

Mathematical Techniques in Evolution and Ecology

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Introduction

To come.

Chapter 1

How to construct a model

1.1 Problems

- P.1.1 In the squirrel example, assume that death through cyclists occurs before immigration, and immigration is followed by birth. Derive the discrete-time recursion equation and compare it to the one we obtained above, where we assumed that death is followed by birth and immigration. Start from the life-cycle diagram. Interpret the difference.
- P.1.2 [Problem 2.5 in OD2007] Consider a model of disease transmission with the following equations:

$$\frac{dS}{dt} = \theta - dD - \beta SI + \gamma I, \quad (1.1a)$$

$$\frac{dI}{dt} = \beta SI - (d + \nu + \gamma)I. \quad (1.1b)$$

The variables S and I denote the number of susceptible and infected individuals. (a) Draw and label a flow diagram for these two variables. (b) Suggest a plausible biological interpretation of the parameters γ and ν .

- P.1.3 [Problem 2.6 in OD2007] In the flu model, suppose that after contracting the flu, people are initially resistant to reinfection, but this immunity eventually wanes. Alter the flow diagram for the flu model that we derived to include a “recovered

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and immune” class with these properties. (b) Suppose that immune individuals have a constant per capita rate of losing immunity. What are the continuous-time equations for this modified model?

Chapter 2

Some classic models in evolution and ecology

2.1 Problems

- P.2.1 We constructed the differential equation for the continuous-time version of the exponential growth model from the flow diagram. (a) Instead, derive the equation from the difference equation of the discrete-time version. Hint: Introduce a small amount of time Δt and take the appropriate limit. (b) Comparing r_d and r_c , explain the difference between the discrete- and continuous-time versions.
- P.2.2 Derive the differential equation for the logistic growth model in continuous time. Start by assuming that the per capita growth rate $r(n)$ declines linearly with population size $n(t)$. Assume that $r(n)$ has a maximum equal to the intrinsic growth rate r_c when there are no competitors (i.e. $r(0) = r_c$), and that it reaches zero when the population is at the carrying capacity (i.e. $r(K) = 0$). Substitute $r(n)$ for r_c in the differential equation of the exponential growth model to obtain the result of interest.
- P.2.3 As an alternative way of deriving the continuous-time version of the haploid selection model, start from the continuous-time version of the exponential growth model. Specifically, let n_A

and n_a be the number of individuals carrying alleles A and a , respectively, and let the growth rate depend on the allelic state, i.e. $r_A = b_A - d_A$ for carriers of A , and $r_a = b_a - d_a$ for carriers of a . From the exponential growth model, we have

$$\frac{dn_A}{dt} = r_A n_A(t), \quad (2.1a)$$

$$\frac{dn_a}{dt} = r_a n_a(t). \quad (2.1b)$$

However, we want to know how the allele frequencies change! As these sum to 1, it is sufficient to follow only the frequency of allele A , $p(t) = n_A(t)/(n_A(t) + n_a(t))$. Start by writing

$$\frac{dp}{dt} = \frac{d \left[\frac{n_A(t)}{n_A(t) + n_a(t)} \right]}{dt}. \quad (2.2)$$

Then use the quotient rule to evaluate the derivative, and use the equations for dn_A/dt and dn_a/dt to express dp/dt in terms of r_A , r_a , and p .

Chapter 3

Primer 1: Functions and approximations

3.1 Problems

P.3.1 Determine the functions $R(n)$ for the reproductive factor in the logistic growth model, such that the shape of the functions is consistent with the first figure above. In each case, choose the parameters such that the intercept is $R(0) = 1 + r$, and the value when $N = K$ is $R(K) = 1$. Specifically, derive

- (a) A function that declines exponentially to zero.
- (b) A quadratic function with a maximum at $n = 0$.
- (c) A reverse-S-shaped function that declines from $1 + r$ to zero. Keep a arbitrary, and use Rule P1.1 above to increase the intercept to $1 + r$. Hint: Start by stretching the generic formula for the S-shaped function along the y axis.

P.3.2 Use Recipe P1.1 to confirm the following linear approximations:

- (a) $e^r \approx 1 + r$, assuming r is small,
- (b) $1/(1 + s) \approx 1 - s$, assuming s is small,
- (c) $\ln(t) \approx t - 1$, assuming t is near 1, and

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(d) $1/x \approx 1/a - 1/(a^2)(x - a)$, assuming x is near a .

Chapter 4

Solutions to problems

4.1 Chapter 1

P.1.1 The life-cycle diagram is as given in Fig. 4.1. The recursion equation is

$$n(t+1) = (1+b) [(1-d)n(t) + m], \quad (4.1)$$

as compared to

$$n(t+1) = (1+b)(1-d)n(t) + m \quad (4.2)$$

for the case of immigration after death and birth. The first equation differs from the second one by an amount of bm , which is the fraction of squirrels born by immigrating (female) squirrels.

P.1.2 The flow diagram is shown in Fig. 4.2. The parameter γ is the rate at which infected individuals recover and become susceptible again. The parameter ν denotes an additional death rate experienced by infected individuals as compared to the death rate d of susceptible ones.

P.1.3 The flow diagram is as given in Fig. 4.3, and the continuous-

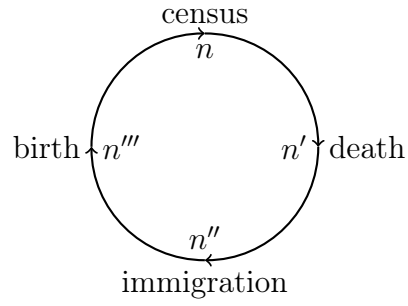


Figure 4.1 Life-cycle diagram for a squirrel model with alternative order of events.

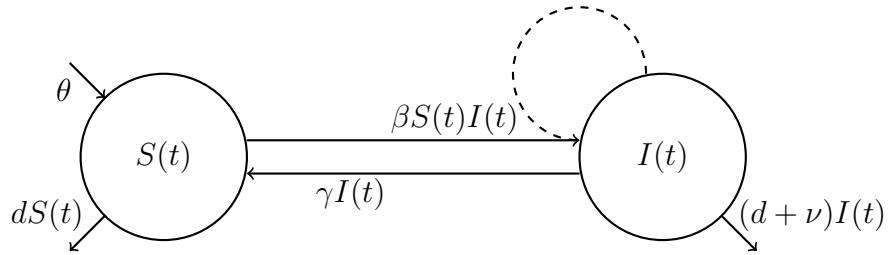


Figure 4.2 Flow diagram for the continuous-time model of Problem P.1.2.

time differential equations are

$$\frac{ds}{dt} = \sigma r(t) - acs(t)n(t) \quad (4.3a)$$

$$\frac{dn}{dt} = acs(t)n(t) - \rho n(t) \quad (4.3b)$$

$$\frac{dr}{dt} = \rho n(t) - \sigma r(t) \quad (4.3c)$$

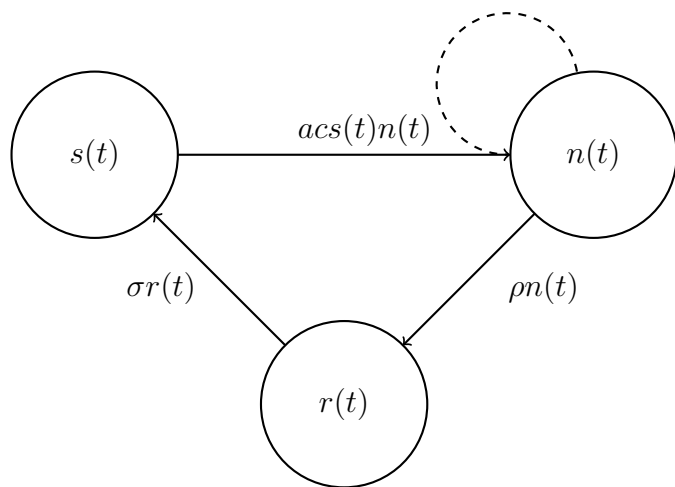


Figure 4.3 Flow diagram for the continuous-time version of the flu model of Problem P.1.3, with an additional class of resistant individuals.

4.2 Chapter 2

P.2.1 We start from the discrete-time difference equation for the exponential growth model,

$$\Delta n = n(t+1) - n(t) = Rn(t) - n(t) = (R-1)n(t) \quad (4.4)$$

We recall that the reproductive factor was given by $R = (1+b)(1-d)$, where b and d are the number of offspring per adult female and the proportion of individuals dying per generation, respectively. Moreover, we recall that the growth rate was defined as $r_d = R - 1 = b - d - bd$. Therefore, the difference equation can be written as

$$\Delta n = (R-1)n(t) = r_d n(t) = (b - d - bd)n(t). \quad (4.5)$$

To derive the differential equation, consider a small amount of time Δt , such that the number of offspring per adult female and the proportion of individuals dying during this time are now $b\Delta t$ and $d\Delta t$, respectively. Hence, $n(t + \Delta t) - n(t) = [b\Delta t - d\Delta t - bd(\Delta t)^2] n(t)$. Applying the definition of the

differential, we find

$$\begin{aligned}
 \frac{dn}{dt} &= \lim_{\Delta t \rightarrow 0} \frac{n(t + \Delta t) - n(t)}{\Delta t} \\
 &= \lim_{\Delta t \rightarrow 0} \frac{[b\Delta t - d\Delta t - bd(\Delta t)^2] n(t)}{\Delta t} \\
 &= \lim_{\Delta t \rightarrow 0} (b - d - bd\Delta t)n(t) \\
 &= (b - d)n(t).
 \end{aligned} \tag{4.6}$$

This implies that the continuous-time growth rate is $r_c = b - d$. Comparing to $r_d = b - d - bd$, we see that the interaction term bd is lost when we switch from discrete to continuous time. This is because, in the limit of a time step becoming small, no more than one type of event occurs. In continuous time, individuals that are born in a small amount of time will not suffer death in that same amount of time.

P.2.2 We start by recalling the continuous-time differential equation for the exponential growth model,

$$\frac{dn}{dt} = r_c n(t). \tag{4.7}$$

To account for the effect of intra-specific competition for limited resources, we want r_c to depend on population size as described in the problem. For this, the intercept needs to be equal to r_c and the slope equal to $-r_c/K$. Hence,

$$r_c(n) = r_c + \left(-\frac{r_c}{K}\right) n(t) = r_c \left(1 - \frac{n(t)}{K}\right). \tag{4.8}$$

Substituting $r_c(n)$ for r_c in Eq. (4.7), we obtain the solution

$$\frac{dn}{dt} = r_c \left(1 - \frac{n(t)}{K}\right) n(t). \tag{4.9}$$

P2.3 Applying the quotient rule of differentiation, we find

$$\begin{aligned}
 \frac{dp}{dt} &= \frac{\frac{dn_A}{dt} (n_A + n_a) - n_A \left(\frac{dn_A}{dt} + \frac{dn_a}{dt}\right)}{(n_A + n_a)^2} \\
 &= \frac{n_a \frac{dn_A}{dt} - n_A \frac{dn_a}{dt}}{(n_A + n_a)^2}.
 \end{aligned} \tag{4.10}$$

Substitution of $dn_A/dt = r_A n_A(t)$ and $dn_a/dt = r_a n_a(t)$ yields

$$\begin{aligned}\frac{dp}{dt} &= \frac{n_a(r_A n_A) - n_A(r_a n_a)}{(n_A + n_a)^2} \\ &= (r_A - r_a) \frac{n_A n_a}{(n_A + n_a)^2} = (r_A - r_a) p(t) q(t),\end{aligned}\tag{4.11}$$

using $q(t) = 1 - p(t) = n_a(t)/(n_A(t) + n_a(t))$. Note that in the derivation we have omitted the dependence of n_A and n_a on t in our notation for simplicity. Comparing the result to the differential equation derived earlier, $dp/dt = s_c p(t) q(t)$, we see that the continuous-time selection coefficient s_c can be interpreted as the difference between the intrinsic growth rates of the two haploid genotypes that compete.