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 $m \ge 1$: 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 > 1Always die out if $m \le 1$

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

 $f(s) = \sum_{k} s^{k} \Pr[\xi = k]$ probability generating function of offspring distribution

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

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

 $m \le 1 \rightarrow certain extinction$

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

Pr[0 offspring] = 0 $\rightarrow Q = 0$



Some terminology

Subcritical branching process: m < 1Extinction 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

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

No explicit solution. Solve numerically or approximate.

Approximation of Q

For slightly supercritical processes: Q close to 1

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

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

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

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

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. $m = \frac{a_m}{1 + c\hat{x}} = \frac{a_m}{1 + c(a-1)/b}$

 $a_m > 0$: 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*

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

Different kinds of multitype processes

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

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

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

Decomposable: absorbing sets, e.g.

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

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

 ρ < 1: Subcritical process: certain extinction in finite time ρ = 1: Critical process: certain extinction, infinite expected time ρ > 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

 $Q_h = \Pr\left[\text{extinction if initial individual has type } h\right]$

pgf of offspring distribution of type *h*:

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

 ξ_{hj} = number of type *j* children produced by a parent of type *h*, then

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

Proof

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

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

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

 $E[\log m_t] > 0$ 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*

$$Q_{t} = \sum_{k} \Pr\left[\text{invader at } t \text{ has } k \text{ offspring}\right] Q_{t+1}^{k}$$
$$= f_{t} \left(Q_{t+1} \right)$$

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
$$Q_{sim} = E[Q_t^n] = E[Q^n]$$
Sequential $Q_{seq} = E\left[\prod_{t=1}^n Q_t\right]$ Independent sites $Q_{ind} = (E[Q_t])^n = (E[Q])^n$

Jensen's inequality:
$$(E[Q])^n \le E[Q^n]$$

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

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

Numerical results



 m_t i.i.d. uniform, $E[m_t] = 1.3$, $Var[m_t] = 0.5$, Poisson distr. offspring