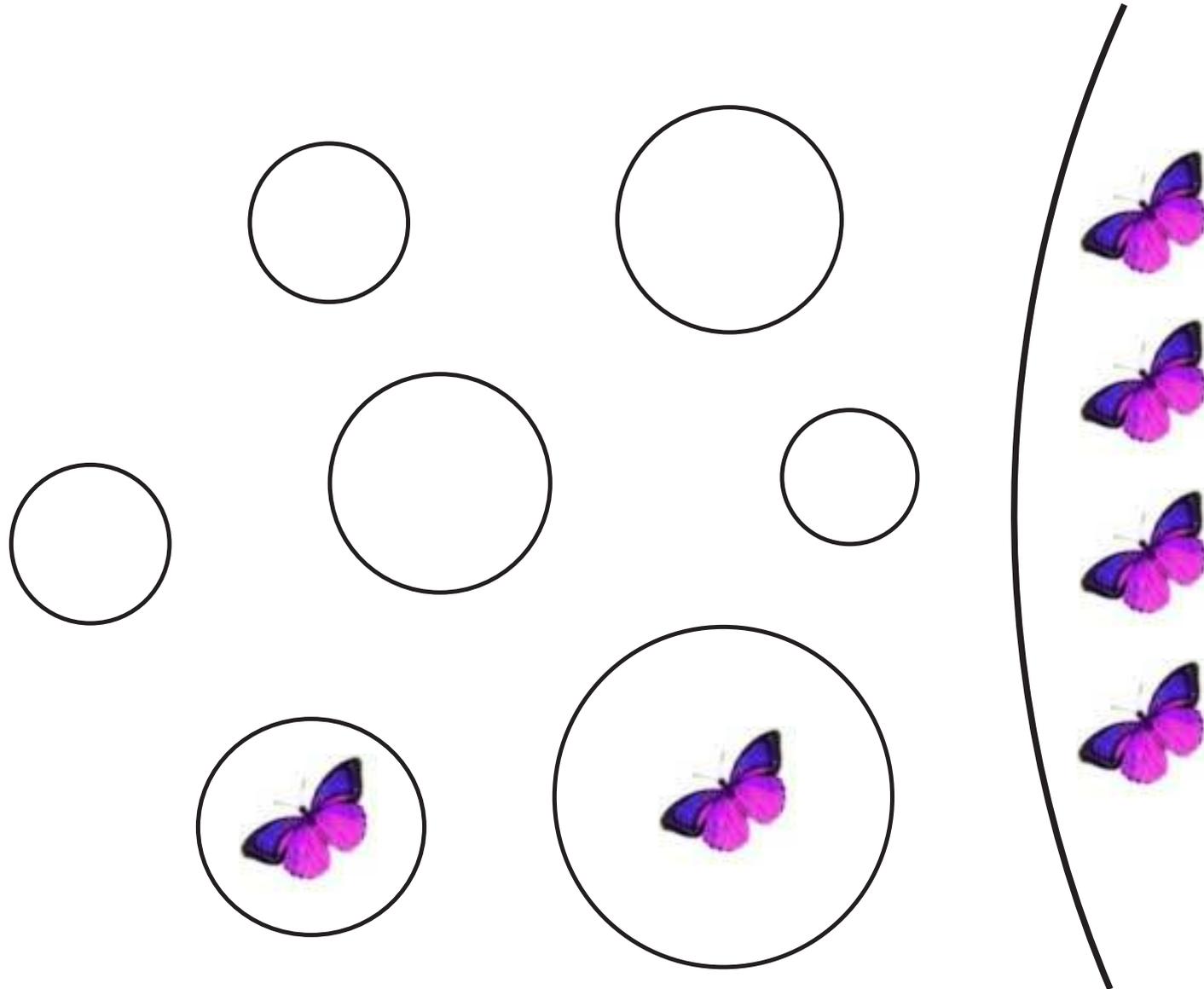
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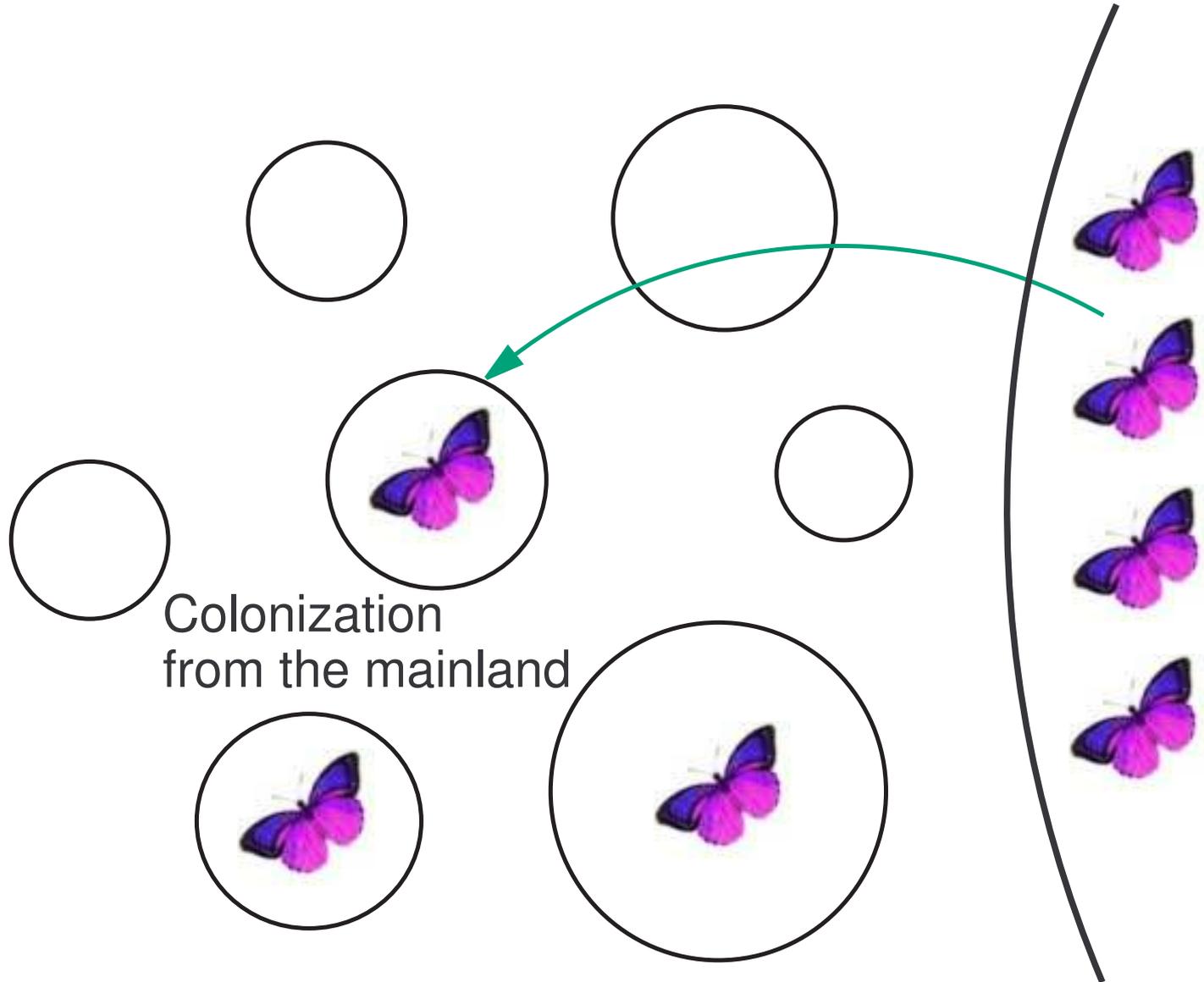
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- Individual patches may suffer local extinction.
- Recolonization can occur through dispersal of individuals from other patches.
- In some instances there is an external source of immigration (mainland-island configuration).

# Mainland-island configuration

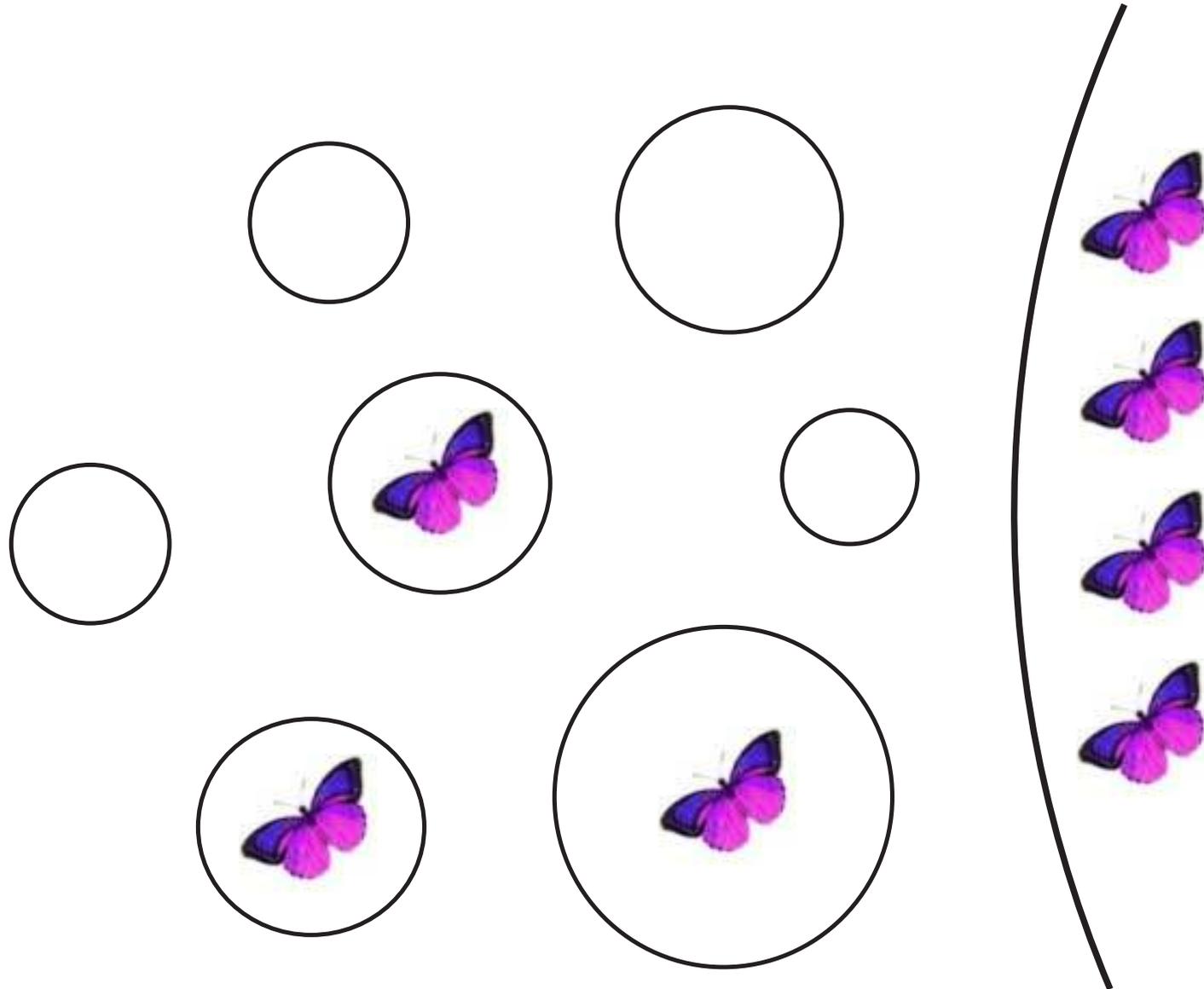


# Mainland-island configuration

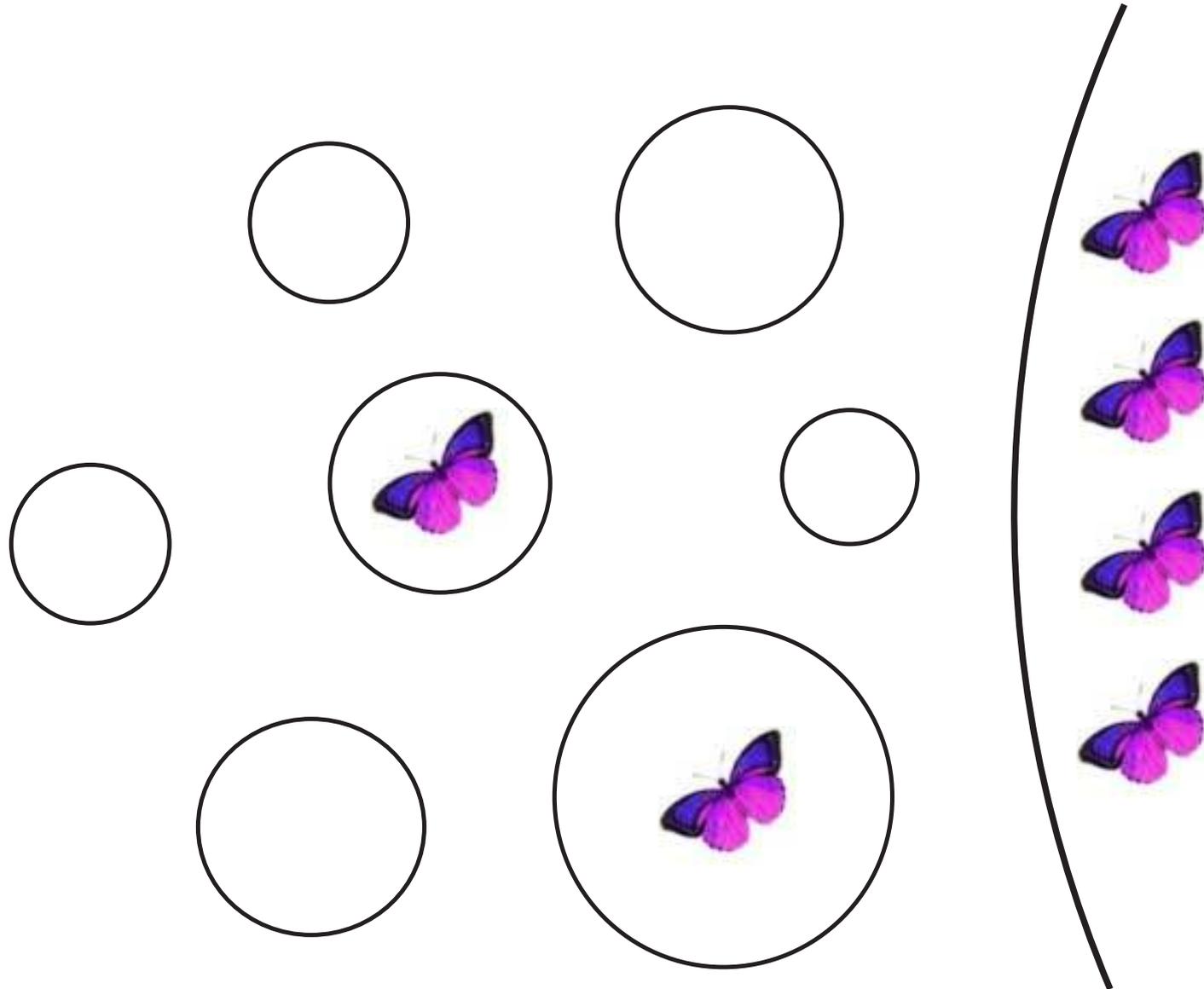


Colonization  
from the mainland

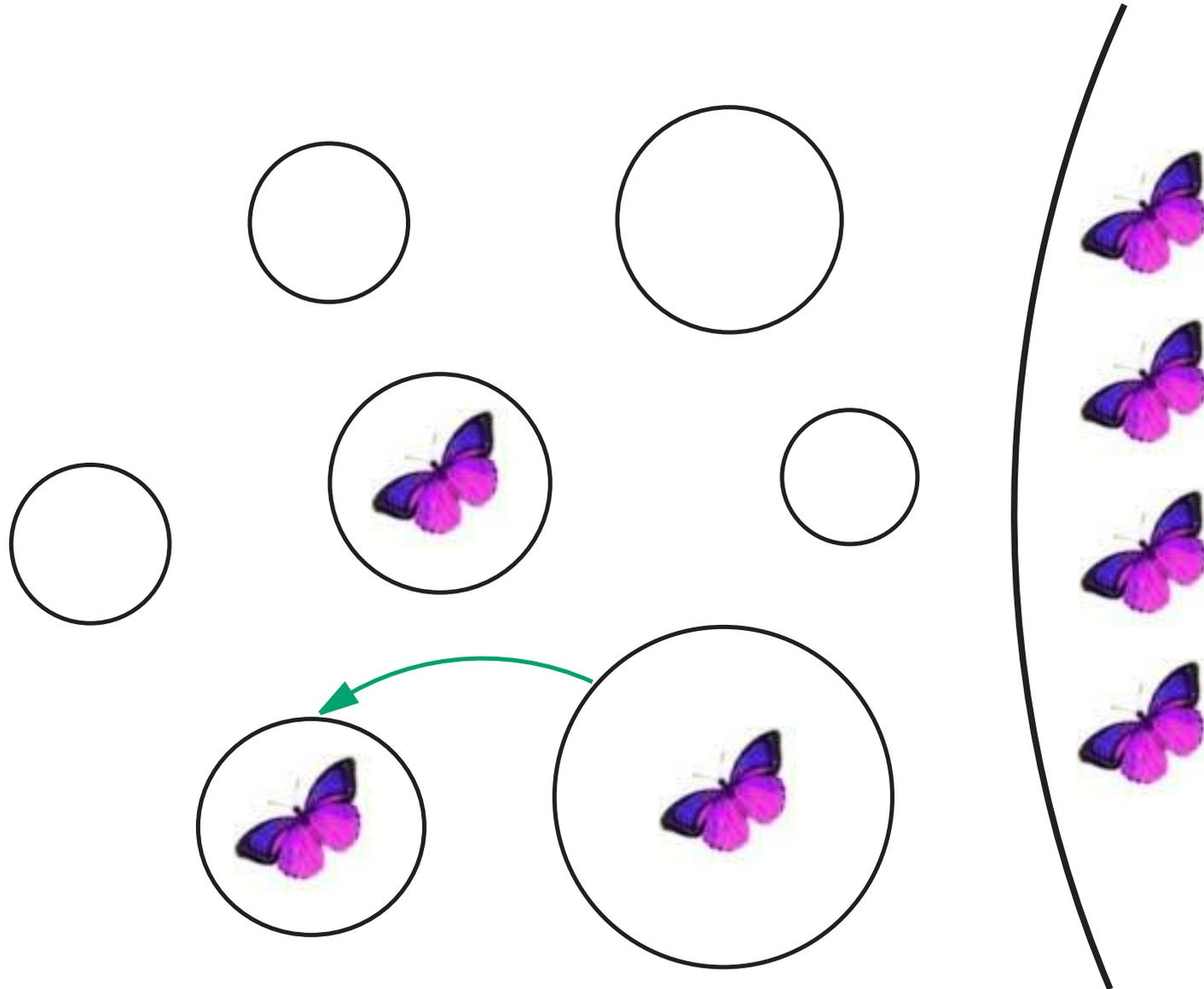
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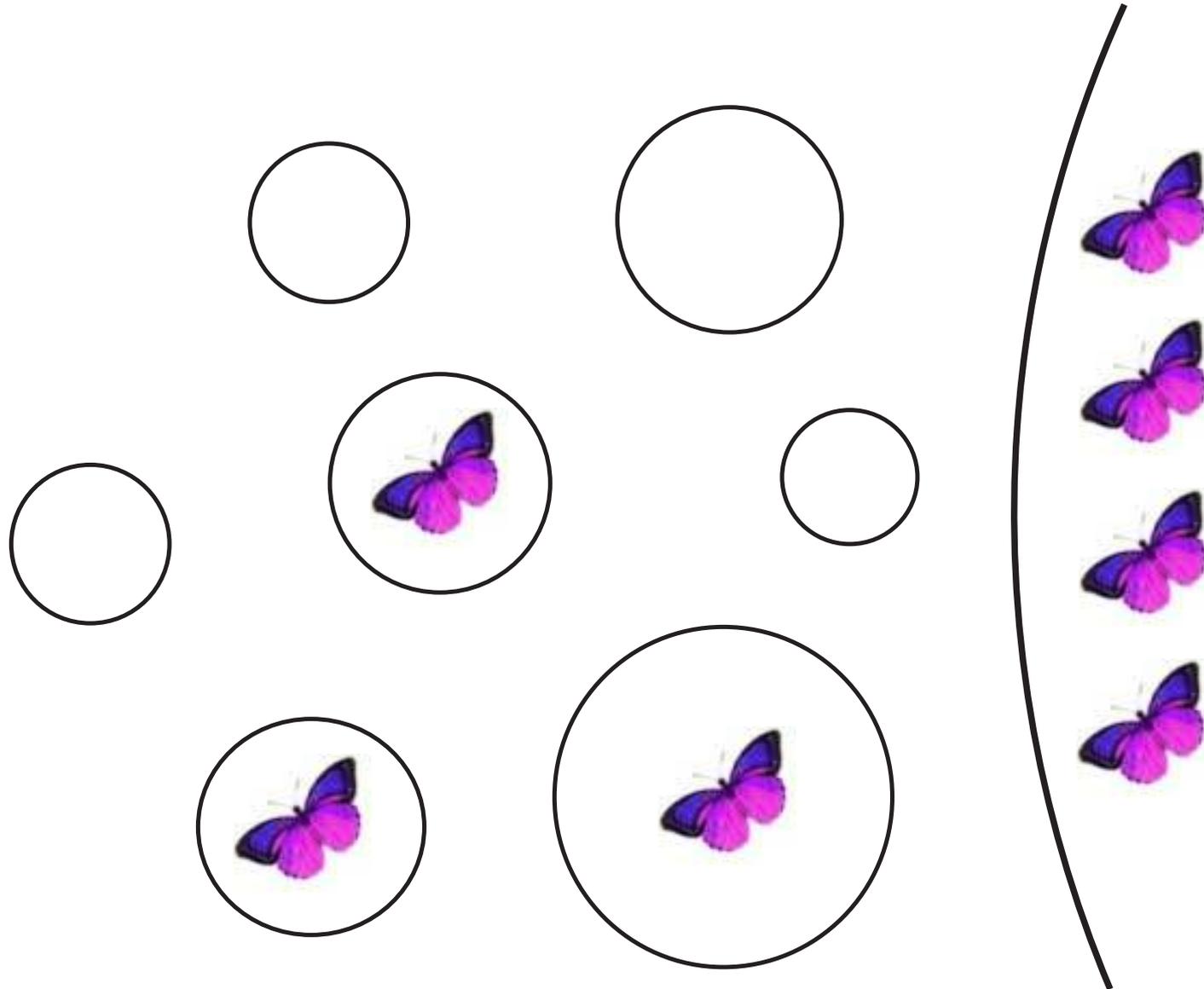
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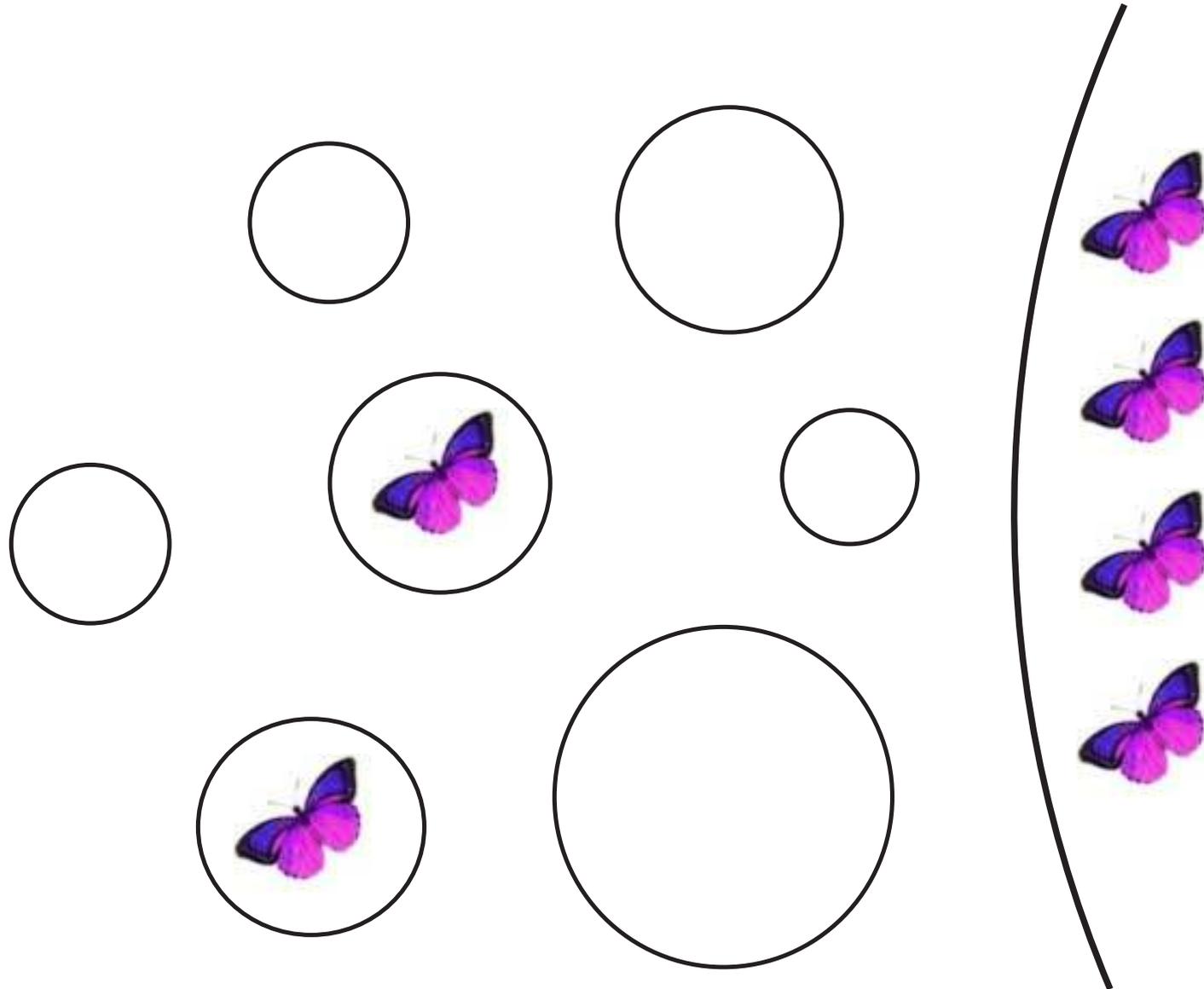
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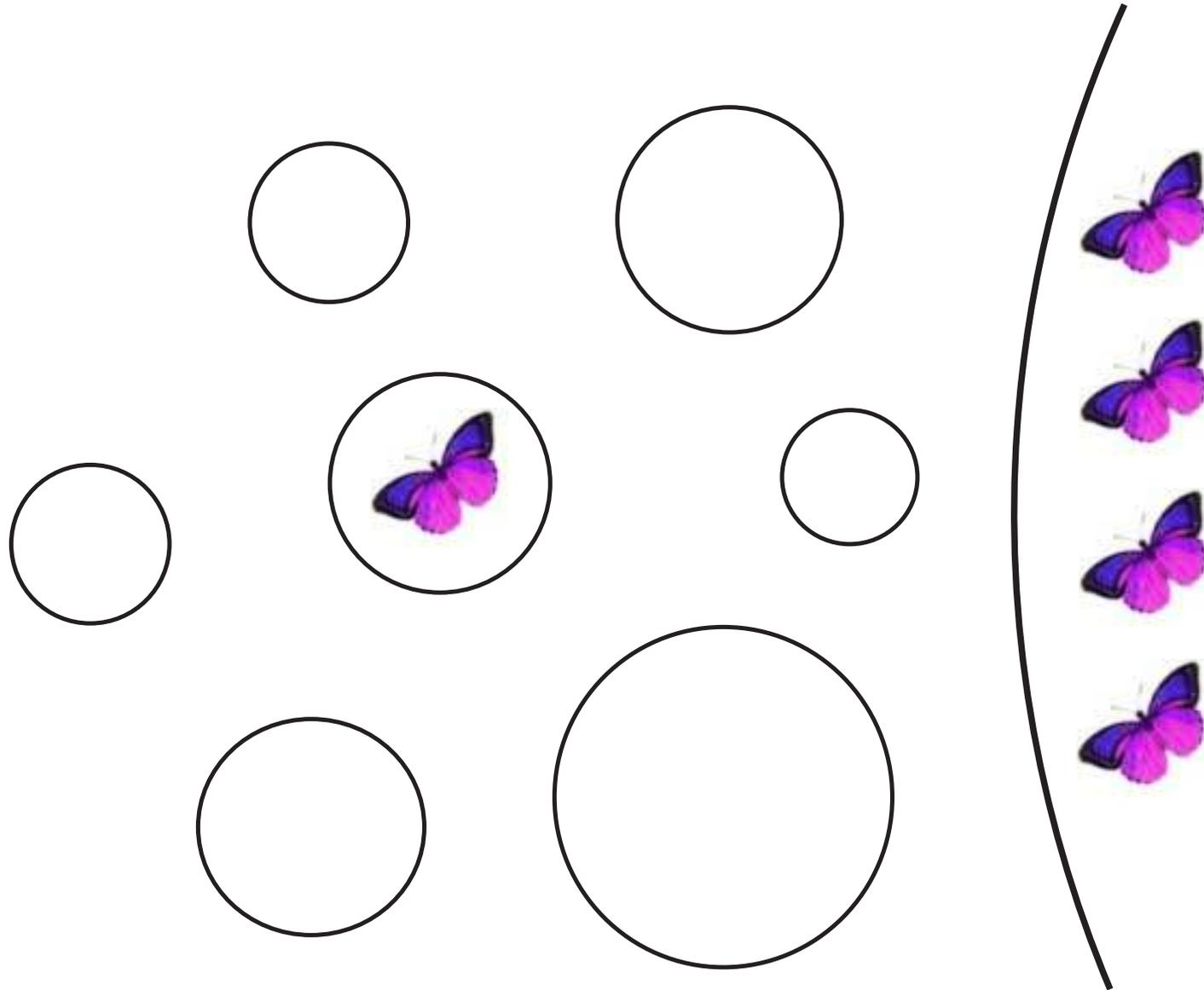
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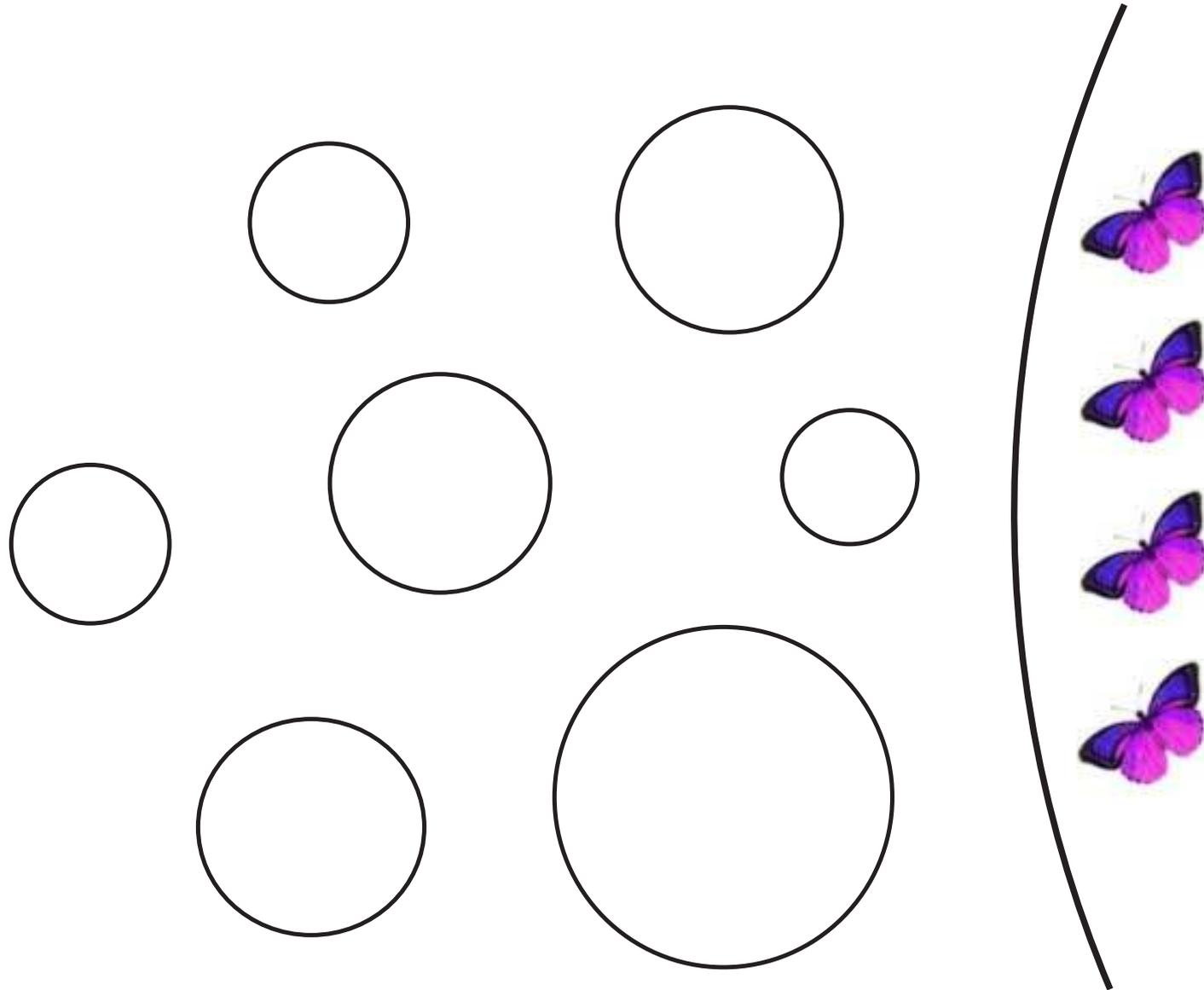
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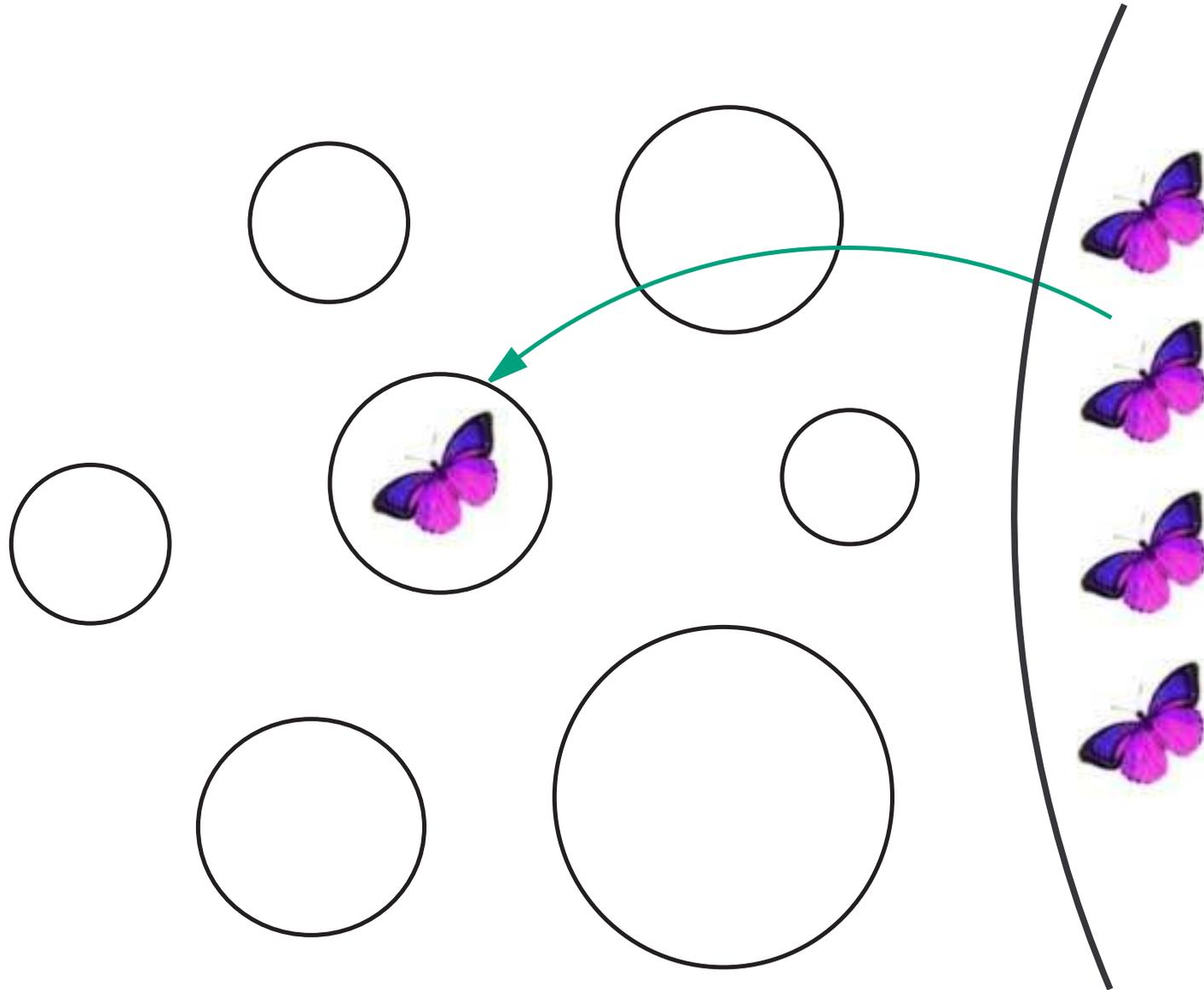
# Mainland-island configuration



# 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

Here we simply record the *number*  $n_t$  of occupied patches at each time  $t$ .

A typical approach is to suppose that  $(n_t, t \geq 0)$  is a Markov chain in discrete or continuous time.

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**Note.** This entails a high degree of homogeneity among patches (in particular the colonization and local extinction processes).

# A continuous-time model

Suppose that there are  $J$  patches. Each occupied patch becomes empty at rate  $e$  and colonization of empty patches occurs at rate  $c/J$  for each suitable pair.

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

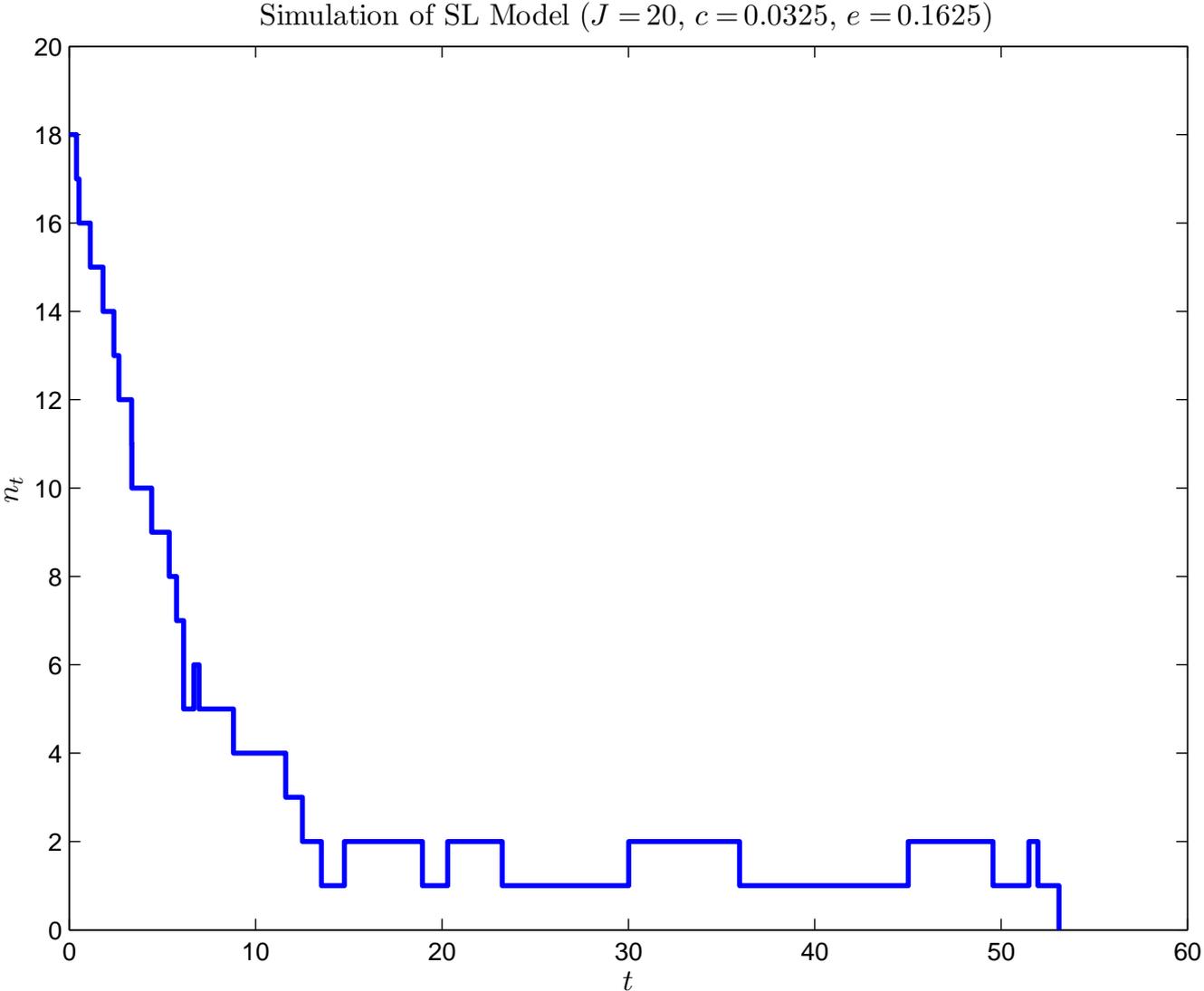
$$n \rightarrow n + 1 \quad \text{at rate} \quad \frac{c}{J}n(J - n)$$

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

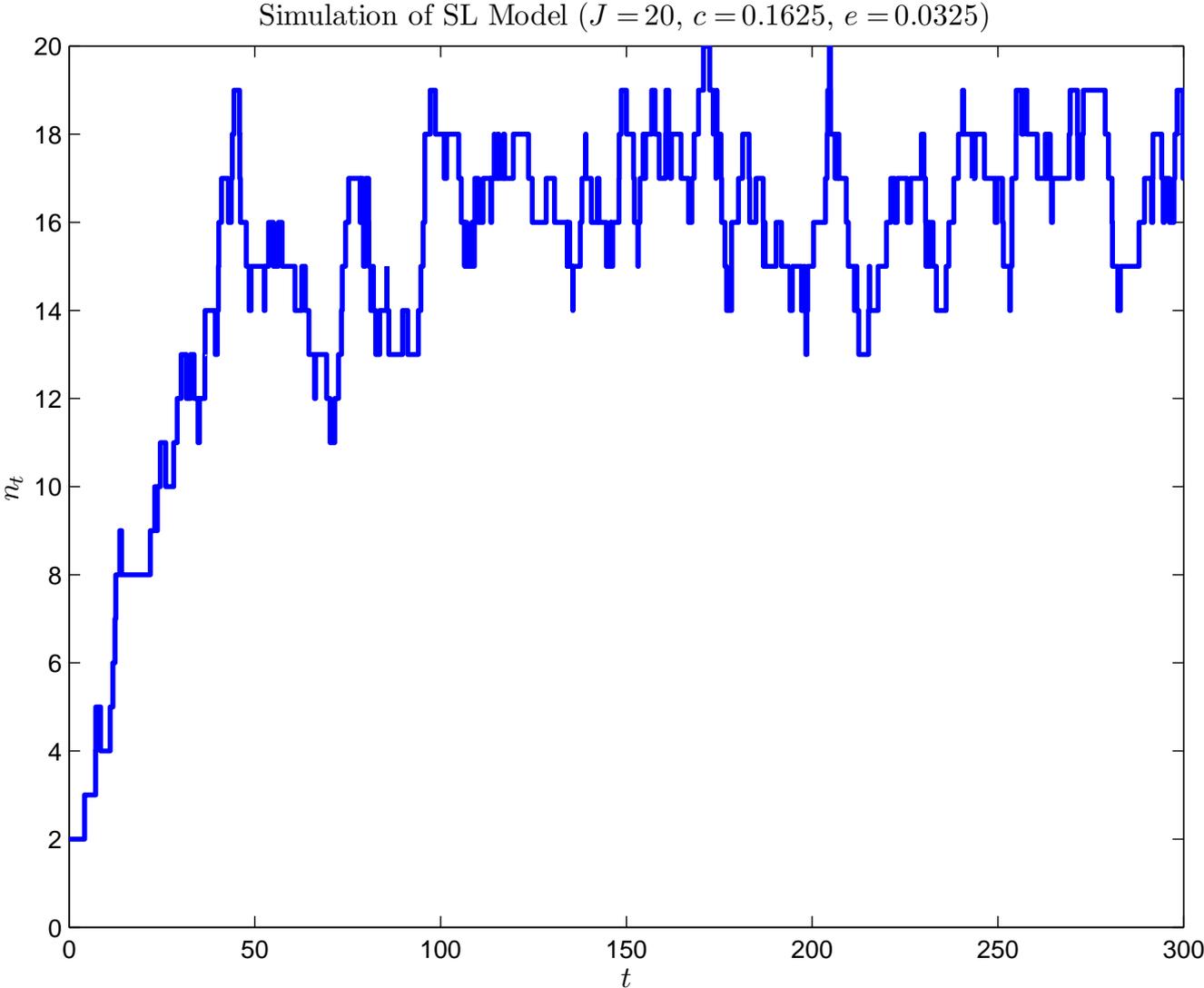
I will call this model the *stochastic logistic (SL) model*, though it has many names, having been rediscovered several times since Feller\* proposed it.

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

# The SL model simulation ( $c < e$ )



# The SL model simulation ( $c > e$ )



# The SL model

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For example, drift:

$$\mathbf{E}(n_{t+s} - n_t | n_t) = n_t \left( c - e - c \frac{n_t}{J} \right) s + o(s).$$

# The SL model

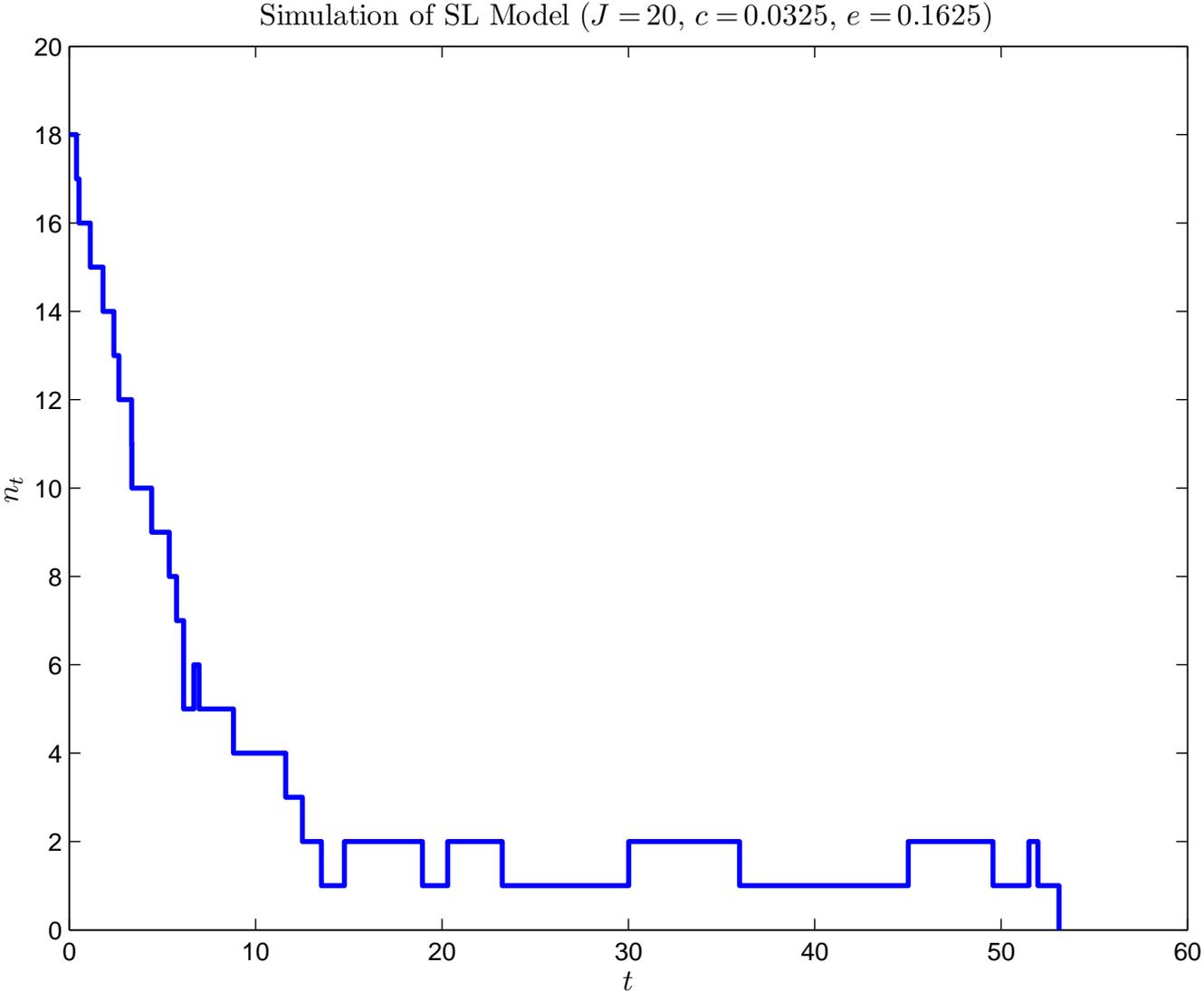
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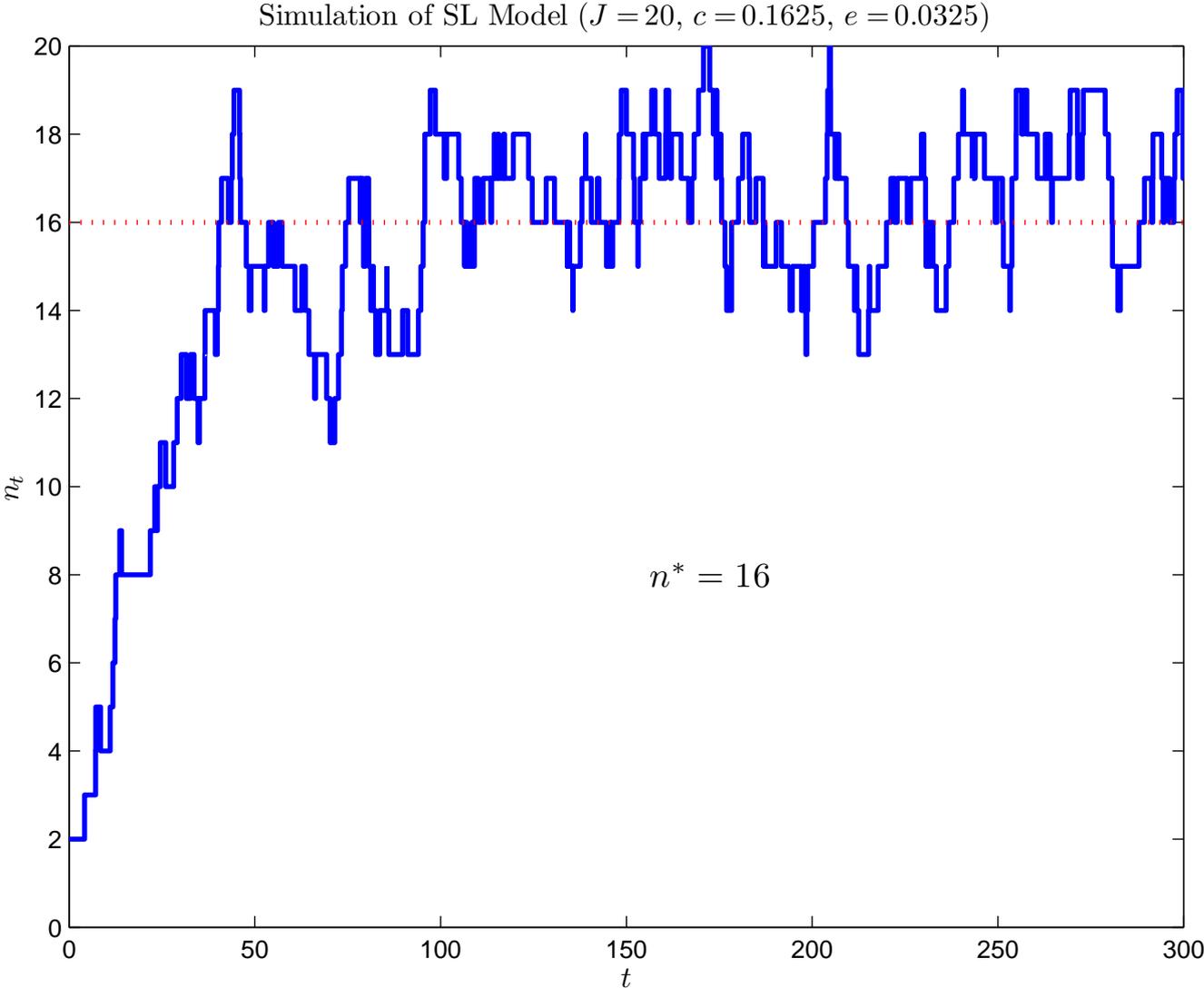
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If  $c > e$ , then the drift is  $< 0$  when  $n_t > n^* := J(1 - e/c)$  and  $> 0$  when  $n_t < n^*$ : the process is “attracted” to  $n^*$ .

# The SL model simulation ( $c > e$ )



# The SL model

Since the SL model is a birth-death process, we have an explicit expression for the *expected time to extinction* starting with  $n$  occupied patches:

$$\tau_n^{(J)} = \frac{1}{e} \sum_{j=1}^n \sum_{k=0}^{J-j} \frac{1}{j+k} \prod_{l=0}^{k-1} \left( \frac{J-j-l}{J\rho} \right),$$

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This expression permits large- $J$  asymptotics . . . .

# The SL model

The following hold in the limit as  $J \rightarrow \infty$ . If  $\rho > 1$  ( $c < e$ ),

$$\tau_1^{(J)} \sim \frac{1}{c} \log \left( \frac{\rho}{\rho - 1} \right)$$

and, for  $n \geq 2$ ,

$$\tau_n^{(J)} \sim \frac{1}{c(\rho - 1)} \left\{ (\rho^n - 1) \log \left( \frac{\rho}{\rho - 1} \right) - \sum_{k=1}^{n-1} \frac{(\rho^{n-k} - 1)}{k} \right\},$$

while if  $\rho < 1$  ( $c > e$ ),

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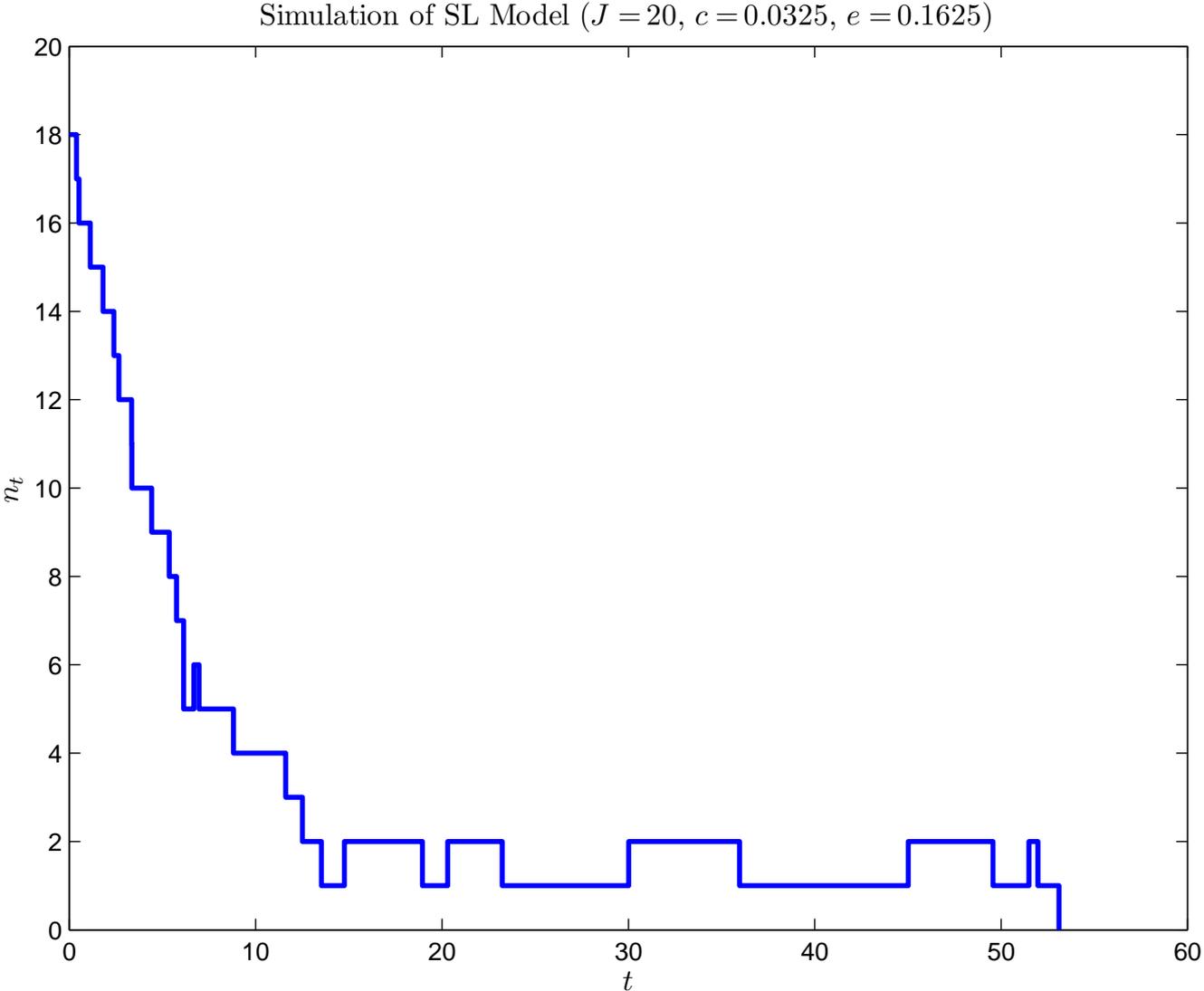
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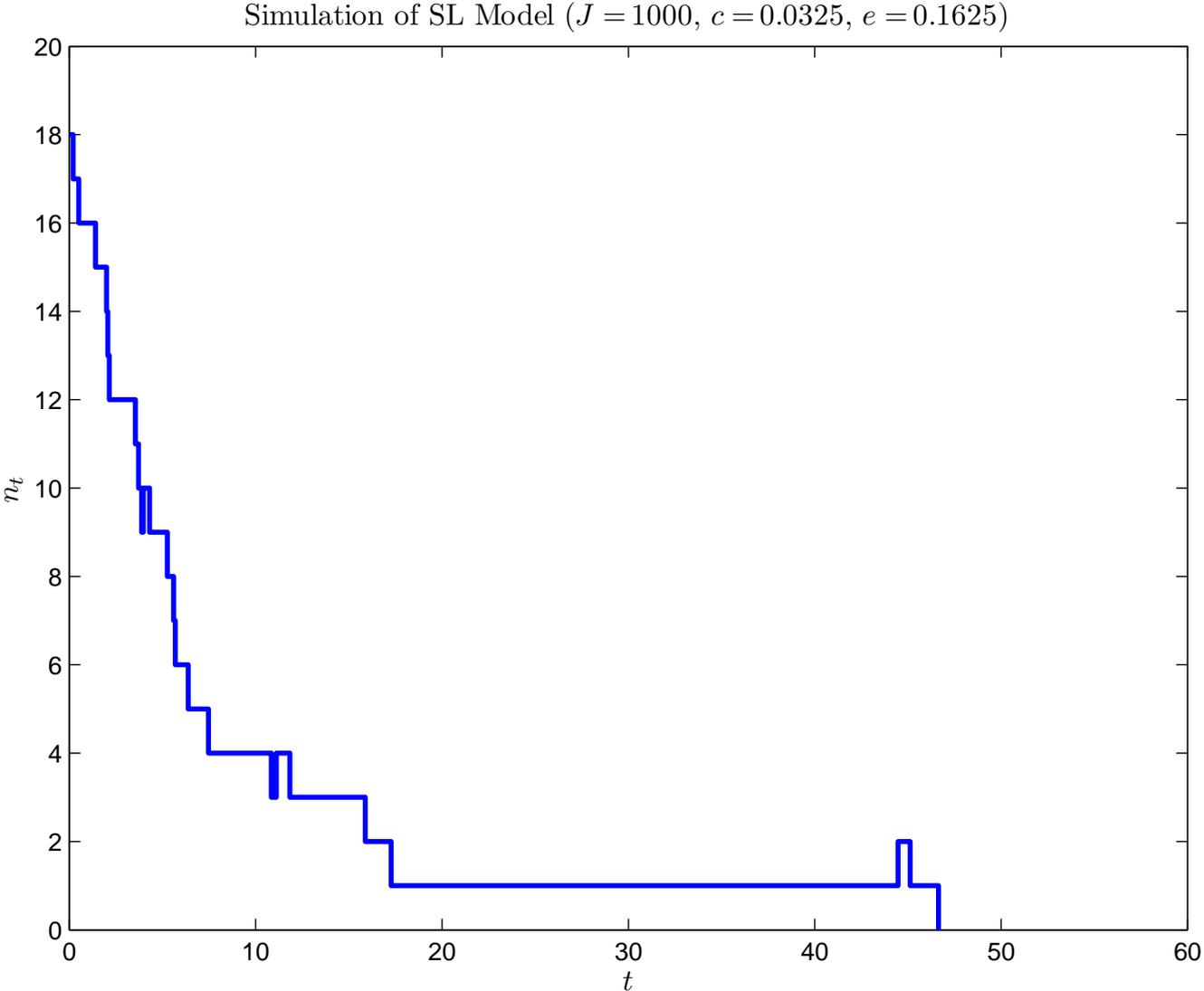
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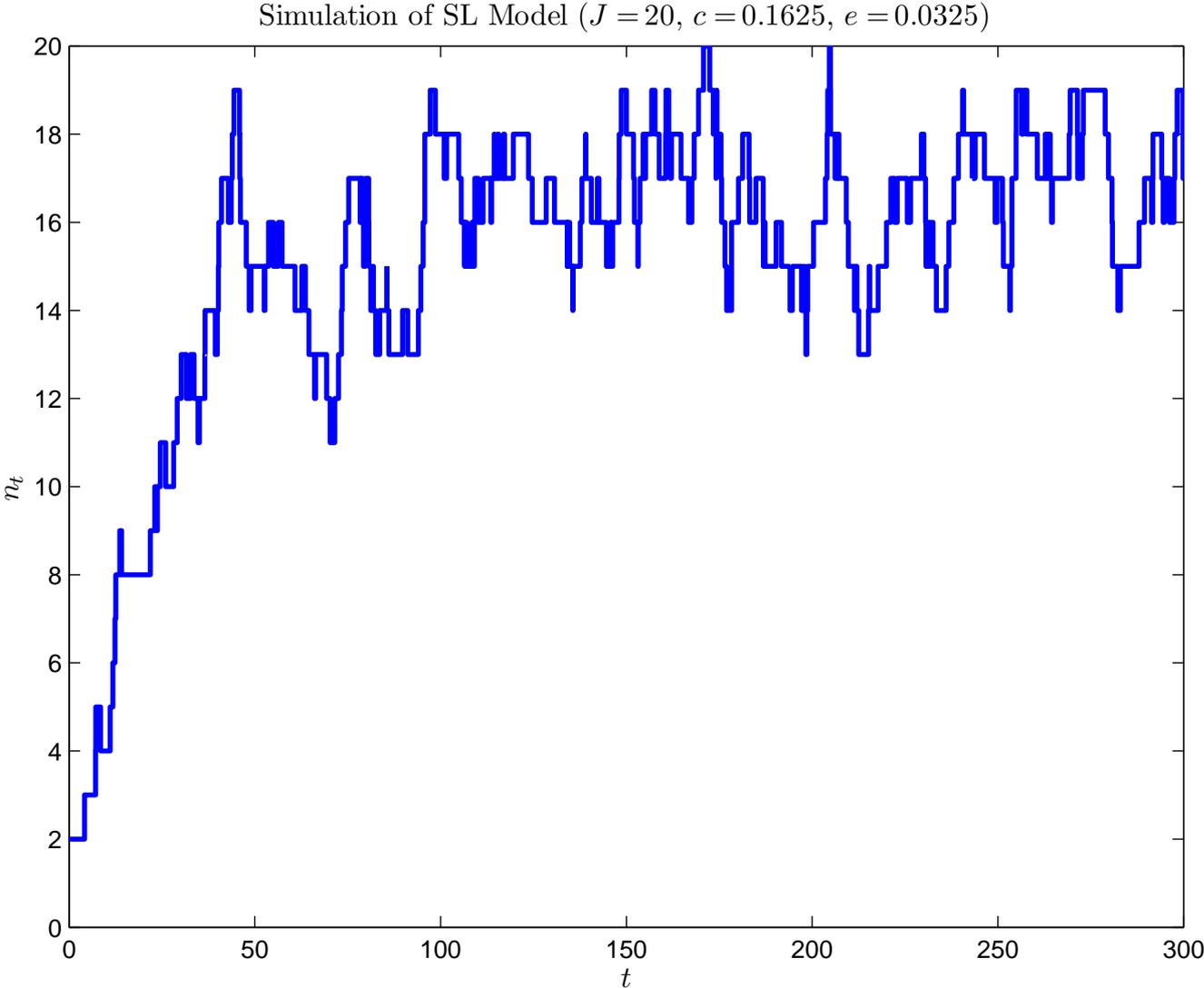
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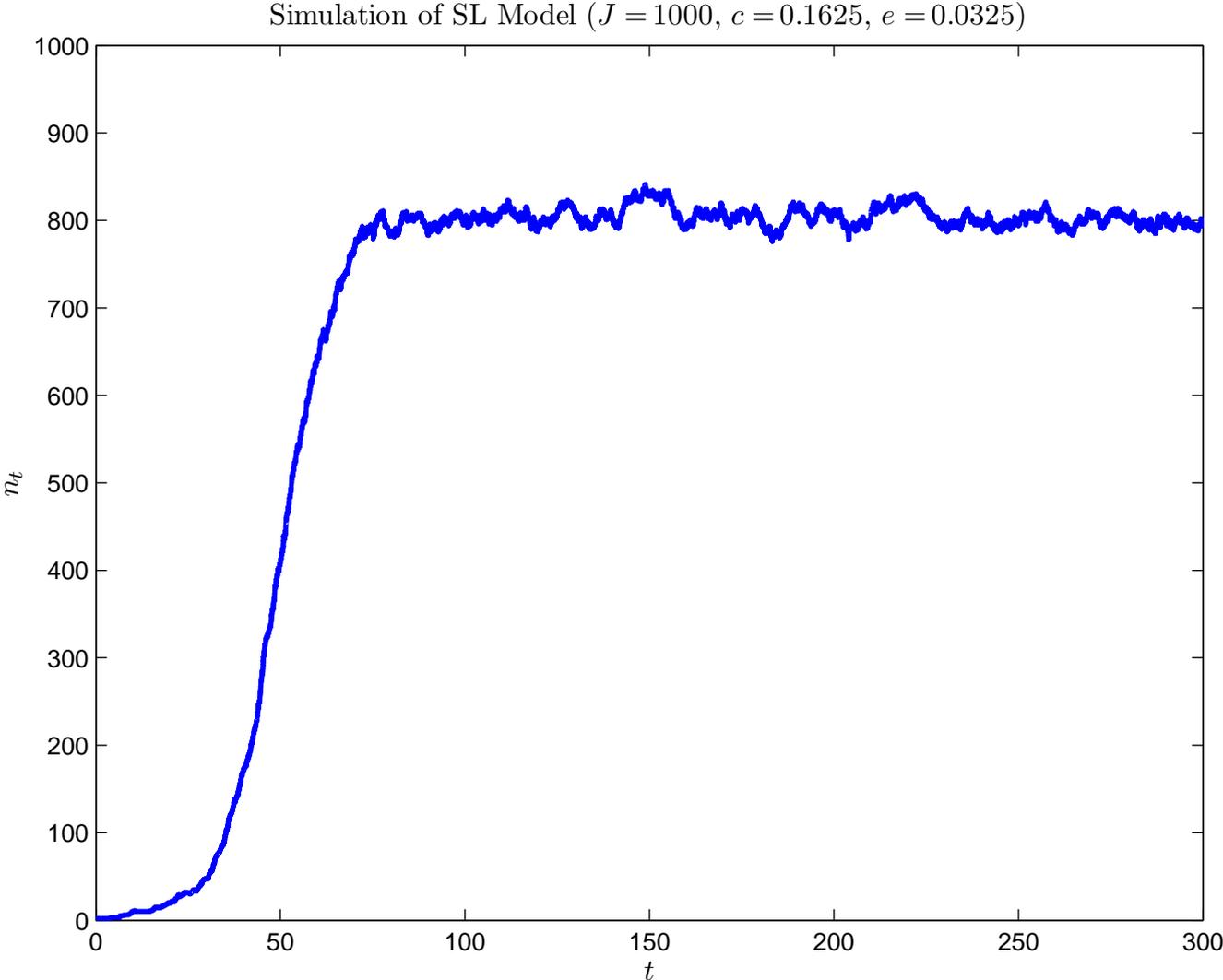
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$$x_t' = cx_t(1 - x_t) - ex_t = cx_t(1 - \rho - x_t),$$

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

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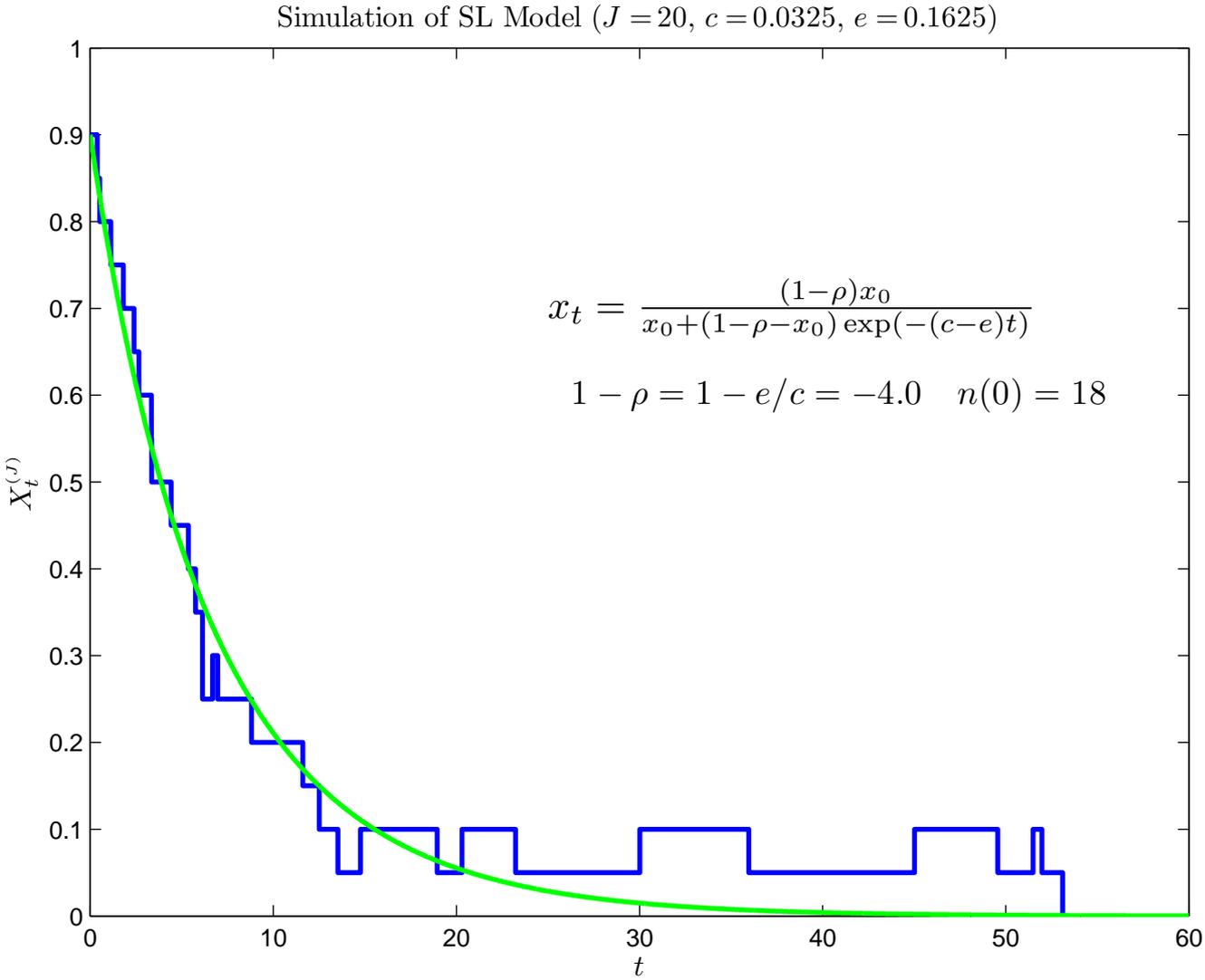
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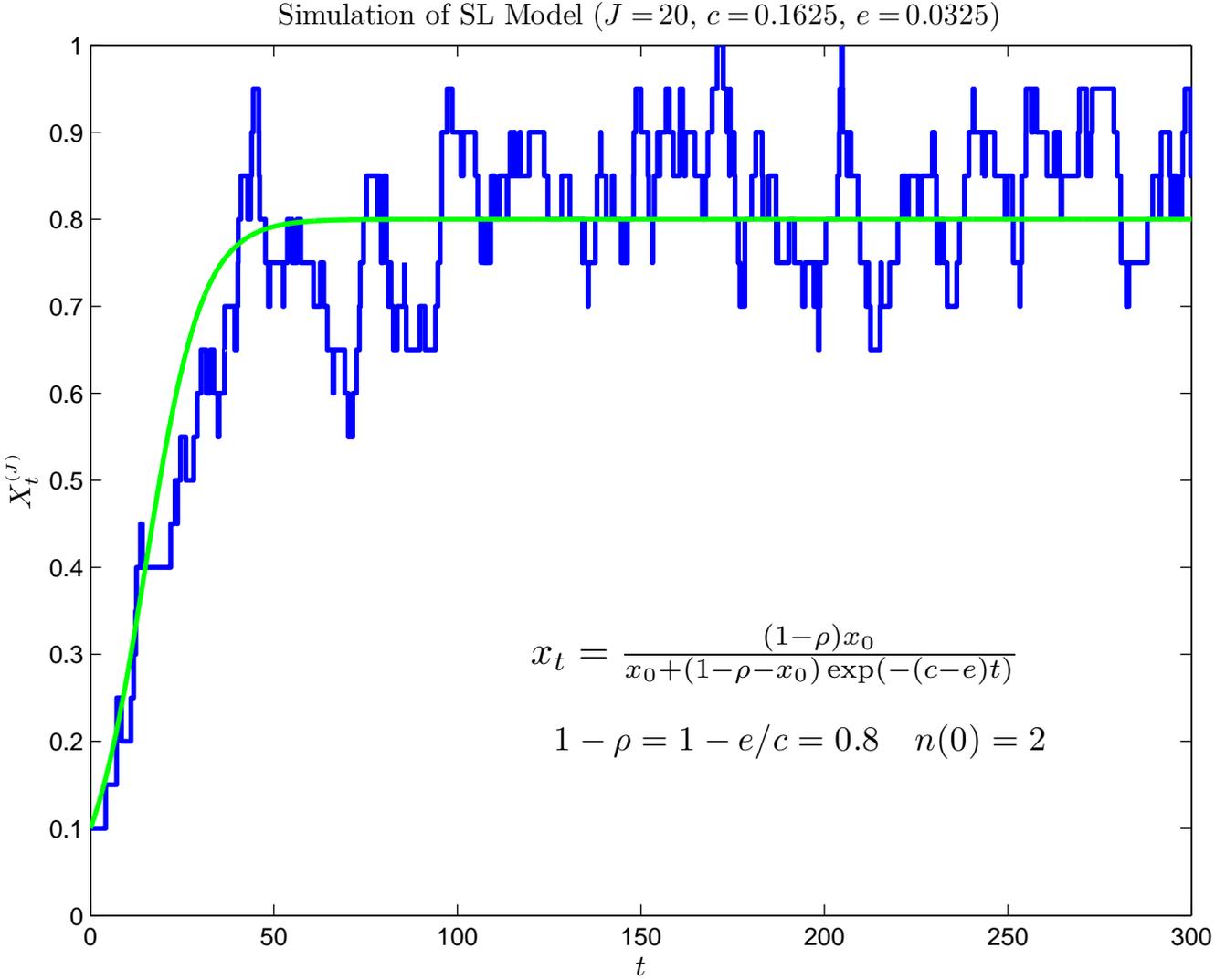
$$x_t = \frac{(1 - \rho)x_0}{x_0 + (1 - \rho - x_0)e^{-(c-e)t}}.$$

There are two equilibria:  $x = 0$  is stable if  $c < e$ , while  $x = 1 - \rho (= 1 - e/c)$  is stable if  $c > e$ .

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



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



# The SL model

This of course is 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.

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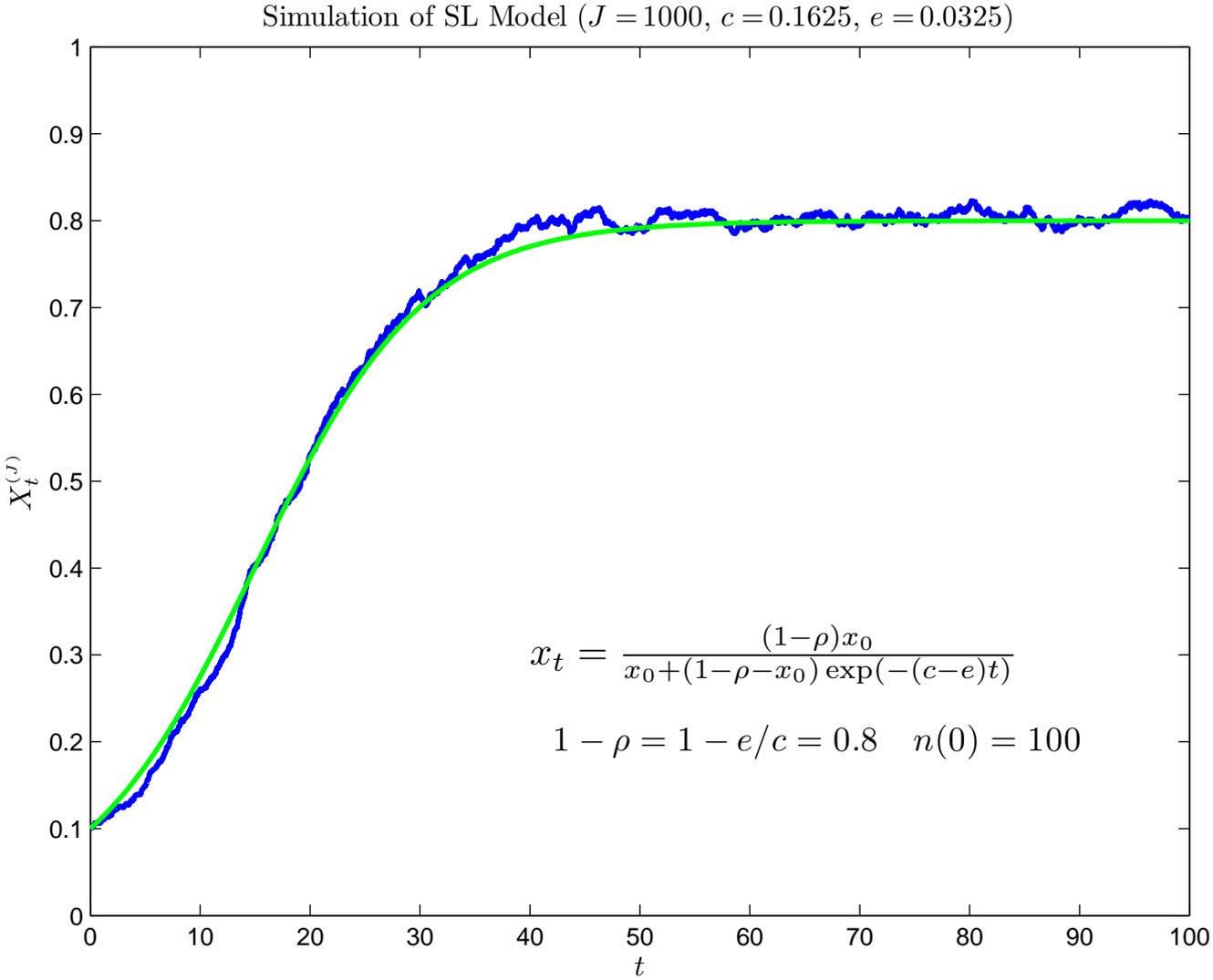
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**Theorem** If  $X_0^{(J)} \rightarrow x_0$  as  $J \rightarrow \infty$ , then the family of processes  $(X_t^{(J)})$  converges *uniformly in probability* on *finite time intervals* to the deterministic trajectory  $(x_t)$ : for every  $\epsilon >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

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

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



# The Mainland-Island model

Recall that there are  $J$  patches. Each occupied patch becomes empty at rate  $e$  and colonization of empty patches occurs at rate  $c/J$  for each suitable pair.

Additionally, immigration from the mainland occurs that rate  $v$ .

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

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

# Network models

We now record the *numbers* of individuals in the various patches: a typical state is  $\mathbf{n} = (n_1, \dots, n_J)$ , where  $n_j$  is the number of individuals in patch  $j$ .

There are two cases: (1) the *open* system, where individuals may enter or leave the network through external immigration and external emigration or removal, and (2) the *closed* system, where there is a *fixed number*  $N$  of individuals circulating.

In the open case individuals are assumed to arrive at patch  $i$  from outside the network as a Poisson stream with rate  $\nu_i$  (if  $\nu_i = 0$  there is no external immigration process at that patch).

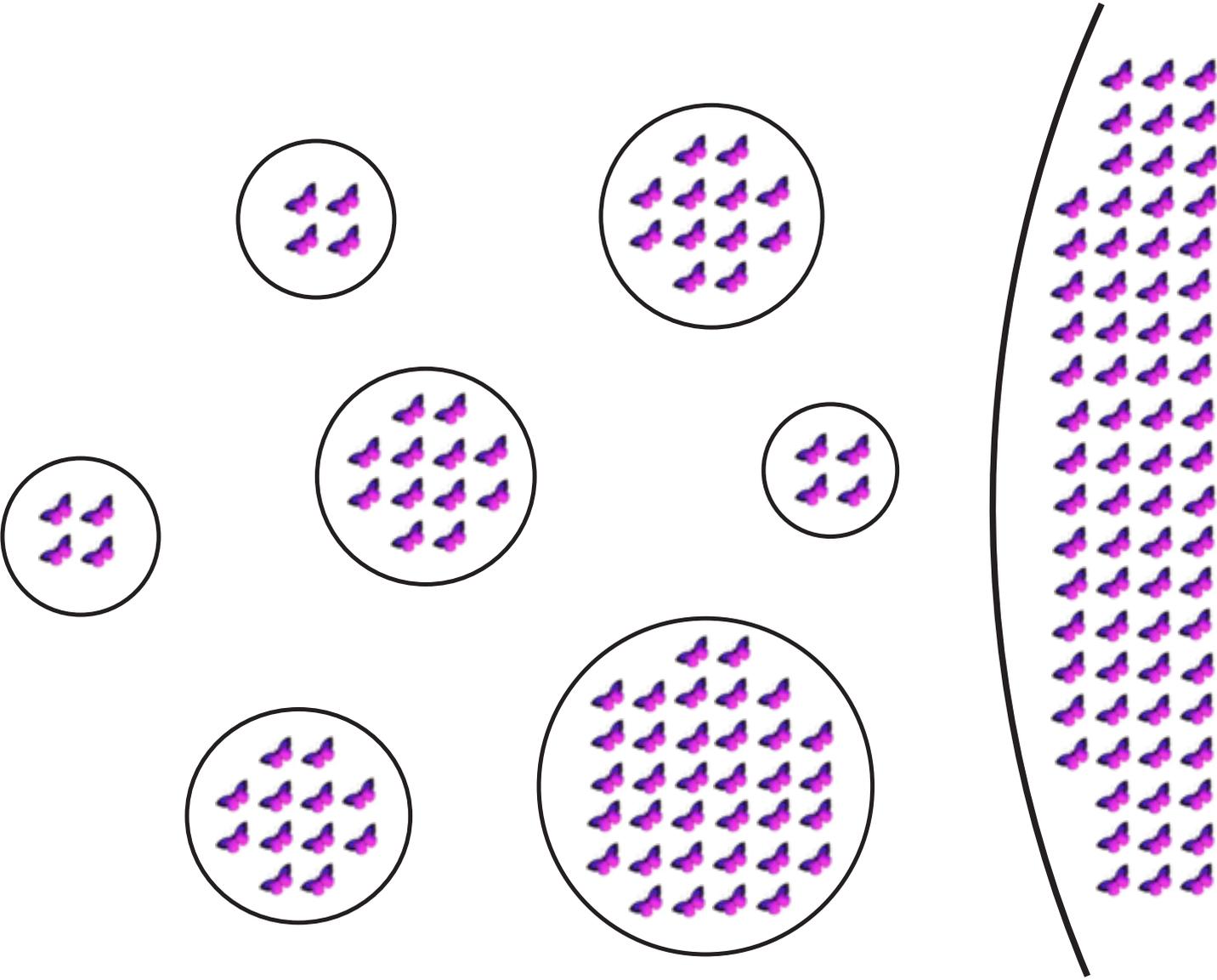
# Network models

We account for spatial structure as follows.

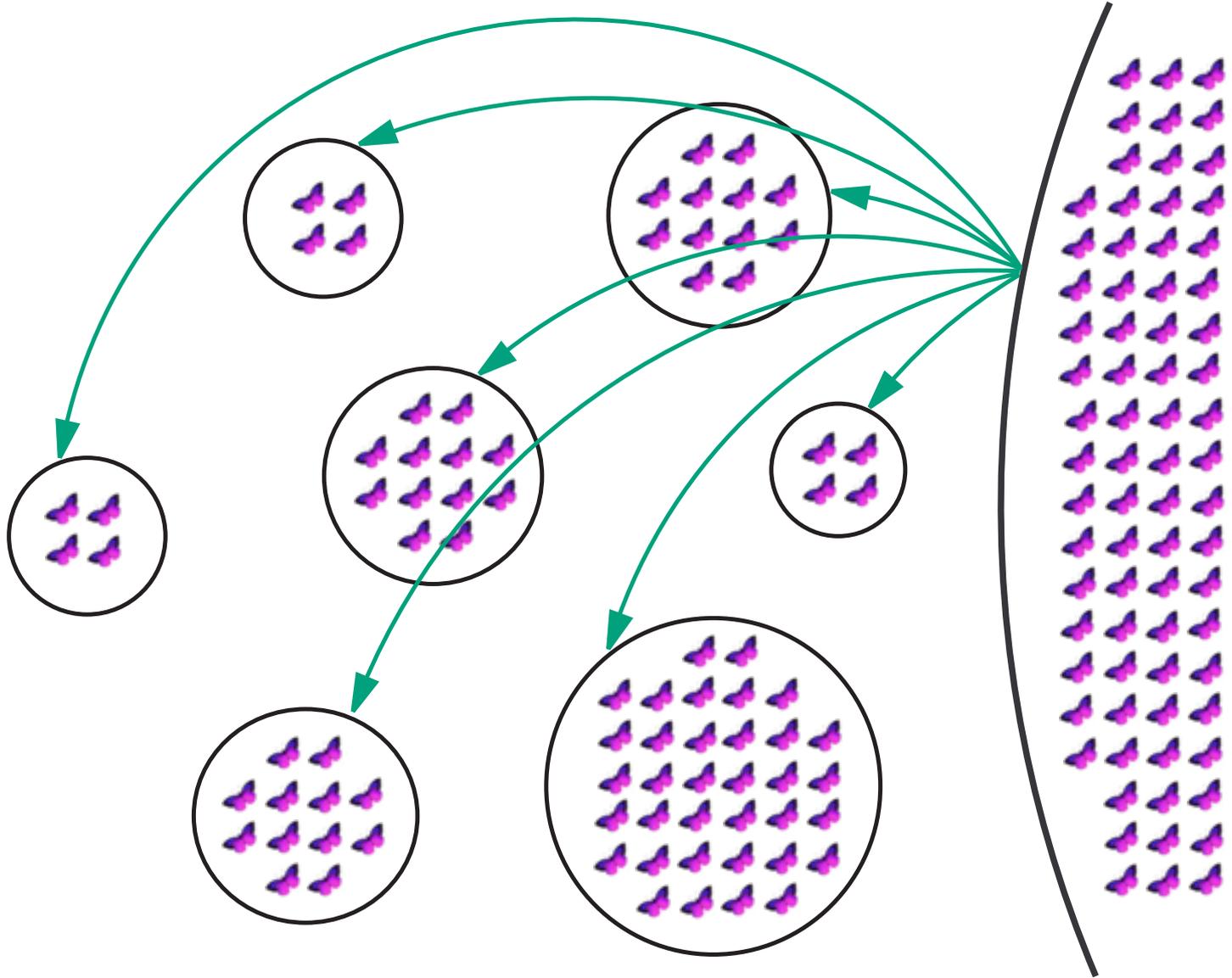
After a sojourn at patch  $i$ , an individual either leaves the network, with probability  $\lambda_{i0}$ , or proceeds to another patch  $j$ , with probability  $\lambda_{ij}$  (in the closed case we take  $\lambda_{i0} = 0$ );  $\lambda_{ij}$  thus specifies the relative proportion of propagules emanating from patch  $i$  that are destined for patch  $j$ ,  $\lambda_{i0}$  being the proportion destined to leave the network. For simplicity, we set  $\lambda_{ii} = 0$ . Clearly  $\sum_j \lambda_{ij} = 1$ .

The matrix  $\Lambda = (\lambda_{ij})$  is termed the *routing matrix*.

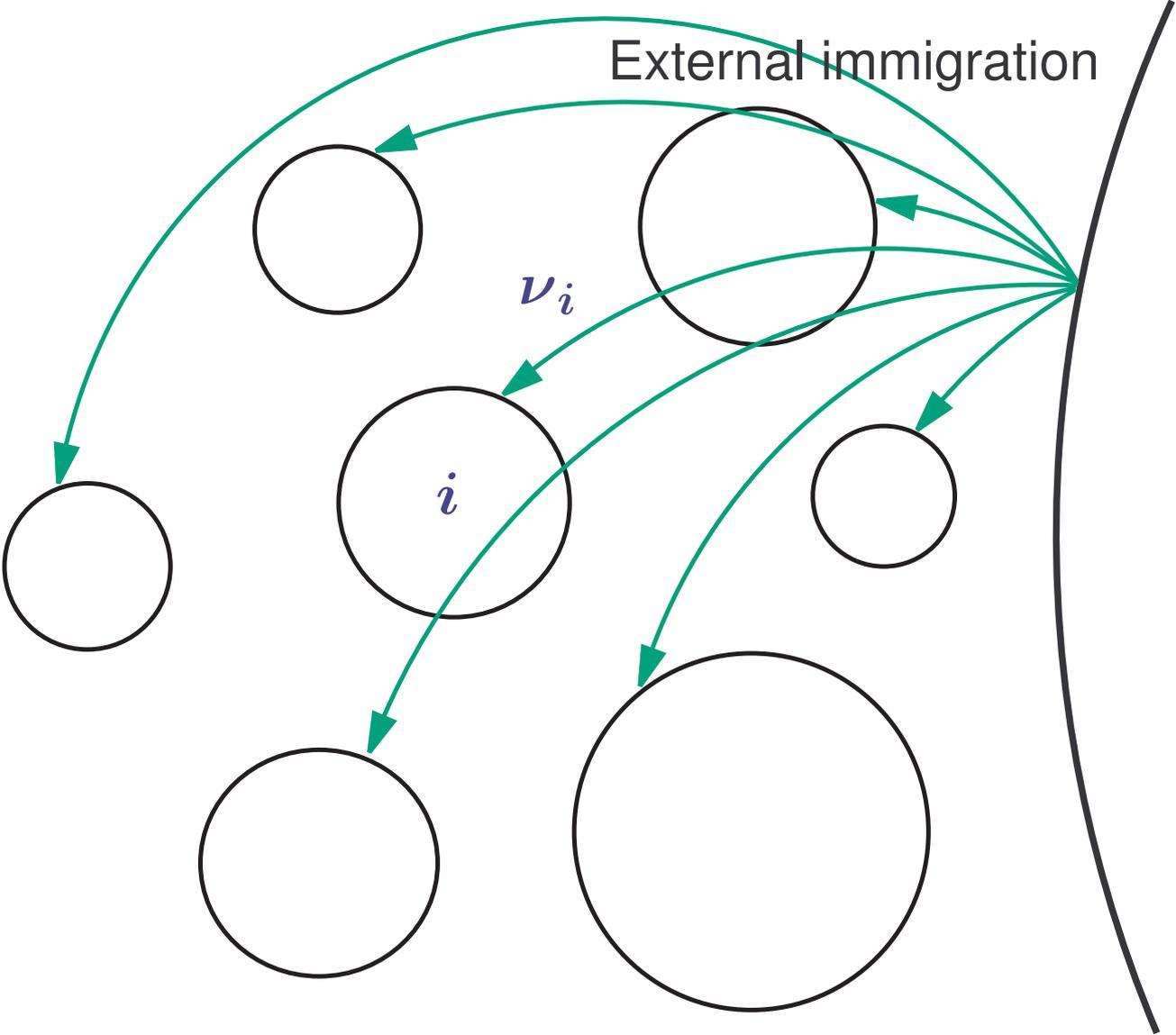
# Open network



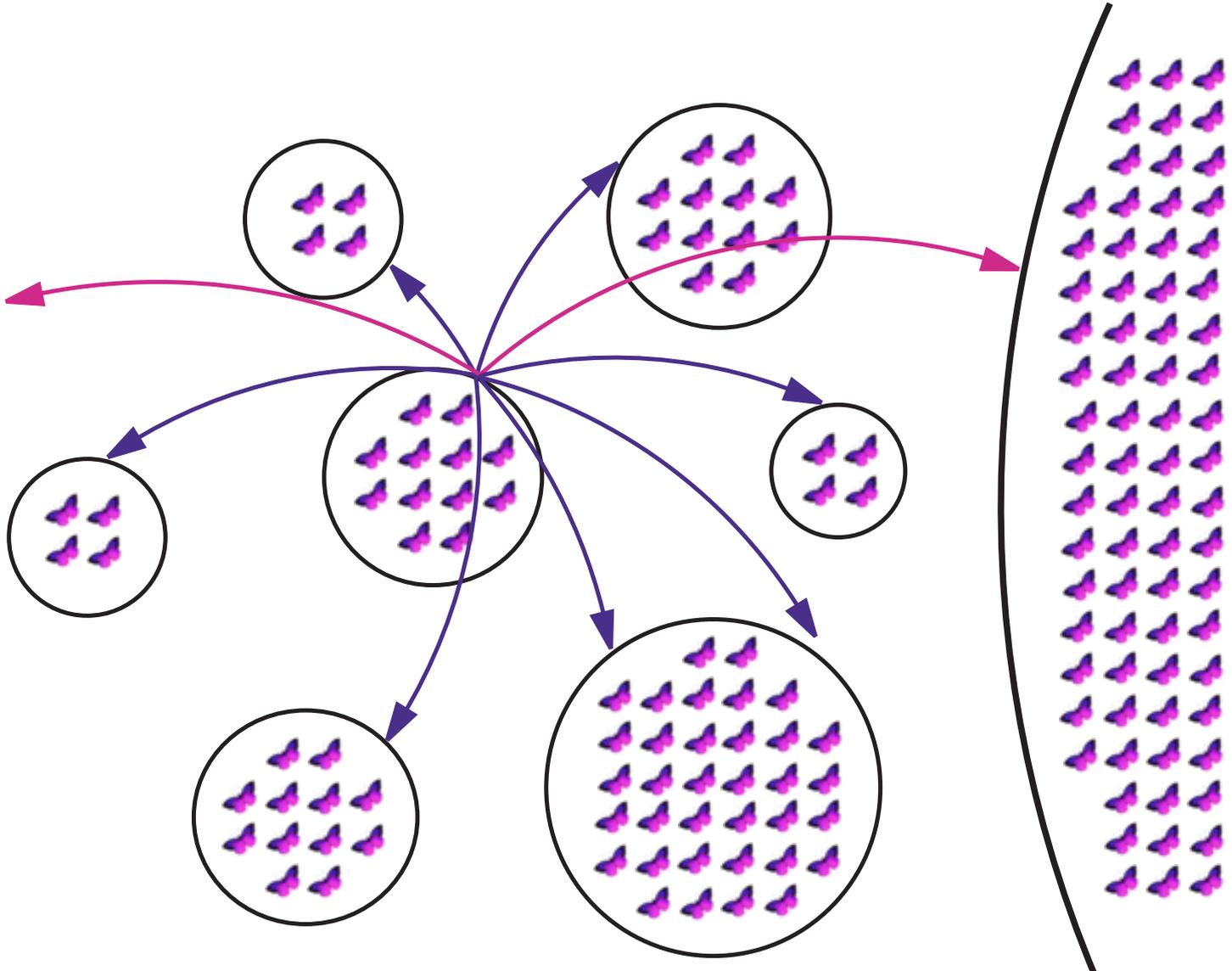
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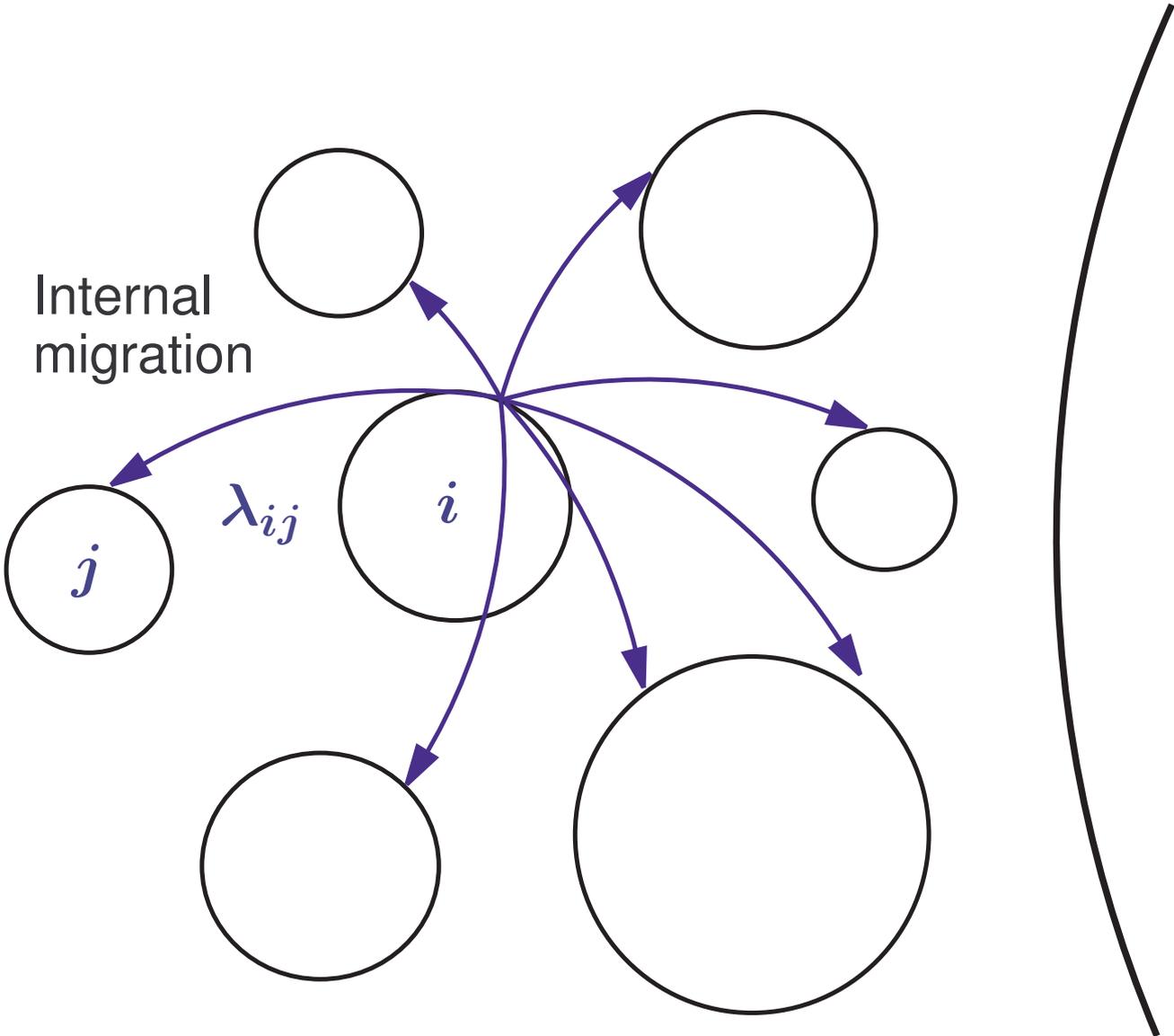
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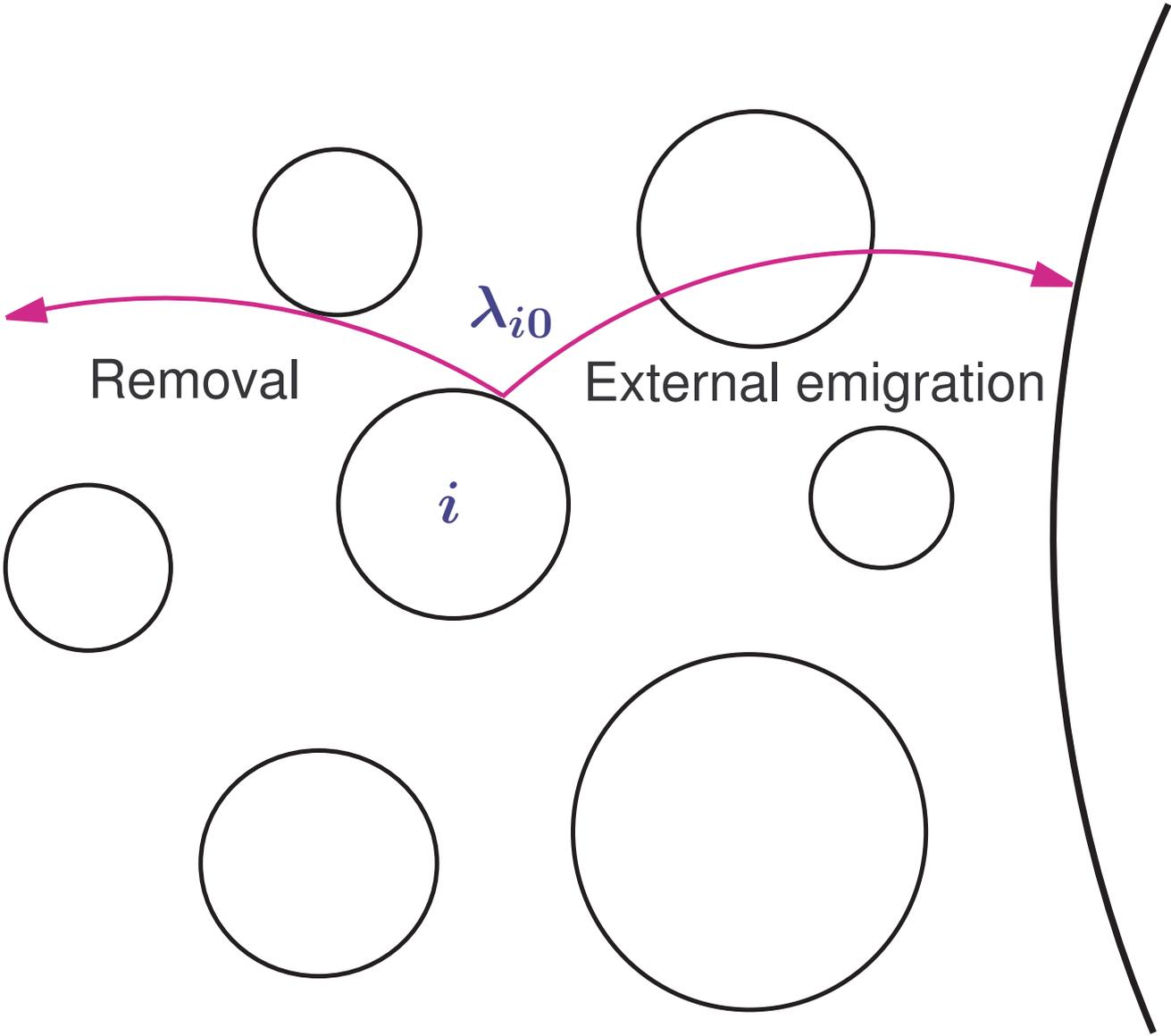
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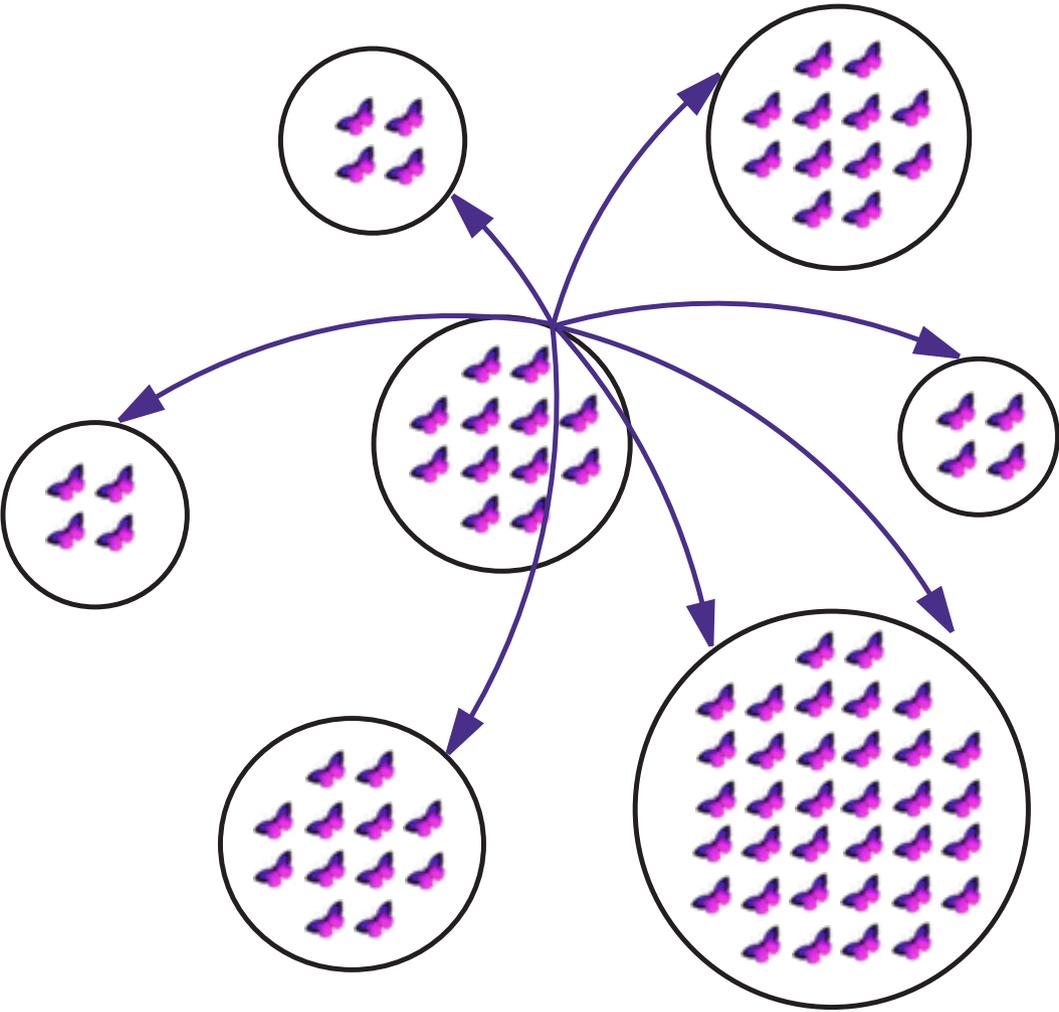
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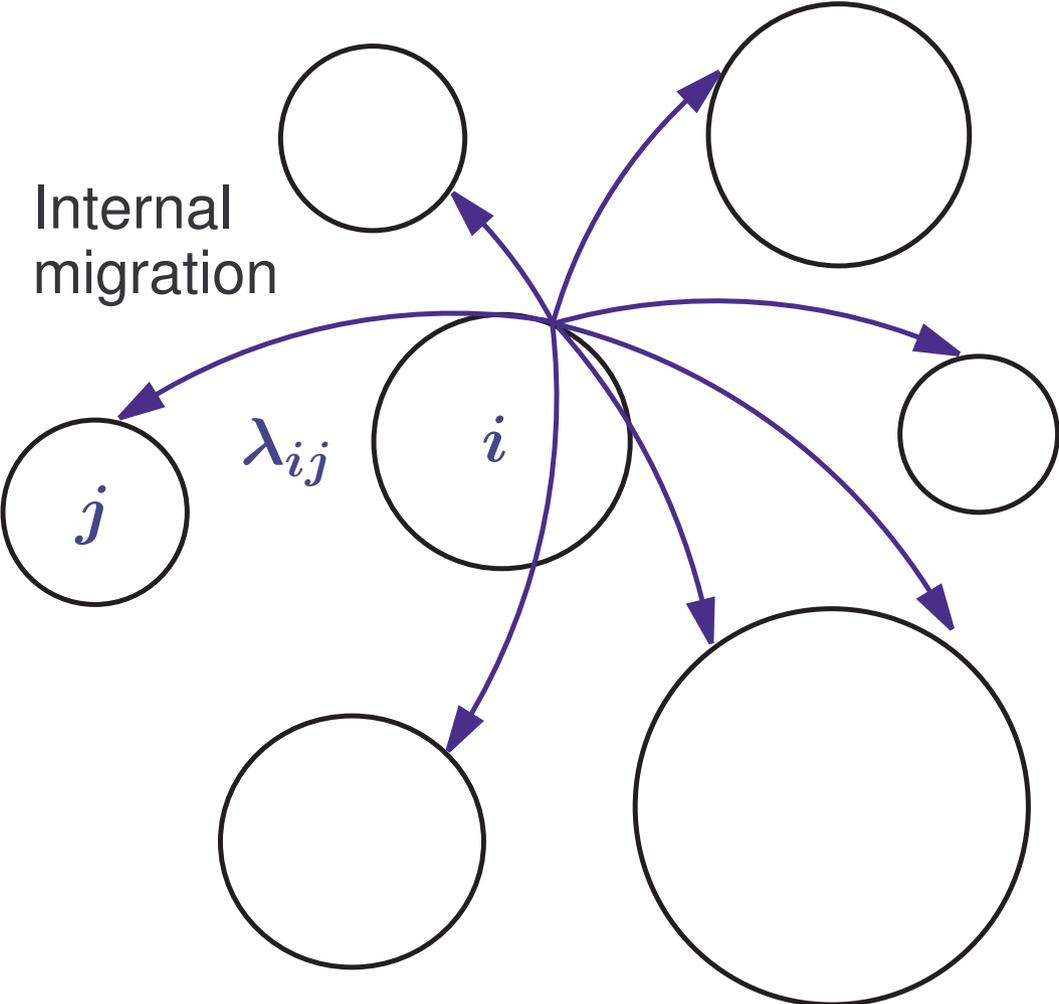
# Open network



# Closed network



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# Network models

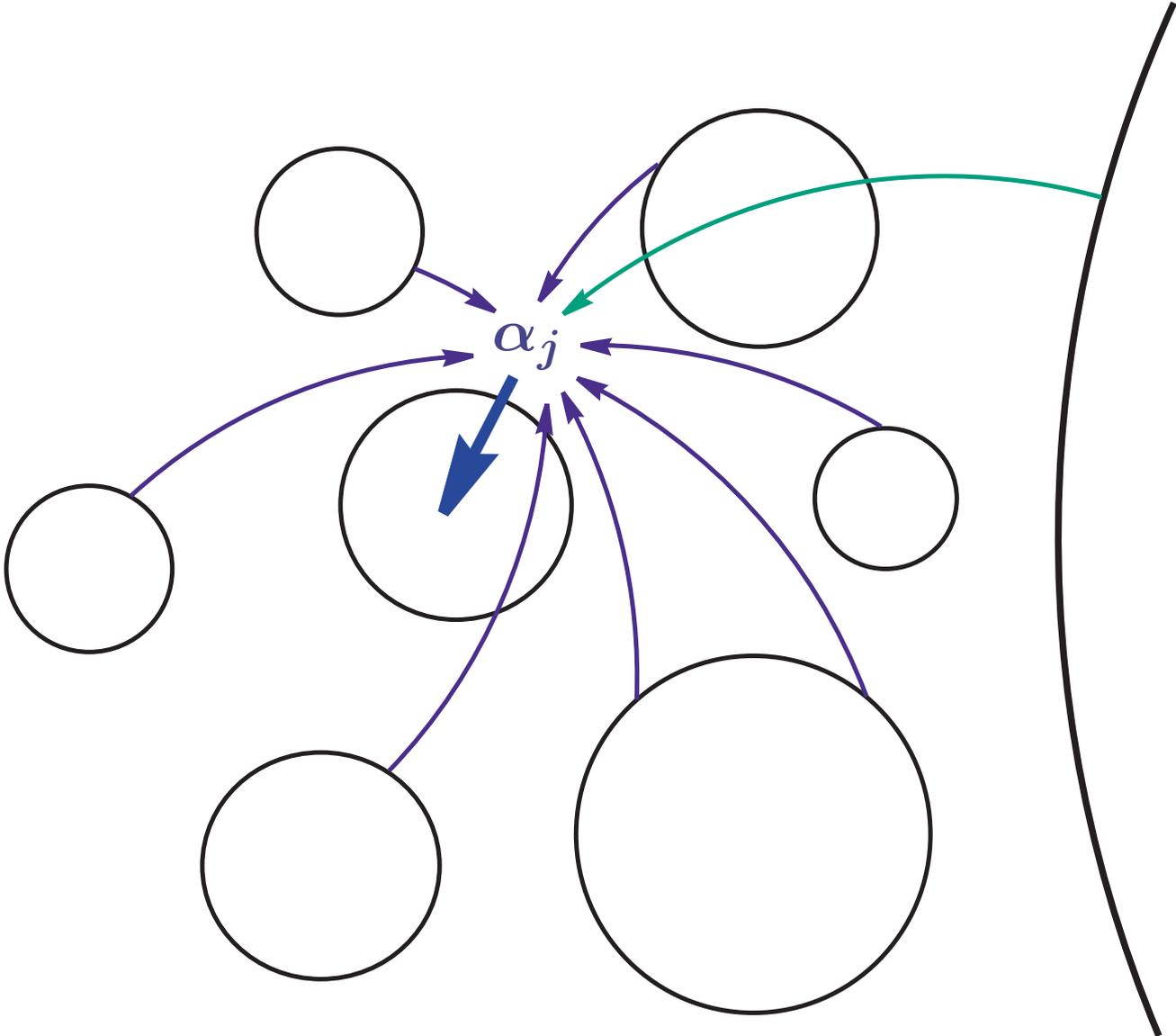
Again for simplicity, we shall assume that  $\Lambda$  is chosen so that an individual can reach any patch from anywhere in the network. In the open case we shall also assume that an individual can reach any patch from outside the network and eventually leave the network starting from anywhere.

In the closed case these conditions ensure that  $\Lambda$  is irreducible and, hence, that there is a unique collection  $(\alpha_1, \dots, \alpha_J)$  of strictly positive numbers which satisfy the *traffic equations*  $\alpha_j = \sum_i \alpha_i \lambda_{ij}$ ,  $j = 1, \dots, J$  (cf. Kirchhoff's law). Here we may assume without loss of generality that  $\sum_j \alpha_j = 1$ .

# Network models

In the open case these conditions ensure that there is a unique positive solution  $(\alpha_1, \dots, \alpha_J)$  to the equations  $\alpha_j = \nu_j + \sum_i \alpha_i \lambda_{ij}$ ,  $j = 1, \dots, J$ . In this case  $\alpha_j$  is the arrival rate at patch  $j$ , while in the closed case  $\alpha_j$  is *proportional to* the arrival rate at patch  $j$ .

# Open network



# Network models: propagation

When there are  $n$  individuals at patch  $j$ , propagation occurs at rate  $\phi_j(n)$  (an arbitrary function for each patch). We assume that  $\phi_j(0) = 0$  and  $\phi_j(n) > 0$  whenever  $n \geq 1$ .

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- $\phi_j(n) = \phi_j$  ( $n \geq 1$ ): the propagation rate is  $\phi_j$ , irrespective of how many individuals are present;
- $\phi_j(n) = \phi_j n$ : the propagation rate at patch  $j$  is proportion to the number of individuals present;
- $\phi_j(n) = \phi_j \min\{n, s_j\}$  ( $n \geq 1$ ): the propagation rate is proportion to the number of individuals present, but there is a fixed maximum rate.

# Network models

I have described the *migration process* of Whittle\*.

\*Whittle, P. (1967) Nonlinear migration processes. Bull. Inst. Int. Statist. 42, 642–647.  
(Constant rates: Jackson, R.R.P. (1954) Queueing systems with phase-type service. Operat. Res. Quart. 5, 109–120.)

The Markov chain  $(\mathbf{n}(t), t \geq 0)$  has state space  $S = Z_+^J$  in open case and transition rates

$$q(\mathbf{n}, \mathbf{n} + \mathbf{e}_j) = \nu_j \quad (\text{external arrival at patch } j)$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i) = \phi_i(n_i) \lambda_{i0} \quad (\text{removal from patch } i)$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i + \mathbf{e}_j) = \phi_i(n_i) \lambda_{ij} \quad (\text{migration from } i \text{ to } j).$$

( $\mathbf{e}_j$  is the unit vector in  $Z_+^J$  with a 1 as its  $j$ -th entry)

# Network models

In the closed case we simply have

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i + \mathbf{e}_j) = \phi_i(n_i) \lambda_{ij} \quad (\text{migration from } i \text{ to } j),$$

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The equilibrium behaviour of migration processes is well understood (but apparently not by ecologists).

Let  $\pi(\mathbf{n})$  be the equilibrium probability of configuration  $\mathbf{n} = (n_1, \dots, n_J)$ .

# Open migration process

**Theorem** An equilibrium distribution exists if

$$b_j^{-1} := 1 + \sum_{n=1}^{\infty} \frac{\alpha_j^n}{\prod_{r=1}^n \phi_j(r)} < \infty \quad \text{for all } j,$$

in which case

$$\pi(\mathbf{n}) = \prod_{j=1}^J \pi_j(n_j), \quad \text{where} \quad \pi_j(n) = b_j \frac{\alpha_j^n}{\prod_{r=1}^n \phi_j(r)}.$$

Thus, in equilibrium,  $n_1, \dots, n_J$  are *independent* and each patch  $j$  behaves *as if* it were isolated with Poisson input at rate  $\alpha_j$ .

# Open migration process: examples

(1)  $\phi_j(n) = \phi_j$  ( $n \geq 1$ ). If  $\rho_j := \alpha_j / \phi_j < 1$ ,

$$\pi_j(n) = (1 - \rho_j)\rho_j^n \quad \text{(geometric).}$$

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(3)  $\phi_j(n) = \phi_j \min\{n, s_j\}$  ( $n \geq 1$ ). If  $\rho_j := \alpha_j / (s_j \phi_j) < 1$ ,

$$\pi_j(n) = \pi_j(0) \frac{(s_j \rho_j)^n}{n!} \quad (n = 1, \dots, s_j)$$

$$\pi_j(n) = \pi_j(s) \rho_j^{n-s_j} \quad (n = s_j + 1, \dots).$$

# Closed migration process ( $N$ individuals)

**Theorem** An equilibrium distribution always exists and is given by

$$\pi^{(N)}(\mathbf{n}) = B^{(N)} \prod_{j=1}^J \frac{\alpha_j^{n_j}}{\prod_{r=1}^{n_j} \phi_j(r)} \quad (\mathbf{n} \in S^{(N)}),$$

where  $B^{(N)}$  is a normalizing constant chosen so that  $\pi^{(N)}$  sums to 1 over  $S^{(N)}$ .

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Note that  $n_1, \dots, n_J$  are *not* independent.

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$$(1) \phi_j(n) = \phi_j \quad (n \geq 1).$$

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The marginal distribution of the number  $n_j$  at patch  $j$  is messy (the form depends on which of the  $\rho_i$ 's are distinct).

# Closed migration process: examples

$$(2) \quad \phi_j(n) = \phi_j n .$$

The equilibrium distribution is *multinomial*:

$$\pi^{(N)}(\mathbf{n}) = \frac{N!}{n_1! n_2! \cdots n_J!} p_1^{n_1} p_2^{n_2} \cdots p_J^{n_J} \quad (\mathbf{n} \in S^{(N)}),$$

where  $p_i = r_i / \sum_{j=1}^J r_j$  and  $r_i = \alpha_i / \phi_i$ .

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where  $p_i = r_i / \sum_{j=1}^J r_j$  and  $r_i = \alpha_i / \phi_i$ .

The marginal distribution of the number  $n_j$  at patch  $j$  is *binomial*:

$$\pi_j^{(N)}(n) = \binom{N}{n} p_j^n (1 - p_j)^{N-n} \quad (n = 0, 1, \dots, N).$$

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For each of the network models—*but where there is homogeneity among the patches*—what is the corresponding/appropriate patch-occupancy model?

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Do we recover the SL model?

Recall that  $n_t$  was the number of occupied patches at time  $t$ , that local extinction occurred at common rate  $e$  and that colonization occurred at common rate  $c/J$  for each of the  $n(J - n)$  occupied-unoccupied pairs:

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

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Even in the epidemiological literature\*, where the SL model—called the Susceptible-Infective-Susceptible (SIS) model—is ubiquitous, there is still controversy about interpretation of the ingredients of the model.

\*Begon, M., Bennett, M., Bowers, R.G., French, N.P., Hazel, S.M. and Turner, J. (2002) A clarification of transmission terms in host-microparasite models: numbers, densities and areas. *Epidemiology and Infection* 129, 147–153.

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We will use the various network models to find out.  
There are some surprises.

# Which patch-occupancy model?

**Symmetric networks** Suppose that  $\phi_j(n) = \phi(n)$  for all  $j$  (all patches produce propagules at the same rate). We consider two cases (i) “**constant**”  $\phi(n) = \phi$  ( $n \geq 1$ ) (constant propagation rate  $\phi$ ) and (ii) “**linear**”  $\phi(n) = \phi n$  ( $\phi$  is the *per-capita* propagation rate).

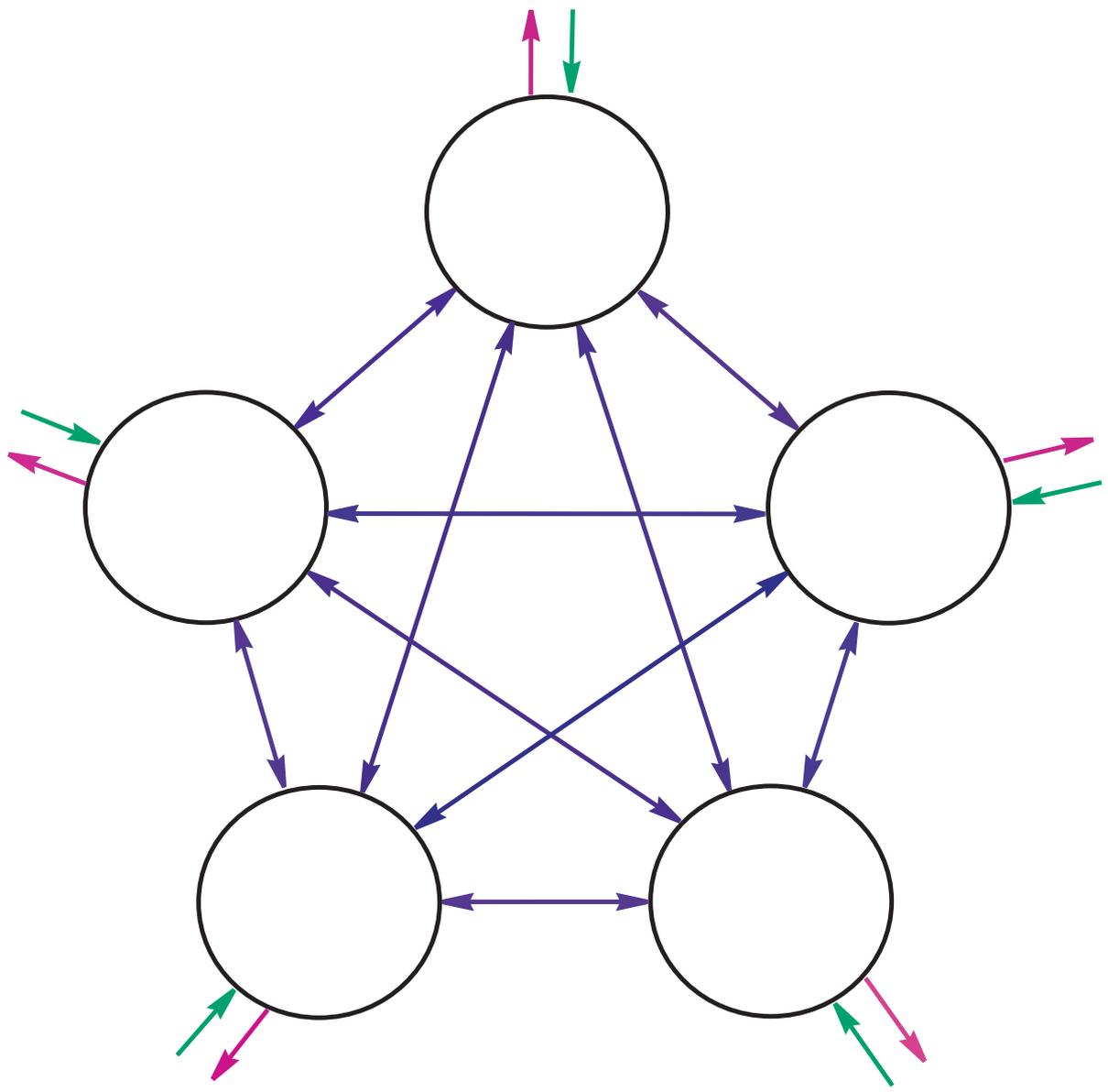
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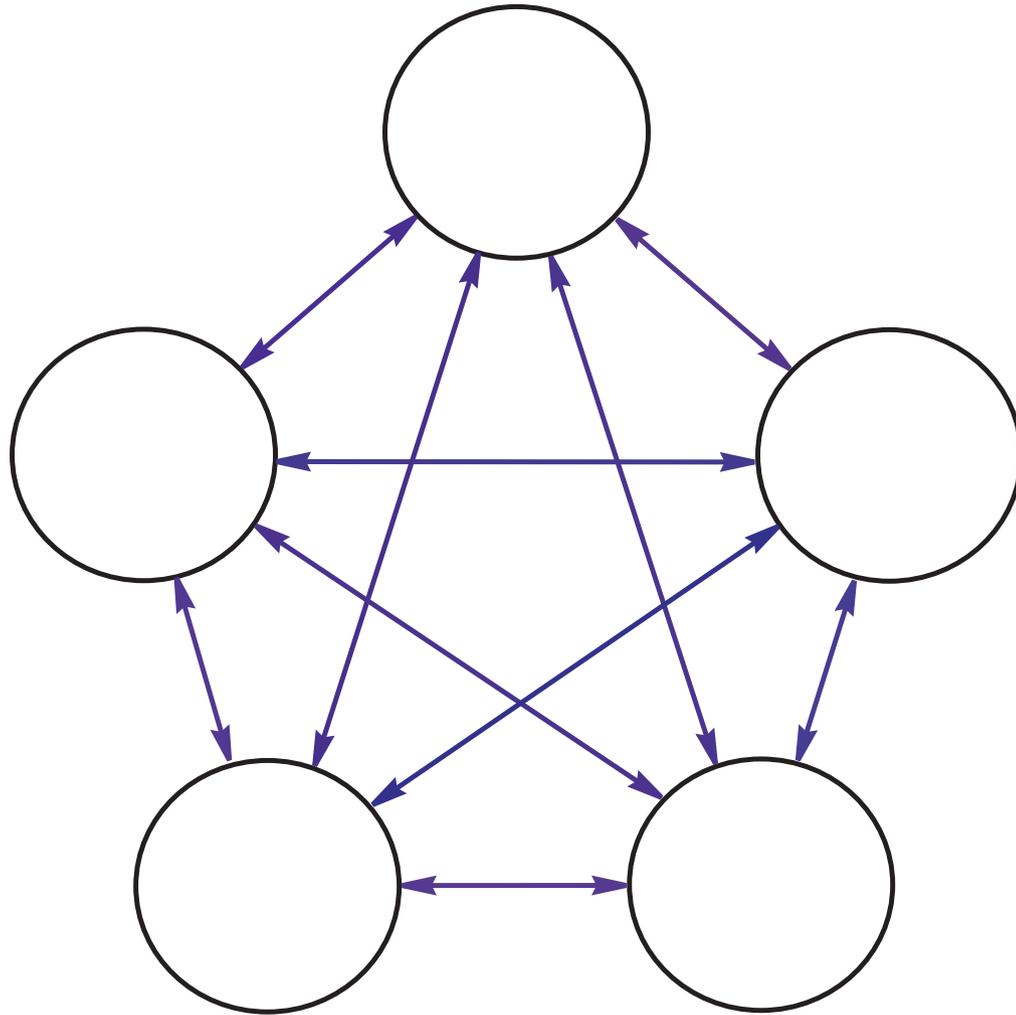
We will also suppose that emigration *to* any patch  $j$  is the same *from* all patches  $i$ :  $\lambda_{ij} = 1/(J - 1)$  in the closed network, and,  $\nu_i = \nu$ ,  $\lambda_{i0} = \lambda_0$  and  $\lambda_{ij} = (1 - \lambda_0)/(J - 1)$  in the open network.

This is sufficient for  $\alpha_j (= \alpha)$  to be the same for all  $j$ :  $\alpha = 1/J$  (closed network) and  $\alpha = \nu/\lambda_0$  (open network).

# Symmetric network (open)



# Symmetric network (closed)



# Which patch-occupancy model?

We will evaluate

- (i) the equilibrium expected colonization rate  $c(m)$ , that is, *the expected arrival rate at unoccupied patches, conditional on there being  $m$  patches occupied*, and,
- (ii) the equilibrium expected local extinction rate  $e(m)$ , that is, *the expected rate at which empty patches appear, conditional on there being  $m$  patches occupied*.

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We might expect that, for some  $c$ ,  $e$  and  $v$ ,

$$(i) \ c(m) = v(J - m) + \frac{c}{J}m(J - m) \quad \text{and} \quad (ii) \ e(m) = em.$$

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External colonization

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Ross (2008)?

We might expect that, for some  $c$ ,  $e$  and  $v$ ,

(i)  $c(m) = v(J - m) + \frac{\phi}{J}m(J - m)$  and (ii)  $e(m) = em$ .

# Which patch-occupancy model?

Let  $C(\mathbf{n}) = \sum_k 1_{\{n_k(t) > 0\}}$  be the number of occupied patches when the network is in state  $\mathbf{n}$ . Then,

$$\begin{aligned} c(m) &= \mathbf{E} \left( \sum_j \left( \nu_j + \sum_{i \neq j} \phi_i(n_i(t)) \lambda_{ij} \right) 1_{\{n_j(t)=0\}} \middle| C(\mathbf{n}) = m \right) \\ &= \sum_j \nu_j \Pr(n_j(t) = 0 | C(\mathbf{n}) = m) \\ &\quad + \sum_j \sum_{i \neq j} \mathbf{E} \left( \phi_i(n_i(t)) 1_{\{n_j(t)=0\}} \middle| C(\mathbf{n}) = m \right) \lambda_{ij}. \end{aligned}$$

(open network)

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# Which patch-occupancy model?

Owing to the symmetry ...

$$c(m) = J\nu \Pr(n_1(t) = 0 | C(\mathbf{n}) = m)$$

$$+ J(J-1) \mathbf{E} \left( \phi(n_1(t)) 1_{\{n_2(t)=0\}} | C(\mathbf{n}) = m \right) \frac{1 - \lambda_0}{J-1}$$

$$= J\nu \left( 1 - \frac{m}{J} \right) + (1 - \lambda_0) J \mathbf{E} \left( \phi(n_1(t)) 1_{\{n_2(t)=0\}} | C(\mathbf{n}) = m \right)$$

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(closed network)

# Which patch-occupancy model?

And, for both the open and closed network,

$$\begin{aligned} e(m) &= \mathbf{E} \left( \sum_i \phi_i(1) 1_{\{n_i(t)=1\}} \mid C(\mathbf{n}) = m \right) \\ &= \sum_i \phi_i(1) \Pr(n_i(t) = 1 \mid C(\mathbf{n}) = m) \\ &= J\phi \Pr(n_1(t) = 1 \mid C(\mathbf{n}) = m) \end{aligned}$$

# Which patch-occupancy model?

Before proceeding, recall that . . .

## Open network

$J$  – number of patches

$\nu$  – common external immigration rate

$\phi(n)$  – common propagation rate when  $n$  individuals present at that patch – two cases:

“constant”  $\phi(n) = \phi 1_{\{n>0\}}$   $\rho := \nu / (\phi \lambda_0)$  ( $< 1$ )

“linear”  $\phi(n) = \phi n$   $r := \nu / (\phi \lambda_0)$

$\lambda_0$  – common external emigration/removal probability

$\lambda_{ij} = (1 - \lambda_0) / (J - 1)$

# Which patch-occupancy model?

## Closed network

$J$  – number of patches

$N$  – number of individuals (fixed)

$\phi(n)$  – common propagation rate when  $n$  individuals present at that patch – two cases:

“constant”  $\phi(n) = \phi 1_{\{n>0\}}$

“linear”  $\phi(n) = \phi n$

$\lambda_{ij} = 1/(J - 1)$

# Equilibrium distributions

Propagation rates	Open network* $\pi_j(n) \ (n \geq 0)$	Closed network $\pi^{(N)}(\mathbf{n}) \ (\mathbf{n} \in S^{(N)})$
Constant	$(1 - \rho)\rho^n$	$\binom{N + J - 1}{J - 1}^{-1}$
Linear	$e^{-r} \frac{r^n}{n!}$	$\frac{N!}{n_1! n_2! \cdots n_J!} \left(\frac{1}{J}\right)^N$

\* $n_1, \dots, n_J$  are independent

# Which patch-occupancy model? $c(m)$

## Closed constant

$$c(m) = \frac{\phi}{J-1} m(J-m) \quad (m = 1, \dots, J)$$

## Closed linear

$$c(m) = \frac{N\phi}{J-1} (J-m) \quad (m = 1, \dots, J)$$

## Open constant

$$c(m) = \nu(J-m) + \frac{\phi(1-\lambda_0)}{(J-1)(1-\rho)} m(J-m) \quad (m = 0, \dots, J)$$

## Open linear

$$c(m) = \nu(J-m) + \frac{\phi(1-\lambda_0)}{J-1} \left( \frac{r}{1-e^{-r}} \right) m(J-m) \quad (m = 0, \dots, J)$$

# Which patch-occupancy model? $e(m)$

## Closed constant

$$e(m) = \phi N \frac{m(m-1)}{(N+m-1)(N+m-2)} \quad (m = 1, \dots, J, N \geq 2)$$

## Closed linear

$$e(m) = \phi N m \frac{b_{m-1}(N-1)}{b_m(N)} \quad (m = 1, \dots, J, N \geq 2)$$

$$b_m(N) = \sum_{k=0}^{m-1} (-1)^k \binom{m}{k} (m-k)^N \quad (m = 1, \dots, J) \quad b_0(N) = \delta_{N0}$$

## Open constant

$$e(m) = \phi(1-\rho)m \quad (m = 0, \dots, J)$$

## Open linear

$$e(m) = \phi \left( \frac{r e^{-r}}{1 - e^{-r}} \right) m \quad (m = 0, \dots, J)$$

# Which patch-occupancy model?

## Closed constant

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## Open

$$c(m) = \nu(J-m) + \frac{c}{J-1} m(J-m) \quad e(m) = em$$

**Constant**  $c = \phi(1 - \lambda_0)/(1 - \rho) \quad e = \phi(1 - \rho)$

**Linear**  $c = \phi(1 - \lambda_0)r/(1 - e^{-r}) \quad e = \phi r e^{-r}/(1 - e^{-r})$

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“Correct” logistic growth

Constant  $c = \phi(1 - \lambda_0)/(1 - \rho) \quad e = \phi(1 - \rho)$

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The SL model with immigration

Open

$$c(m) = \nu(J-m) + \frac{c}{J-1} m(J-m) \quad e(m) = em$$

**Constant**  $c = \phi(1 - \lambda_0)/(1 - \rho)$   $e = \phi(1 - \rho)$

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$$c(m) = \frac{N\phi}{J-1} (J-m) \quad e(m) = \phi N m \frac{b_{m-1}(N-1)}{b_m(N)}$$

## Open

$$c(m) = \nu(J-m) + \frac{c}{J-1} m(J-m) \quad e(m) = em$$

**Constant**  $c = \phi(1 - \lambda_0)/(1 - \rho) \quad e = \phi(1 - \rho)$

**Linear**  $c = \phi(1 - \lambda_0)r/(1 - e^{-r}) \quad e = \phi r e^{-r}/(1 - e^{-r})$

# Which patch-occupancy model?

For the open network with linear propagation rates (only), we can do much better.

We can evaluate the expected colonization rate and the expected local extinction rate as *time-dependent quantities*. This yields a corresponding *time-inhomogeneous* SL model:

$$c_t(m) = \nu(J - m) + \frac{c_t}{J - 1}m(J - m) \quad e_t(m) = e_t m.$$

Here  $c_t = \phi(1 - \lambda_0)r_t/(1 - e^{-r_t})$ ,  $e_t = \phi r_t e^{-r_t}/(1 - e^{-r_t})$ , where  $r_t = \nu(1 - e^{-\phi\lambda_0 t})/(\phi\lambda_0)$ .

# Local population dynamics

We have not attempted to account for local population dynamics (within patches).

Here is a simple embellishment that separates emigration from death:

$$q(\mathbf{n}, \mathbf{n} + \mathbf{e}_j) = \nu_j$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i) = d_i n_i + \phi_i(n_i) \lambda_{i0}$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i + \mathbf{e}_j) = \phi_i(n_i) \lambda_{ij}$$

# Local population dynamics

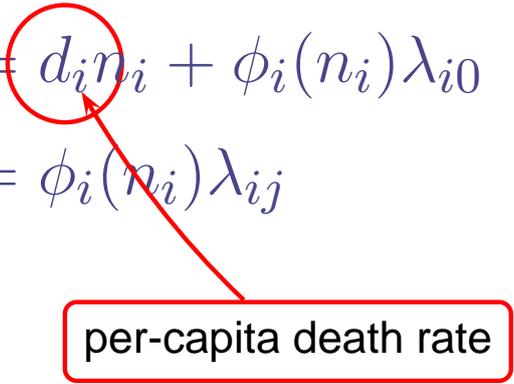
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per-capita death rate

# Local population dynamics

For example, with linear propagation rates ...

$$q(\mathbf{n}, \mathbf{n} + \mathbf{e}_j) = \nu_j$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i) = d_i n_i + \phi_i n_i \lambda_{i0} = \phi_i n_i \lambda'_{i0}$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i + \mathbf{e}_j) = \phi_i n_i \lambda_{ij}$$

where  $\lambda'_{i0} = \lambda_{i0} + d_i/\phi_i$ .

(This can be accommodated within the present setup with some minor adjustments.)

# Local population dynamics

And, something a little more complicated ...

Let  $S = \{0, \dots, N_1\} \times \dots \times \{0, \dots, N_k\}$  and define non-zero transition rates as

$$q(\mathbf{n}, \mathbf{n} + \mathbf{e}_i) = \nu_i + b_i \frac{n_i}{N_i} (N_i - n_i)$$

$$q(\mathbf{n}, \mathbf{n} - \mathbf{e}_i + \mathbf{e}_j) = \phi_i(n_i) \lambda_{ij}$$

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Local population dynamics are in accordance with the stochastic logistic model.