

# Stochastic models in ecology and evolution

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## Introduction

- **huge** topic (Ulam's analogy: the study of nonlinear systems as the study of "non-elephant animals". Ditto for deterministic systems.)
- stochasticity generally not covered thoroughly in math biology classes (but see Allen (2003), Kot (2001)); harder math, (?) less clear-cut results (?)
- stochasticity in a model: *epistemological* rather than an *ontological* purpose
- analysis harder . . . especially for nonlinear systems (most of the interesting ones, although there are interesting linear approximations, e.g. during invasions)
- definitions can be a bit tricky: is a diffusion model a "stochastic" model or not? Not explicitly . . . but implicitly based on a "random walk" or Brownian motion process (movement of individuals in space, or of populations in state space)
- well-posed stochastic models typically converge to deterministic ones in *some* limit (e.g. Kurtz (1970)), either in a large-population case or as the expected solution of an ensemble of realizations
- defs: *endogenous* ( $\approx$  demographic) vs. *exogenous* ( $\approx$  environmental) stochasticity
- state space can get very large (e.g. metapopulation models)

## Examples

- discrete-state populations (Markov chains, etc.) easier than continuous populations (stochastic differential equations)
- discrete time easier than continuous time

Simple population models: analogues of logistic equation. Mortality = binomial  $(\mu + \alpha N)$ . Fecundity = Poisson  $(fN)$ . (Different from SI model, which has infection binomial with probability  $(1 - \exp(-\beta I \Delta t))$  — maximum value = 1.)

(Discrete models have their own issues: results may be sensitive to the order of events — fecundity before mortality, or vice versa? Problems go away as  $\Delta t \rightarrow 0$ .)

Continuous-time model: use *master equation*

|       |                       |                             |
|-------|-----------------------|-----------------------------|
| birth | $N \rightarrow N + 1$ | $fN \Delta t$               |
| death | $N \rightarrow N - 1$ | $(\mu + \alpha N) \Delta t$ |

(note  $r = f - \mu$ ,  $K = (f - \mu)/\alpha$ ;  $R = f/\mu$  disappears from deterministic analogue but not in stochastic case)

## Analytical techniques

- branching processes
- moment generating functions
- Kolmogorov (Fokker-Planck) "forward equation", diffusion approximations
- (quasi-)stationary distributions (Keeling and Ross, 2008)
- moment equations/moment closure (e.g. Isham (2005))

## Numerical techniques

- Gillespie model
- exact solutions of Kolmogorov forward equations (leading eigenvectors)
- moment equations

## Phenomena

Are stochastic models just “deterministic models with added noise”?

- Jensen’s inequality (static, not dynamic, but important): e.g. geometric mean in exponential growth
- extinction and fixation: absorbing boundaries in population genetics and ecology (first passage times etc.)
- neutral theories of population genetics (Ewens) and ecology (Hubbell): related to classical “urn problems”
- interaction of nonlinearity with noise: fractal basin-hopping, (de)stabilization of attractors by noise (stochastic repellors) (Coulson et al., 2004) Rand & Keeling

## See also

Some talks by Rick Durrett (applied probability, Cornell): <http://www.math.cornell.edu/~durrett/Talks/Talks.html>, and the corresponding paper <http://www.math.cornell.edu/~durrett/Talks/waldpaper.pdf>

Lecture notes: <http://www.amath.washington.edu/courses/423-winter-2007/outline.pdf>

Bailey (1990); Renshaw (1993); Lande, Engen, and Sther (2003)

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