

Quantitative Genetics Approaches to Modeling Adaptive Evolution

Colleen T. Webb
Department of Biology
Colorado State University

Game Plan

- Quantitative genetics background
- Two approaches
 - Deterministic Gradient Equations (*sensu* Lande)
 - Stochastic Adaptive Dynamics (*sensu* Dieckmann)
- Pros and cons of these approaches
- Examples of theory applications
- Example of a real world system

Modeling Goals

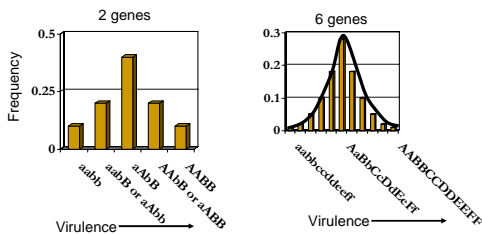
- Explore adaptive evolutionary processes underlying evolutionary patterns
- Explicitly link evolutionary consequences of ecological interactions
- Usually applied to theory problems (but not always)

Defining Quantitative Traits

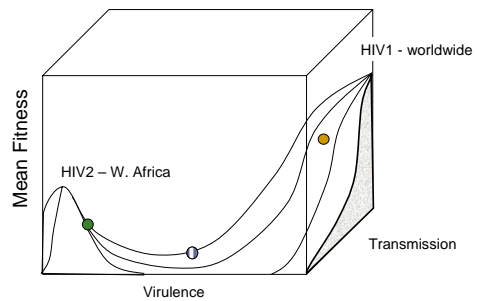
- Continuous, measurable, non-categorizable
- Genotypic and environmental effects determine trait value
- Usually follow phenotype
- Modeling at phenotypic level

Distribution of Quantitative Traits

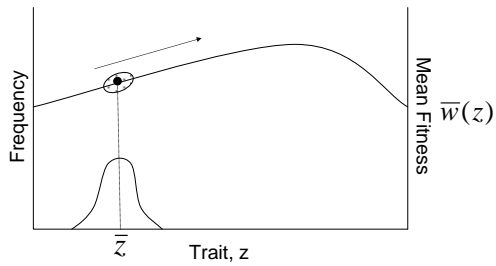
- It's Normal, Gaussian, Bell shaped



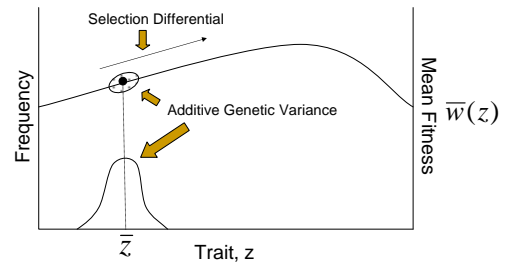
Adaptive Landscapes



Describing Movement across the Landscape



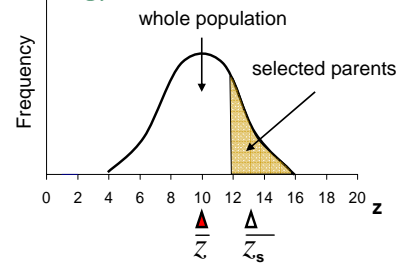
Describing Movement Depends on...



Deterministic Modeling Approaches

- Describe rate of change for mean value of a quantitative trait
- Based on Wright's work for discrete traits
- Lande 1976
- Breeder's Equation

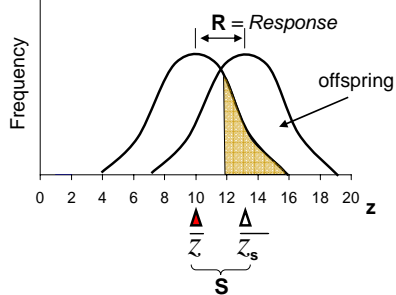
Terminology



\bar{z} is the mean of the whole population

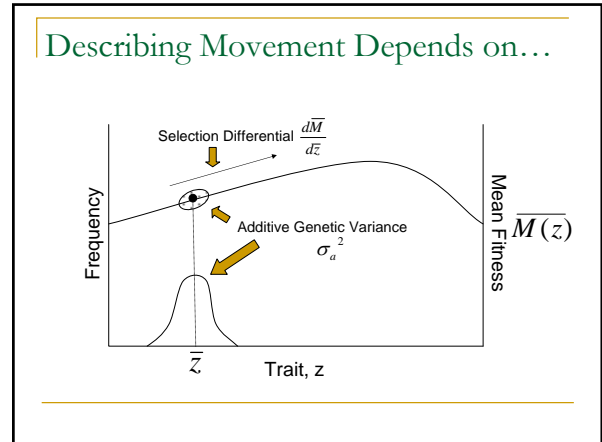
\bar{z}_s is the mean of the population after selection (the parents that get to reproduce)

Terminology



Putting It Together: The Breeder's Equation to Lande's Gradient Equation

$$\begin{aligned}
\frac{d\bar{z}}{dt} &\approx \Delta\bar{z} \\
&= \bar{z}_{original} - \bar{z}_{offspring} \\
&= R \\
&= (\bar{z}_s(t) - \bar{z}(t))h^2 && \text{Breeder's Equation} \\
&= \frac{\sigma_p^2}{\bar{w}} \frac{d\bar{w}}{d\bar{z}} h^2 && \text{Magic (i.e., Lande 1976)} \\
&= \sigma_a^2 \frac{1}{\bar{w}} \frac{d\bar{w}}{d\bar{z}} \\
&= \sigma_a^2 \frac{d \ln(\bar{w})}{d\bar{z}} \\
&= \sigma_a^2 \frac{d\bar{M}}{d\bar{z}}
\end{aligned}$$



Lande's Equation Combines These...

$$\frac{d\bar{z}}{dt} = \sigma_a^2 \frac{d\bar{M}(z)}{d\bar{z}}$$

- Lots of assumptions
- User beware
- Assumptions mostly reasonable
- Most accurate under weak selection

- ### Stochastic Simulation Approaches
- Adaptive Dynamics
 - Dieckmann, Geritz, others
 - Simulates "mutational stochasticity"
 - Based on ESS theory
 - Underlying fitness function

- ### ESS Theory in One Slide!
- Compare the fitness of a rare mutant to a resident type to see if the mutant can invade
 - We compare fitness(mutant|resident) to fitness(resident|resident)
 - If fitness(mutant|resident) is greater than fitness(resident|resident) then, mutant invades
 - Somewhat simplified!

- ### Adaptive Dynamics Simulation
- Mutants appear in the population stochastically
 - Fitnesses compared
 - Type in population is updated

- These approaches have different assumptions and the potential to give quite different results, but in practice they frequently give similar results

Deterministic Approaches

- Pros
 - Analytical approach makes it easy to understand dynamics
 - Tries to realistically mimic mutation and selection processes
- Cons
 - Constant variance assumption and disruptive selection
 - Applicability to frequency dependent selection?
 - Standing variance vs. mutation-selection balance

Stochastic Approaches

- Pros
 - Can use any type of fitness function
 - Relatively easy to analyze
 - Models of speciation
- Cons
 - Potentially computationally intensive
 - Mutation-selection process unnatural

Fitness Functions Are Everything

- Measure fitness function experimentally
- Pull fitness function from underlying ecological model (most common)
 - $\frac{dx}{dt} = f(x, z)x$
 - f describes per capita growth rate, which is a measure of fitness

Making Ecology Explicit

- Usual assumption is that underlying ecological model is at equilibrium
- Ecology can be made explicit
 - Deterministic model with interaction between ecological and evolutionary timescales
 - Stochastic model with both mutational and demographic stochasticity

Applications

- Most often used in theoretical systems, but can be applied to real systems
- Two theory examples
 - Darwinian Extinction/Evolutionary Suicide
 - Evolutionary Branching (underlying phylogenetic patterns)
- Real system example
 - Plague in prairie dogs

Darwinian Extinction

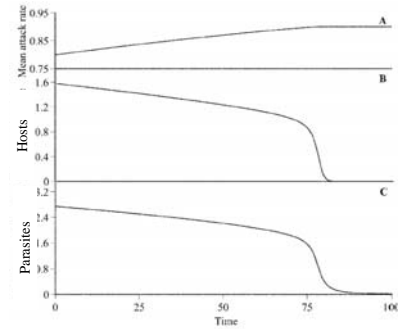
- Lande's approach with explicit ecology

$$\frac{dx}{dt} = x(-1 + rx - x^2 - zy)$$

$$\frac{dy}{dt} = y(-d - y + (z+1)x)$$

$$\frac{dz}{dt} = \sigma_a^2 \frac{df}{dz} = \sigma_a^2 x$$

Webb 2003



Webb 2003

Evolutionary Branching

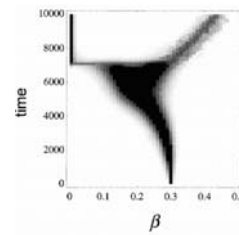
- Adaptive Dynamics

$$\frac{dx_i}{dt} = r(\beta_i)x_i - q(\sum x_i + y)x_i - \beta_i x_i y$$

$$\frac{dy}{dt} = \sum \beta_i x_i y - \Gamma y$$

Boots and Haraguchi 1999

Tradeoffs and Evolutionary Impacts



Boots and Haraguchi 1999

Plague and Prairie Dogs

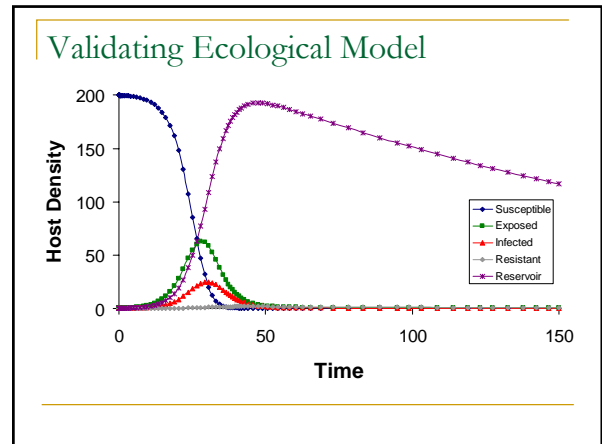
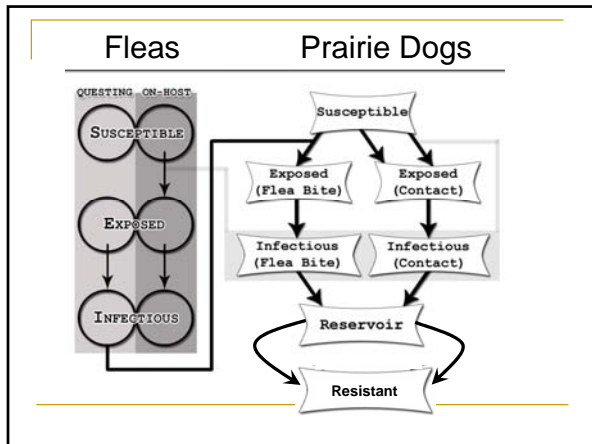
with Brandon Goodell and Mike Antolin

- Highly virulent with 95-100% mortality
- It's been this way for a long time



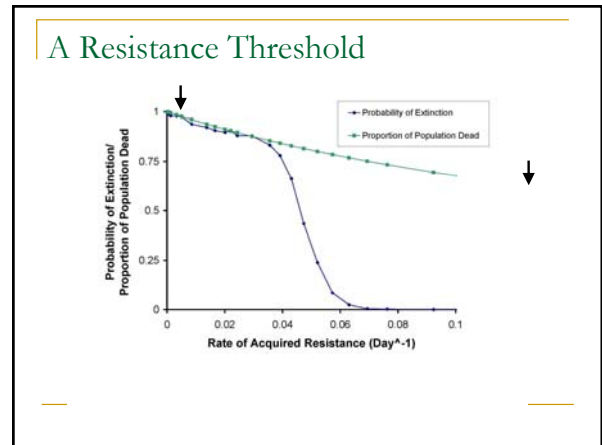
Motivation

- Variation in levels of resistance among hosts
- In prairie dogs: Strong selection pressure, weak response – What's up?



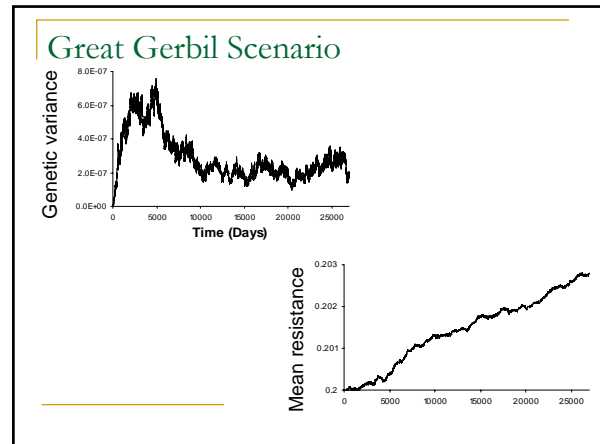
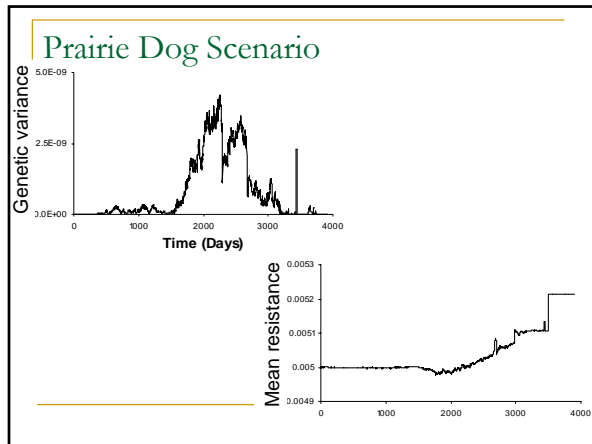
Quantitative Validation

	Model	Field Data
Probability of Extinction	98%	12 of 12 epizootics
Time to Extinction	~ 14 weeks	~ 6-8 weeks



- Adding Genetic Variation**
- Simulation model
 - Stochastic births and deaths (demographic stochasticity)
 - When birth occurs, offspring has probability of slightly different resistance (mutational stochasticity)

- Adding Selection**
- Plague returns to a town approximately every 7 years
 - Pulsed plague from the distribution of plague return times



- ### When is demographic stochasticity important?
- Race between evolutionary and demographic stochasticity processes
 - Theory suggests critical population size is 10-100 individuals
 - Prairie dog models - about 6 resistant individuals usually survive
 - Great gerbil models – many individuals survive
- Gomulkiewicz and Holt 1995
Lande 1993

- ### Initial conditions matter
- Different initial conditions (i.e., levels of resistance) is what drives differences in population persistence
 - Threshold exists such that some populations/species are unable to maintain evolutionary response despite strong selection pressure

- ### Acknowledgements
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