

EEB 2208 TOPIC 15

POPULATION VIABILITY ANALYSIS

Reading for this topic

Primack: Chapters 13

Sodhi and Ehrlich: Chapter 16, pp 328-330

1. Population viability analysis (PVA)

A) THE BASICS OF DEMOGRAPHIC MODELING

- i) Population viability analysis is a special case application of the commonly-used population modeling that is frequently done by ecologists (you might have encountered this in general ecology or in a course on wildlife management). Human demographers use similar methods to determine the rate of change of human population sizes.
- ii) In its simplest terms, all that needs to be done is to figure out how big a population is now, and to predict how its size will change. To do this we need to know how many individuals will be added and how many will be lost at each time step.
- iii) To figure out how many to add, we need to estimate the number of breeders in the population and how many young each of those breeders will produce.
- iv) To determine how many are lost, we need to know the mortality rate (i.e., how many will die), preferably with some sort of breakdown for different age classes (e.g., juveniles usually have much higher death rates than adults).
- v) Then, just to complicate things, we also have to account for whether there will be immigrants (which get added) or emigrants (which get subtracted). If an entire population is being modeled (as is often the case with endangered species), immigration/emigration can be ignored.
- vi) To put this in simple mathematical terms:

$$N_{t+1} = N_t + \text{Births} - \text{Deaths} + \text{Immigrants} - \text{Emigrants}.$$

Where, N_t is the current population size; N_{t+1} is the population size one time-step in the future (usually time-steps are measured in years, but they do not have to be).

- vii) In very simplistic terms, if $(B+I)$ is larger than or equal to $(D+E)$ then the population will be OK. If $(D+E)$ is larger than $(B+I)$, the population will decline, and eventually go extinct.

B) INCORPORATING VARIATION

- i) A key element of any PVA is that it should provide information about the potential variability in the system. When we estimate the numbers that go into a model, we know that there is uncertainty about each of them. This uncertainty arises both because we often do not have very good information (i.e., our estimates have error), but also because of **stochasticity** inherent in the system. For example, not every individual produces exactly the same number of young, not every individual gets to breed, not every year is as good as the others, and so on.
- ii) This stochasticity is incorporated by telling the computer not to use a single fixed number for each part of the equation, but rather to pick a number from a range of possible numbers. Some numbers may be more

likely to be picked than others (can you think where these numbers might lie in a distribution?), but it is not certain how any particular simulation will turn out. Models that incorporate variability in this way are referred to as **stochastic**, rather than **deterministic** (a deterministic model will produce exactly the same result every time – i.e., the result is “determined” by the model’s design and input data).

- iii) **Genetic stochasticity** will be covered in more detail in the conservation genetics lecture. Briefly, drift is a random process by which genetic variation is lost from a population. As populations get smaller, variation is lost at a faster rate. Mutation is another stochastic process that restores variation, but the mutation rate does not vary with population size. So, the smaller the population size, the more the losses due to drift will outweigh the gains through mutation.
- iv) **Demographic stochasticity** refers to random variation in birth and death rates. For example, imagine you have a population in which there is a high chance of an individual failing to breed in any one year. If the population is large, then there will almost certainly be some breeding every year (just as if you roll a pair of dice 1000 times then you will almost certainly get two sixes at least once). But if population size is sufficiently small then it is possible that no individuals will breed in some years – simply due to chance (i.e., if you only roll a pair of dice ten times, then you could easily not get two sixes).
- v) **Environmental stochasticity** refers to the effects of random variation in environmental conditions. For example, there might be a string of “bad” years for a species. If the population is big, this might not be a big deal – there will be a decline in population size, but nothing that the population cannot recover from once conditions improve. But, if the population is small, a string of bad years might be just enough to cause it to decline to zero – from which there is no recovery.
- vi) Finally, many models incorporate an additional type of variation: **catastrophes**. These are major events that have a drastic impact on a population (e.g., they cause total reproductive failure, or they kill a large proportion of the individuals). Catastrophes are really just the extremes that arise from environmental variation and could be included within environmental stochasticity. But, for several reasons it is often easier to incorporate them into models separately (if you want to know the details, ask me about it – basically, it is just that the math is harder if you don’t do it this way).

C) ESTIMATING EXTINCTION RISK

- i) Once the model has been created, all the researcher needs to do is run repeated simulations and determine how many simulations end with the population going extinct.
- ii) To determine the probability that a population of a given size will go extinct within a given number of years, one simply needs to set the starting population size and then determine the proportion of the model runs that went extinct within that time frame. E.g., if 1000 separate simulations are each run for 50 years, and the computer population went extinct in 423 of the simulations, then the probability of extinction is predicted to be 0.423 (i.e., just over 40%) over a 50 year period.
- iii) To determine how extinction risk changes with population size, one can simply repeat this process with different starting population sizes, and compare the number of extinctions at each size.
- iv) Similarly, to determine how effective different management actions are likely to be, you can simply change the numbers in the model (the parameter estimates) based on how they are expected to change with the

target management action, and compare the extinction probabilities under each scenario.

- v) We will do a demonstration of these simulations in class – actually seeing how it works makes it much easier to understand than just reading about it.

D) CASE STUDY: HAWAIIAN STILT

- i) Once a model has been created, we first need to ask if we can trust the model results. One way to do this is to see if the model accurately predicts what has happened in the past. Another alternative is to see whether plausible (i.e., accounting for error in the way the estimates were made) changes in the numbers that go into the model radically change the results.
- ii) For example, in a PVA on Hawaiian stilts (described in the Primack text book), the model predicted that the population would increase over time until all of the habitat was filled. This prediction matched reality – populations have steadily increased on all islands where stilts are found and have subsequently started to level off, suggesting habitat saturation.
- iii) When the model was constructed, some of the numbers had to be made up (note that “made up” does not necessarily mean that the numbers were randomly picked from thin air – it is often possible to make well-informed guesses).
- iv) For example, there was no information on survival rates in these birds, so data from other similar species were used instead (assuming that survival in stilts would be similar to survival in these closely related species). A **“sensitivity analysis”** was conducted in which the survival values were systematically varied to see how much they affected the model results. This analysis suggested that the survival estimates would have to be wrong by quite a lot for it to alter the conclusions.
- v) Hence, even though some of the numbers were not well known (and thus the model is far from perfect), the conclusion was that it probably did an OK job of predicting extinction risk despite the imperfections.