Diffusion Approximations for Ecological Models

by

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Zen Maxims for Survival in a Modern University

Those not relevant to my talk

• Don’t be irreplaceable; if you can’t be replaced, you can’t be promoted.

• Give a man a fish and he will eat for a day; teach him how to fish, and he will sit in a boat and drink beer all day.

• If you lend someone $20, and never see them again, it was probably worth it.

• If you tell the truth, you don’t need a good memory.

• A closed mouth gathers no foot.
Zen Maxims for Survival in a Modern University

Those relevant to my talk

• Good judgement comes from bad experience, and a lot of that comes from bad judgement.

• It may be that your sole purpose in life is simply to serve as a warning to others.

• Before you criticize someone, you should walk a mile in their shoes; that way, when you do criticize them, you’ll be a mile away and you’ll have their shoes.
**DIFFUSION APPROXIMATIONS**

**Technical definition.** A *diffusion process* is a continuous-time (strong) Markov process whose sample path $X(t)$ is an almost-surely continuous function of $t$.

Diffusion processes are usually realised as approximations of jump processes (in discrete or continuous time) by way of a limiting procedure involving space-time scaling: $\Delta x \sim \sqrt{\Delta t}$. 

![Sample path of a diffusion model for population density](image-url)
Why are they useful models? They are popular in ecological modelling because their parameters can be estimated simply from very little data, and they offer explicit expressions for important quantities of interest, such as the expected time to extinction.

Difficulties. Formulae arising from diffusion models are frequently used in situations where this is clearly inappropriate:

- The underlying process, which the diffusion is approximating, might not itself be appropriate for modelling the system in question.
- The approximation procedure is often not well understood, and its physical meaning not taken into account.
- The diffusion approximation might provide estimates that are at variance with the underlying model.
**EXAMPLE**

**The stochastic logistic model.** This model is ubiquitous in the literature on population modelling, but also appears in a variety of different contexts: for example, chemical kinetics, genetics and epidemics. It is a birth-death process $n(t)$ taking values in $S = \{0, 1, \ldots, N\}$ with birth and death rates

$$
q(n, n+1) = \frac{\lambda}{N} n(N-n), \\
q(n, n-1) = \mu n,
$$

where $N$ is to be interpreted as the carrying capacity and $\lambda, \mu > 0$. Absorption at state 0 (representing the event of extinction) occurs with probability 1. We will consider the interesting case of positive drift: $\lambda > \mu$.

**A diffusion approximation.** There is a diffusion approximation for the *density* $X(t) = n(t)/N$, which becomes more and more accurate as $N$ gets large.
Illustration. Here is a simulation of the logistic model, together with its diffusion approximation \((N = 500, \lambda = 0.2, \mu = 0.1\) and \(X(0) = x_0 = 0.1\)):

![Graph showing population density over time](image)

The mean path is shown (solid), with \pm two standard deviations (dashed).
**Specification of the approximation.** \( X(t) \) has an approximate normal \( N(\mu_t, \sigma_t^2/N) \) distribution, where

\[
\mu_t = \frac{qx_0}{x_0 + (q - x_0)e^{-\lambda qt}}, \quad t \geq 0,
\]

\( q = 1 - \rho, \ \rho = \mu/\lambda \ (< 1), \) and

\[
\sigma_t^2 = M_t^2 \int_0^t \left( G(\mu_s)/M_s^2 \right) ds,
\]

where \( G(x) = \lambda x(1 - x) + \mu x, \) \( 0 \leq x \leq 1, \) and

\[
M_t = \frac{q^2 e^{-\lambda qt}}{(x_0 + (q - x_0)e^{-\lambda qt})^2}.
\]

Note that \( \sigma_t^2 \to \rho \) and \( t \to \infty. \)

Note also that \( q \) is an equilibrium point, which is asymptotically stable \( (\mu_t \to q). \) It can be shown that

\[
\sigma_t^2 = \rho(1 - e^{-2\lambda qt}) + O(|x_0 - q|),
\]

for \( x_0 \) near \( q. \)
**Extinction times.** The diffusion approximation accurately models the density $X(t)$, but how good is it in predicting extinction times? Let $\tau_i$ be the expected time to extinction starting in state $i$.

![Graph showing expected time to extinction](image)

The approximation (dashed) for $\tau_i$ is shown, together with the exact values (solid) obtained directly from the logistic model ($N = 100$, $\lambda = 0.1111$, $\mu = 0.1$, $n(0) = n_0 = i$, $x_0 = x = i/N$).
Shown is the diffusion approximation (dashed) for $\tau_i$, the exact values (solid) and a large-$N$ asymptotic expansion (dash-dot):

$$\tau_i \sim \frac{\rho(1 - \rho^i)}{\mu(1 - \rho)^2} \left( \frac{e^{-(1-\rho)}}{\rho} \right)^N \sqrt{\frac{2\pi}{N}}.$$  

($\rho = 0.1538$, $\mu = 0.1$, $x_0 = 0.05$, and $n_0 = i = [0.05 \times N].$)
RECENT WORK*

Summary. A class of Markovian models is specified, called *asymptotically density dependent* models, which permit a diffusion approximation via a simple limiting procedure. This procedure allows one to identify the most appropriate approximating diffusion and to decide whether the diffusion approximation is appropriate for describing the population in question. Result are presented in a form that most easily permits their direct application to population models. Additionally, results are presented which can be used to assess the accuracy of diffusion approximations: they specify for *how long* and *over what ranges* the underlying Markovian model is faithfully approximated.

THE LURE OF EXPLICIT FORMULAE

Foley* proposes the following formula for the expected extinction time in terms of

\[ L_t = -\log(R_t), \]

where \( R_t \) is the growth rate at time \( t \):

\[ \tau_i \sim \frac{1}{2sr} \left( N^{2s} \left( 1 - i^{-2s} \right) - 2s \log(i) \right), \]

where \( i = n_0 \), \( s = r/v \), and \( r \) and \( v \) are the expected value and the variance of \( L_t \) (both assumed to be constant).

This formula is nothing more than an expression for the expected time it takes Brownian motion with drift \( r \) (reflected at \( \log(N) \)) to reach state 0 from a given state \( i \), and thus it would be naive to expect Foley’s formula to be universally applicable.

*P. Foley, Predicting extinction times from environmental stochasticity and carrying capacity, Conservation Biology, 8, 124–137, 1994.
Whether this formula could possibly predict extinction times accurately depends firstly on whether the population in question could be modelled as a discrete-state process that jumps \textit{up or down by 1} at each time point (\textit{independently} at each time point) and with jump probabilities that \textit{do not depend} on the population size, and, secondly, whether there have been sufficiently many transitions for the approximation procedure to work (this procedure mirrors the way in which the Central Limit Theorem works, with the position of the jump process at any given time instant being expressed as the sum of a large number of independent increments).

For example, the logistic model does not satisfy all of these conditions; most importantly, the jump probabilities are state dependent.

And, you will not be surprised to learn that . . .
Shown is the diffusion approximation (dashed) for $\tau_i$, the exact values (solid) and those predicted using Foley’s formula (dash-dot) ($N = 100, \lambda = 0.1111, \mu = 0.1, n_0 = i, x_0 = x = i/N$).