A Probabilistic Population Model with Cyclic Birth Rates

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Background

- · A model is used to simulate population changes in a group of organisms living in a single region
- Random processes are inherent to actual living populations, but they can be extremely complex, so difficult to describe mathematically
- To simplify: approximate possible events on an individual scale, and then simulate them for a larger group (implementing mean field theory)

An extension of a mean field Bolker-Pacala Model

- Each event is modeled by an exponential random variable whose parameters we control
- This makes more efficient simulation of a large and complex population possible

Model Parameters

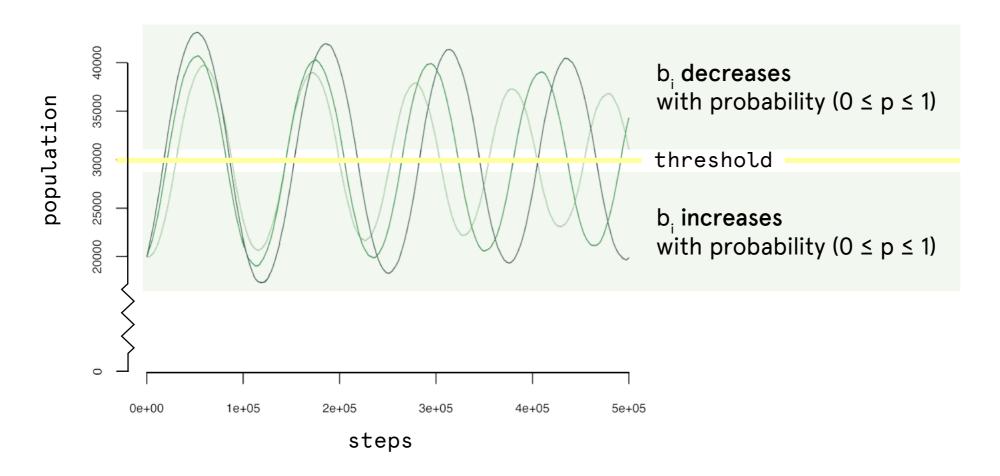
birth rate after ith event
 μ death rate (natural)
 a death rate (competition)
 birth rate after ith event
 j initial population
 population threshold
 μ of locations in region
 resource growth rate
 p probability of change in b

Probability of population increase + Probability of population decrease = 1

$$\frac{b_i j}{b_i j + \mu j + aj^2 L} + \frac{\mu j + aj^2 L}{b_i j + \mu j + aj^2 L} = 1$$

Cyclic Birth Rate

 Births become more likely when the population is below a certain threshold, and less likely when the population is above that threshold



- These birth rate cycles result in oscillations, whose period and amplitude are affected by the input parameters
- Two methods for implementing: input-based vs. density-based

Key Features of Population Oscillations

Max (peak) and min values

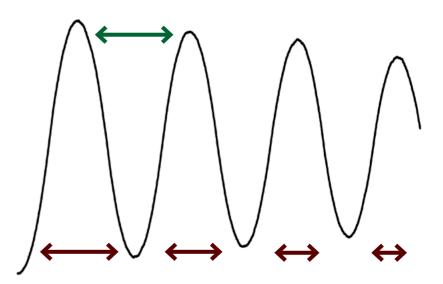
Overall relatively consistent for any given starting population (despite randomness inherent to model)

Dampening effect

Stronger as competition rates increase

Distance between peaks

(Related to the "slowing factor")
Varies according to p, the probability
that the birth rate will change

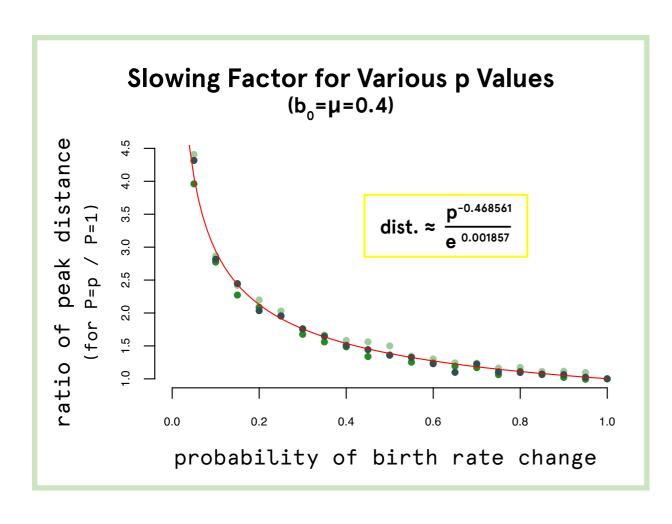


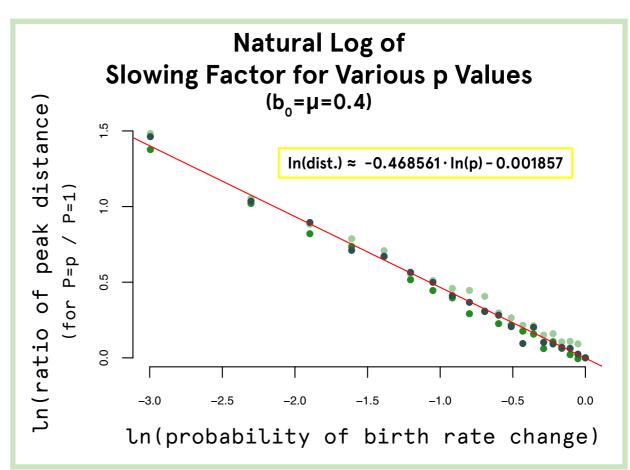
Condensing effect

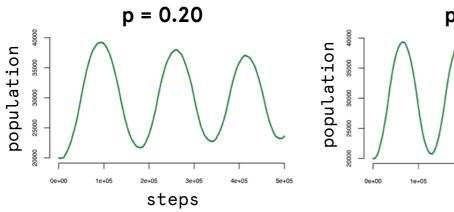
Stronger as birth rate exceeds death rate

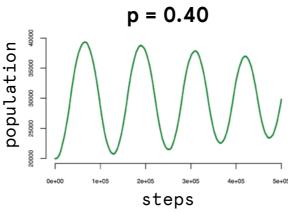
Slowing Factor Caused by p

(for initial birth rate = death rate)

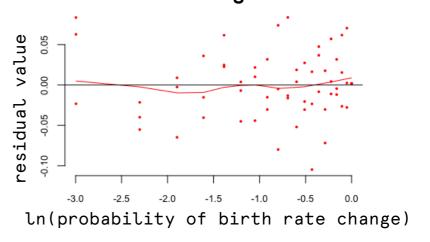






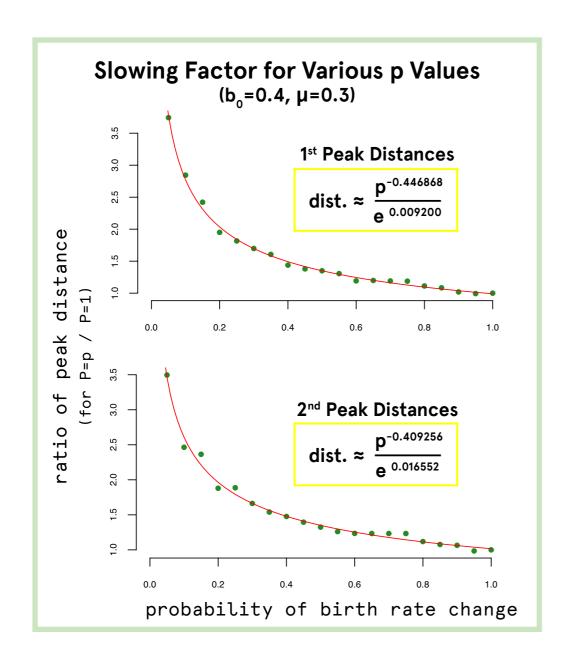


Residuals vs. Fitted Logarithmic Curve



Slowing Factor Caused by p (for initial birth rate > death rate)

- Note that condensing effect begins to have an impact when b_i > µ
 (peak distances are no longer constant for any given p)
- · So, we compare peaks separately (1st-to-1st, 2nd-to-2nd)



Input Parameters		Exponent of p for 2 nd Peak Distance (= slope of log line)
$\mu = 0.4$ $(b_0 = 0.4)$	-0.449470	-0.416982
$\mu = 0.3$ $(b_0 = 0.4)$	-0.446868	-0.409256
$\mu = 0.2$ $(b_0 = 0.4)$	-0.492693	-0.42614
$\mu = 0.1$ $(b_0 = 0.4)$	-0.60194	

Conclusions and Further Study

- We are able to approximate power functions that relate the probability of birth rate change to a slowing factor in the oscillations
- Because of the stochastic processes in the simulation, these equations should be improved by conducting many more trials
- While these models do incorporate realistic events in a population, the assignments of event probabilities should be more fine-tuned to increase their meaningfulness
 - Parameters for b_i, u, a, and r should reflect actual ecological dynamics (e.g. plant species in a terrain, or human social castes in a country)
- Switching to the more realistic density-based model, where p changes in response to the threshold, may provide different solutions
- In future, these models could be extended to show interactions of multiple population groups in the same ecological system