## Modelling invasions and calculating establishment success chances

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## Biological examples of invaders

- **❖** Exotic species
- **❖** Biocontrol agents
- ❖ Mutants
- **❖** Tumour cells
- ❖ Insecticide or pesticide resistance genes
- Artificially modified genes
- ❖ Pathogens

# What kind of models for invasion studies?



Fate is largely determined by chance, e.g.

- variation in offspring numbers
- hybridization and backcross chance (in introgresion)
- interaction with resident individuals

Stochastic processes

## Some classical population models

discrete time 
$$
x(n+1) = m \cdot x(n)
$$

continuous time 
$$
\frac{dx}{dt} = m \cdot x
$$

*m*: (mean) number of offspring per individual

- Deterministic
- *x*: density, continuous

## **Predictions**



# **Implications**

 $x(n+1) = m \cdot x(n)$ 

when *m* < 1: never success, always extinction when  $m \geq 1$ : always establishment, never extinction independent of initial population size (as long as  $x(0) > 0$ )

Typical result of deterministic models

# Small populations

individuals are discrete entities -> jumps in *x* inter-individual variation in offspring establishment chance depends on population size



## Some results

Populations may be successful or not if *m* > 1 Always die out if  $m \leq 1$ 

Examples: fate of 5 populations of 10 individuals (Geometric offspring)



# Basic model: branching process (Galton and Watson)

- Independent reproduction
- Nonoverlapping generations
- Identical offspring distributions



#### Popular offspring distributions



## Example: 20 runs



Poisson(1.1) distributed offspring numbers

## Example: 20 runs



## Calculation of extinction probability

Example "splitting process"



*Q*: probability of extinction if we start with 1 individual

## Roads to extinction



## Calculation of *Q*

$$
Q = p + (1-p)Q^2
$$
  $\longrightarrow$  solutions: 1 and  $\frac{p}{1-p}$ 

smallest root = extinction probability



## General calculation of *Q*



# Facts about Q

 $f(s) = \sum s^k \Pr \Big[ \xi = k \Big]$ *k*  $\sum s^k \Pr[\xi = k]$  probability generating function of offspring distribution

For 
$$
s \in [0,1]
$$
:  
\n $f(s) \ge 0$   
\nall derivatives of  $f(s) \ge 0$   
\n $f(0) = Pr[\xi=0]$   
\n $f(1) = 1$   
\n $f'(1) = m$ 

 $Q$  is smallest root of:  $Q = f(Q)$ 

 $m \leq 1 \rightarrow$  certain extinction

 $m > 1$  and Pr[0 offspring]  $> 0 \rightarrow 0 < Q < 1$ 

 $Pr[0 \text{ offspring}] = 0 \rightarrow Q = 0$ 



# Some terminology

Subcritical branching process: *m* < 1 Extinction certain, expected extinction time finite

Critical branching process: *m* = 1 Extinction certain, expected extinction time infinite

Supercritical branching process: *m* > 1 Positive establishment chance

#### Example: Poisson(*m*) offspring

$$
f(Q) = \sum_{k=0}^{\infty} e^{-m} \frac{m^k}{k!} Q^k = e^{-m} e^{mQ}
$$

$$
\implies Q = e^{-(1-Q)m}
$$

No explicit solution. Solve numerically or approximate.

# Approximation of *Q*

For slightly supercritical processes: *Q* close to 1

$$
f(Q) = f(1) + f'(1)(Q - 1) + \frac{1}{2}f''(1)(Q - 1)^2 + O((Q - 1)^3)
$$
  

$$
f(s) = E[s^x] \Rightarrow f(1) = 1, f'(1) = E[x], f''(1) = E[x(x - 1)]
$$
  

$$
f(Q) \approx 1 + m(Q - 1) + \frac{1}{2}E[x(x - 1)](Q - 1)^2
$$
  

$$
(1 - Q) \approx \frac{2(m - 1)}{E[x(x - 1)]} \approx \frac{2(m - 1)}{\text{Var}[x]}
$$

# Applications of the GWBP to biology

Main assumptions:

- 1. All reproductive individuals are equivalent, with identical offspring distributions
- 2. Individuals do not affect each other's reproduction
- 3. Offspring distributions do not change in time

At first sight not so realistic. However......

# 1. All *reproductive* individuals are equivalent

Clonal reproduction: unicellulars, e.g. bacteria, yeast

Hermaphrodites, e.g. monoecious plants

Two-sex species: only count females, provided: enough males available no genetic difference in reproduction, e.g. heterozygous mutants in a homozygous resident population





# Invasion of mutants: count only heterozygotes



mutant: heterozygote, mates with homozygous resident -> offspring are heterozygous too

#### Mutant invasion as a GWBP





# 2. Individuals do not affect each other's reproduction

Initial growth in environment with abundant resources.

Invasion in a large resident population that keeps the resource supply at a fixed level. Invaders do not mate and compete with each other, but only with residents.

# Example

Resident population large -> deterministic density-dependent model, e.g.

$$
x(n+1) = \frac{ax(n)}{1+bx(n)}
$$

*a* > 1: initial per-capita growth  $b > 0$ : intra-specific competition

Equilibrium: 
$$
\hat{x} = \frac{a-1}{b}
$$

Invader model GWBP with e.g.  $m =$ *a m*  $1 + c\hat{x}$ = *a m*  $1 + c(a-1)/b$ 

 $a_m$  > 0: per-capita growth of invader without competition  $c > 0$ : inter-specific competition

# 3. Offspring distributions do not change in time

Non-overlapping generations: reproduction only once in a lifetime.

Repeated reproduction, adults equivalent to juveniles, and constant mortality chance (no age-dependence), e.g. determined by predation risk.

# Overlapping generations as a GWBP



## From extinction to invasion probability

BP model: *Q* = extinction probability -> 1−*Q* = establishment success chance But: modelpopulations that do not go extinct grow infinitely large

Problems: (1) Realistically: populations have limited size (2) At large numbers invaders will affect each other

(1) Numerical analysis reveals: 1−*Q* is a good approximation for the chance to grow up to a large, fixed level.

(2) If 1−*Q* > 0 invasion is possible, but invaders might not take over a resident population completely (coexistence). This has to be examined separately.

#### Generalizations of the GWBP

Multitype processes Time-inhomogeneous processes Bisexual BP Population size -dependent





#### Mean matrix

 $m_{hj}$  = expected number of offspring of type *j* produced by 1 individual of type *h*

$$
M = \begin{pmatrix} m_{11} & m_{12} & \dots & m_{1d} \\ m_{21} & \cdot & \cdot & \cdot \\ \vdots & \cdot & \cdot & \vdots \\ m_{d1} & \cdot & \dots & m_{dd} \end{pmatrix}
$$

## Different kinds of multitype processes

Indecomposable: each type can *eventually* produce every other type, e.g.:

$$
M = \left(\begin{array}{cc} 0 & 3 \\ 1 & 2 \end{array}\right)
$$

Type 1 produces only type 2 offspring, but can have grandchildren of both types.

Decomposable: absorbing sets, e.g.

$$
M = \left(\begin{array}{cc} 2 & 1 \\ 0 & 1 \end{array}\right)
$$

Type 2 can only ever produce type 2

#### Periodic indecomposable processes

#### Example:

$$
M = \begin{pmatrix} 0 & 1 \\ 1 & 0 \end{pmatrix} = M^3 = M^5 = \dots = M^{2n+1}, n = 0, 1, 2, \dots
$$

$$
M^{2} = \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix} = M^{4} = M^{6} = \dots M^{2n}
$$

transformation to nonperiodic process: only consider process at even *n*, with mean matrix

$$
M'=M^2
$$

# Extinction of indecomposable multitype processes

ρ: largest eigenvalue of *M*

 $p$  < 1: Subcritical process: certain extinction in finite time  $p=1$ : Critical process: certain extinction, infinite expected time  $p > 1$ : Supercritical process: extinction probability < 1

Extinction probability depends on initial type. If no extinction occurs expected numbers of all types grow with rate ρ.

## Extinction of decomposable processes

Extinction probability depends on initial type, may be 1 for some types and less for others.

In processes that don't go extinct, some types may go extinct, while others grow, different types can grow at different rates, e.g.

$$
M = \left(\begin{array}{cc} 0.1 & 0 \\ 0 & 2 \end{array}\right)
$$

ρ=2, but if first individual has type 1, extinction is certain. If first individual has type 2, the process is supercritical, and non-extinct populations grow at rate 2.

# Calculation of extinction probabilities for multitype processes

 $Q_h$  = Pr extinction if initial individual has type *h* 

pgf of offspring distribution of type *h*:

$$
f_h(s_1, \ldots s_d) = \mathbf{E}\bigg[s_1^{\xi_{h1}}s_2^{\xi_{h2}}\ldots s_d^{\xi_{hd}}\bigg]
$$

ξ*hj* = number of type *j* children produced by a parent of type *h*, then

$$
Q_h = f_h(Q_1, \ldots, Q_d)
$$

## Proof

$$
\Pr\left[\text{extinct if initial type is } h\right]
$$
\n
$$
= \sum_{x_1} \dots \sum_{x_d} \Pr\left[\xi_{h1} = x_1, \dots, \xi_{hd} = x_d\right] Q_1^{x_1} Q_2^{x_2} \dots Q_d^{x_d}
$$
\n
$$
= \mathbb{E}\left[Q_1^{\xi_{h1}} Q_2^{\xi_{h2}} \dots Q_d^{\xi_{hd}}\right]
$$

## Example: spore formation



# Generalizations of the GWBP: Changing environments

Smith (1968), Smith & Wilkinson (1969): Inhomogeneous BP



Expected # offspring:  $m_t$   $m_{t+1}$ 

#### Extinction of inhomogeneous processes

 $E[log m_t] \leq 0$  Certain extinction:  $Q = 1$ 

 $\text{E}[\text{log}m_t]$  >  $0$   $\;$  Extinction probability  $Q$  is a random variable with E[*Q*]<1

## Q is a random variable: example



## Extinction depends on invasion time



# Numerical calculation of *Q*

 $Q_t$ : Pr[1 invader at  $t$  fails],  $f_{t}(s)$ : pgf of offspring distribution at  $t$ 

$$
Q_t = \sum_k \Pr[\text{invader at } t \text{ has } k \text{ offspring}] Q_{t+1}^k
$$
  
=  $f_t(Q_{t+1})$ 

Backward iteration (i.i.d.  $m_t$  values): Start with array of (arbitrary) *Q*-values in (0,1) Simulate random *m*-values Calculate *Q*-values 1 timestep *before* Continue until distribution is stable



## Invasion mode and extinction risk

Simultaneous

\n
$$
Q_{sim} = E\left[Q_t^n\right] = E\left[Q^n\right]
$$
\nSequential

\n
$$
Q_{seq} = E\left[\prod_{t=1}^n Q_t\right]
$$
\nIndependent sites

\n
$$
Q_{ind} = \left(E\left[Q_t\right]\right)^n = \left(E\left[Q\right]\right)^n
$$

Jensen's inequality: 
$$
(E[Q])^n \le E[Q^n]
$$
  
\n $Q_{ind} \le Q_{sim}$   
\nHölder's inequality:  $E\left[\prod_{t=1}^n Q_t\right] \le \left(\prod_{t=1}^n E[Q_t^n]\right)^{\frac{1}{n}} = E[Q^n] Q_{seq} \le Q_{sim}$ 

Haccou & Vatutin (TPB, 2003):  $Q_{ind} \leq Q_{seq}$  if  $m_t$  are independent

## Numerical results



 $m_{_t}$  i.i.d. uniform,  ${\rm E}[m_{_t}]=1.3, {\rm Var}[m_{_t}]=0.5,$  Poisson distr. offspring