

PopBio Help

This document briefly describes the equations used by each of the simulations in the PopBio program. A more detailed application manual is planned for a future release. In the mean time, if you have a question about the PopBio program that is not answered here please contact Ethan Benatan at benatan@reed.edu.

Population growth

Variables

- x = [population]
- x_0 = [initial population]
- r = [rate of increase]
- K = [carrying capacity]
- L = [growth time lag]

Equations

Exponential growth

The basic assumption of population growth modeling is that the rate of population growth is proportional to the current population – that is,

$$(1) \quad \frac{dx}{dt} = rx$$

The solution to (1) is

$$(2) \quad x(t) = x_0 e^{rt}$$

Equation (2) is used for computing $x(t)$ deterministically; the stochastic mode uses equation (1), recalculating the value of r at each time step.

Logistic growth

To incorporate the carrying capacity, we add a term to dx/dt :

$$(3) \quad \frac{dx}{dt} = rx \left(1 - \frac{x}{K}\right)$$

The solution to this equation is

$$(4) \quad x(t) = \frac{Kx_0 e^{rt}}{K + x_0 (e^{rt} - 1)}$$

As with exponential growth, the deterministic mode uses the second equation, while the stochastic mode uses the first, recalculating K and r at every time step. If the growth lag in a stochastic calculation is non-zero, then for $t \geq L$, $x(t - L)$ is used in place of x in equation (3).

Predator-Prey

Variables

- N = [prey population]
- P = [predator population]
- a = [prey birth rate]
- b = [prey capture probability]
- c = [predator energy efficiency] $\cdot b$
- d = [predator death rate]
- K = [prey carrying capacity]
- e = [prey critical density]
- m = [maximum predator attack rate]
- W = [1/2 attack rate population]
- w = [1/2 attack rate density]

Lotka-Volterra model

$$dN/dt = aN - bNP$$

$$dP/dt = cNP - dP$$

Leslie model

$$dN/dt = aN(1 - N/K) - bNP$$

$$dP/dt = cP(1 - eP/N)$$

Holling models

$$dN/dt = aN(1 - N/K) - b(N, P)P$$

$$dP/dt = c b(N, P)P - dP$$

$$b(N, P) = mN/(W + N) \quad \text{(Holling I functional response)}$$

$$b(N, P) = mN/(wP + N) \quad \text{(Holling II functional response)}$$

Population Genetics

Variables

- p = [A allele frequency]
- S = [Population size]
- w_{11} = [Genotype AA fitness]
- w_{12} = [Genotype Aa fitness]
- w_{22} = [Genotype aa fitness]

Equations

Given p_b (with $q_n \equiv 1 - p_n$), compute p_{n+1} using the following equation:

$$p_{n+1} = p_n q_n \frac{(w_{11} - w_{12})p_n + (w_{12} - w_{22})q_n}{w_{11} p_n^2 + 2w_{12} p_n q_n + w_{22} q_n^2} + r_n \sqrt{\frac{3p_n q_n}{2N}}$$

If genetic drift is enabled, then for each n , r_n is a random number in the range $(-1, 1)$; otherwise, $r_n = 0$ for all n .

Life Table

Variables

- N = [number of age groups]
- x = [age group index between zero and $N - 1$ inclusive]
- n_x = [initial population of age group x]
- P_x = [probability of an individual surviving to age group x]
- m_x = [fecundity of age group x]

Equations

The columns of the Life Table output matrix are computed using the following equations.

$$\begin{aligned}p_x &= P_{x+1}/P_x \\q_x &= 1 - \frac{p_{x+1}}{p_x} \\d_x &= P_x - P_{x+1} \\e_x &= P_x^{-1} \sum_{j=x+1}^{j < N} P_j\end{aligned}$$

The Age Structure animation is computed in the following way. At every time interval the population in age group x advances into age group $x + 1$ and size of the advancing population is given by

$$n_{x+1,t+1} = n_{x,t}p_{x,t}$$

After each age group has been advanced, the first age group is populated according to the equation

$$n_{0,t+1} = \sum_{j=0}^{j < N} n_{x,t}m_x$$

The survivorship curve visualizes the probability that individuals will survive into each age group and the total population curve shows $\sum n_x$ over time.

Competition

Variables

- x = [species population]
- r = [rate of increase]
- K = [carrying capacity]
- α_{12} = [species 1 competition coefficient]
- α_{21} = [species 2 competition coefficient]

Equations

$$\begin{aligned}\frac{dx_1}{dt} &= r_1 x_1 \left(1 - \frac{x_1 + \alpha_{12} x_2}{K_1} \right) \\ \frac{dx_2}{dt} &= r_2 x_2 \left(1 - \frac{x_2 + \alpha_{21} x_1}{K_2} \right)\end{aligned}$$