



# Overview

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Eco-Evolutionary Dynamics

Modelling Frameworks

Mathematical Connections

Biodiversity Dynamics

Adaptive Speciation

Niche Theory



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2+

# Adaptive dynamics



# Adaptive Dynamics Theory

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The theory of adaptive dynamics extends evolutionary game theory and quantitative genetics theory in a number of respects:

- Evolving traits are continuous
- Trait dynamics are described
- Mutational covariances and constraints can be examined
- Arbitrary density and frequency dependence is allowed
- Coevolution is integrated
- Structured population dynamics are allowed
- Non-equilibrium population dynamics are allowed
- Fitness landscapes are derived

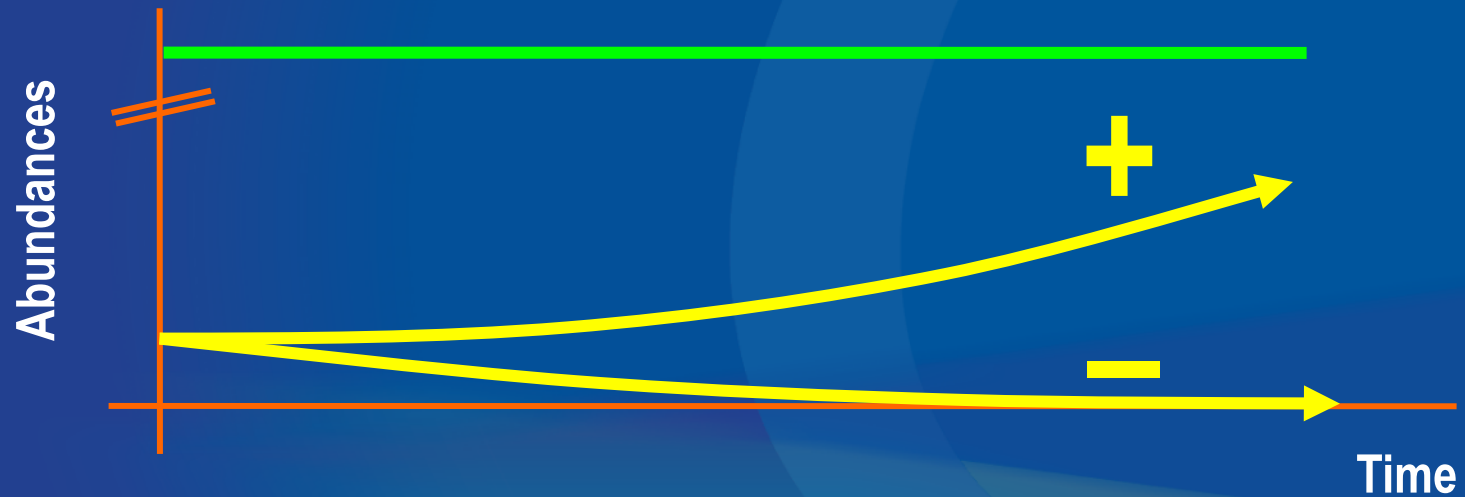
Extensions  
of EGT

Extensions  
of both  
EGT  
and QGT

# Invasion Fitness

Metz et al. (1992)

- Initial per capita growth rate of a small **variant population** within a large **resident population** at ecological equilibrium:



Matsuda & Abrams (1985)  
 Metz et al. (1992)

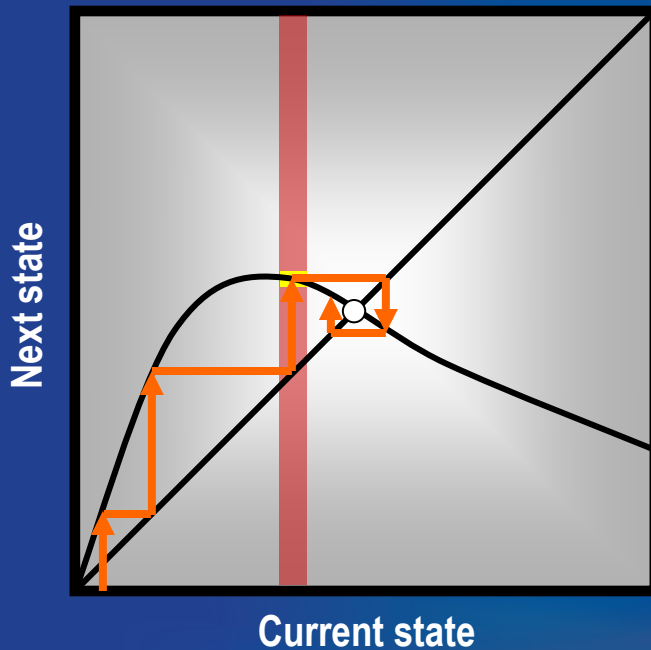
# Pairwise Invasibility Plots



- +** Invasion of the variant into the resident population is possible
- Invasion is impossible
- ↑** One trait substitution
- ↗** Trait substitution sequence
- Evolutionarily singular phenotype

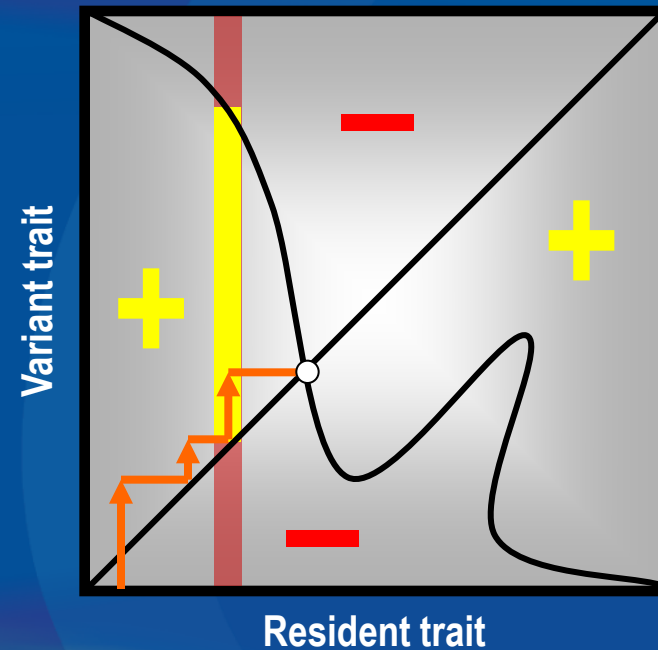
# Comparison with Recursions

## ■ Recursion relations



Size of vertical steps deterministic

## ■ Trait substitutions

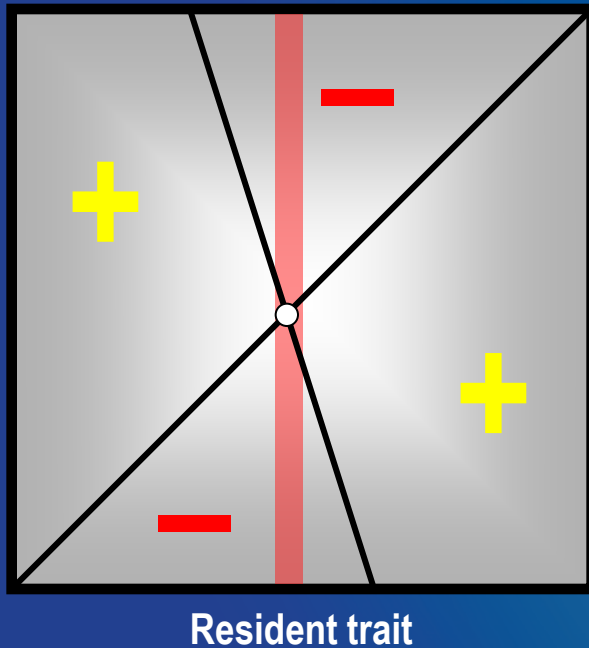


Size of vertical steps stochastic

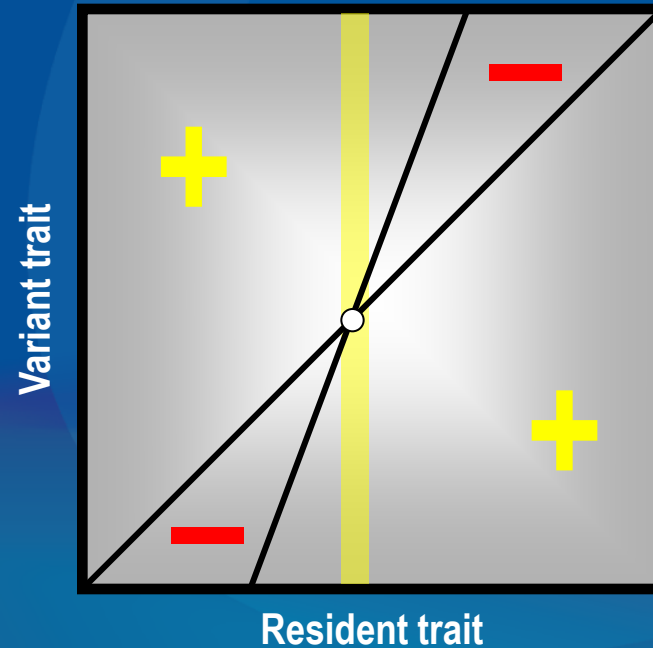
# Reading PIPs: Evolutionary Stability

- Is a singular phenotype immune to invasions by neighboring phenotypes?

Yes:



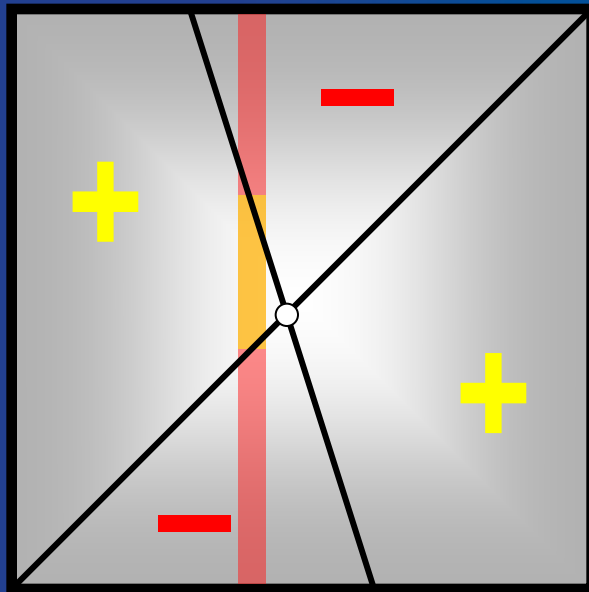
No:



# Reading PIPs: Convergence Stability

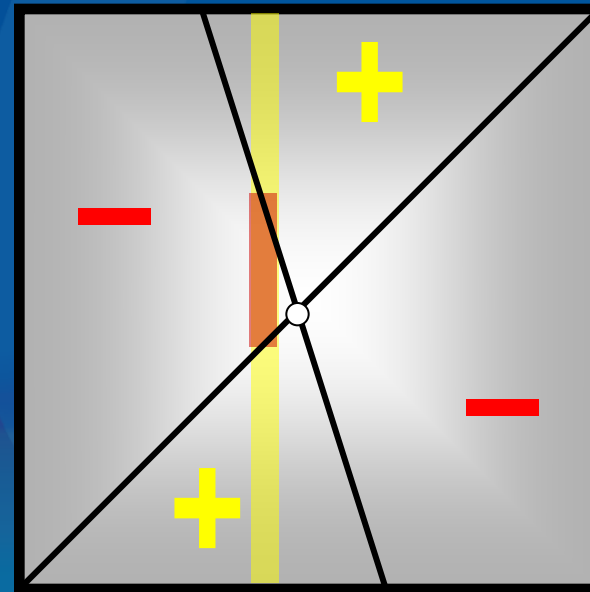
- When starting from neighboring phenotypes, do successful invaders lie closer to the singular one?

Yes:



Resident trait

No:



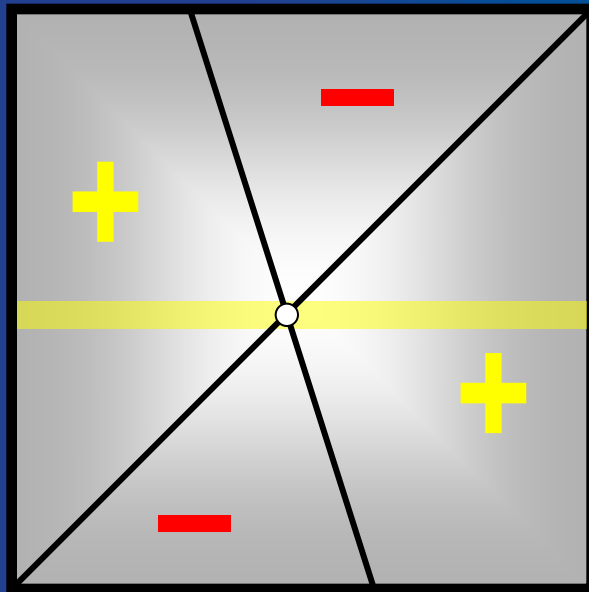
Resident trait



# Reading PIPs: Invasion Potential

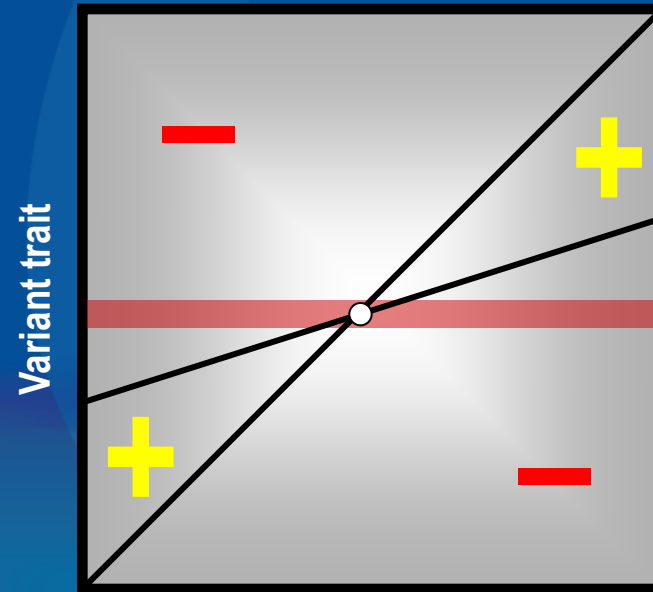
- Is the singular phenotype capable of invading into all its neighboring types?

Yes:



Resident trait

No:

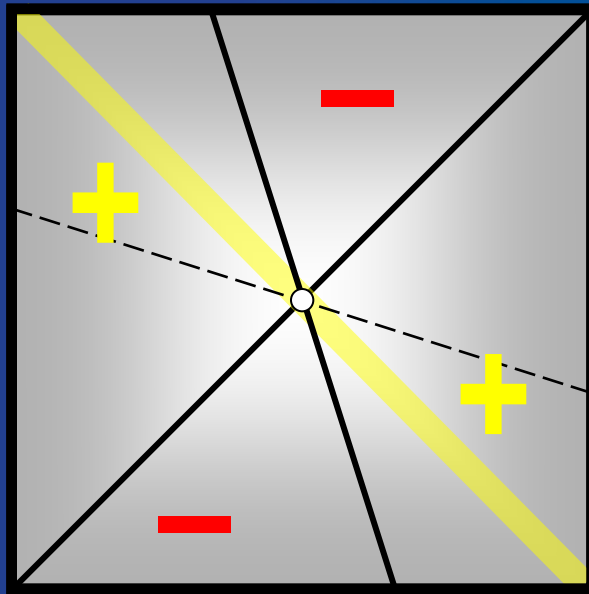


Resident trait

# Reading PIPs: Mutual Invasibility

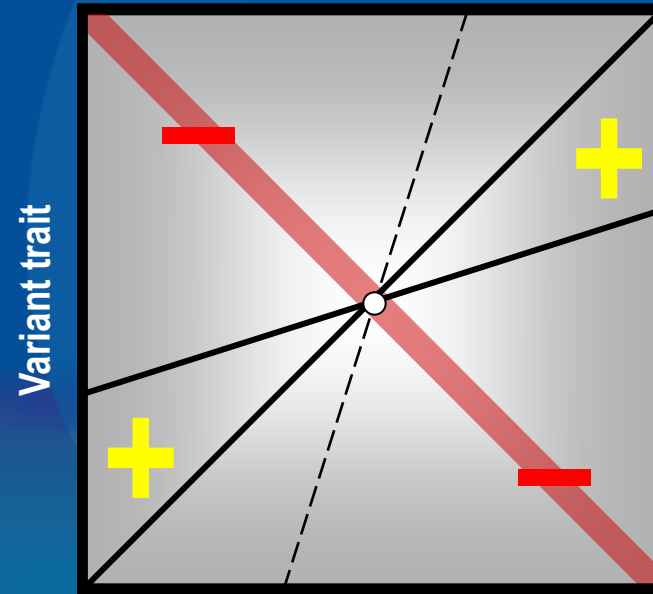
- Can a pair of neighboring phenotypes on either side of a singular one invade each other?

Yes:



Resident trait

No:



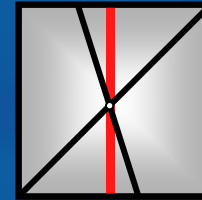
Resident trait

# Reading PIPs: Four Independent Properties

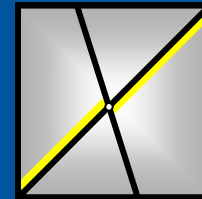
Geritz et al. (1997)

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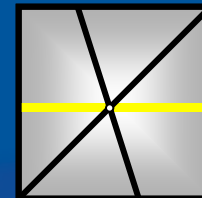
■ Evolutionary stability



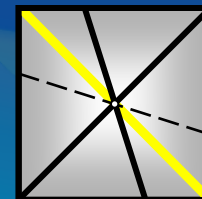
■ Convergence stability



■ Invasion potential

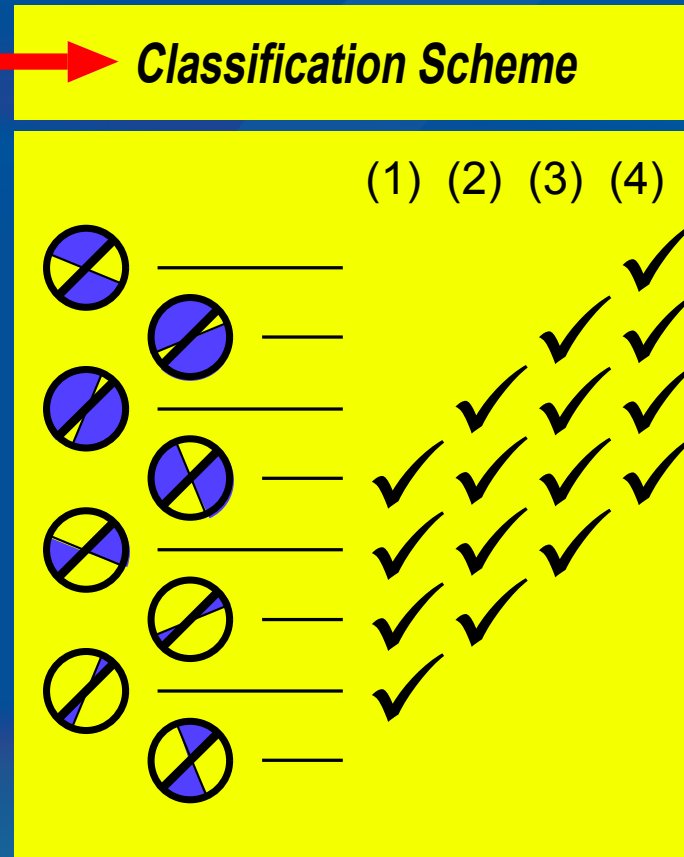
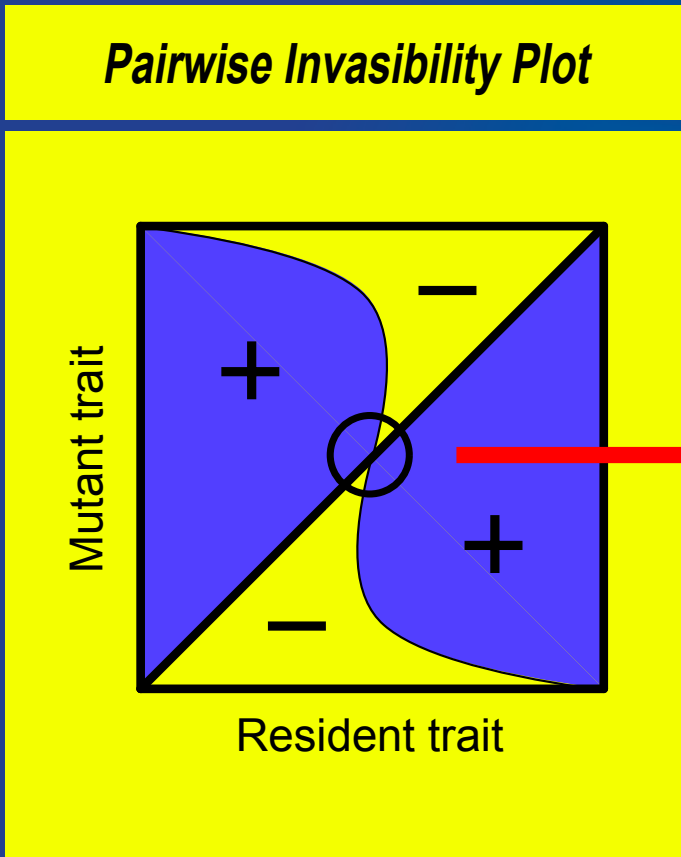


■ Mutual invasibility



# Reading PIPs: Eightfold Classification

Geritz et al. (1997)  
Dieckmann (1997)

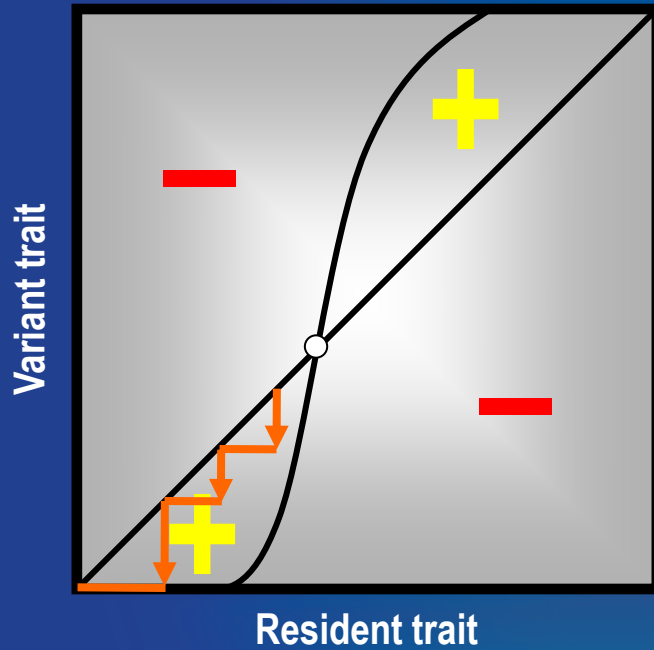


↑ ↑ ↑ ↑ ↑ ↑ ↑ ↑  
Evolutionary bifurcations

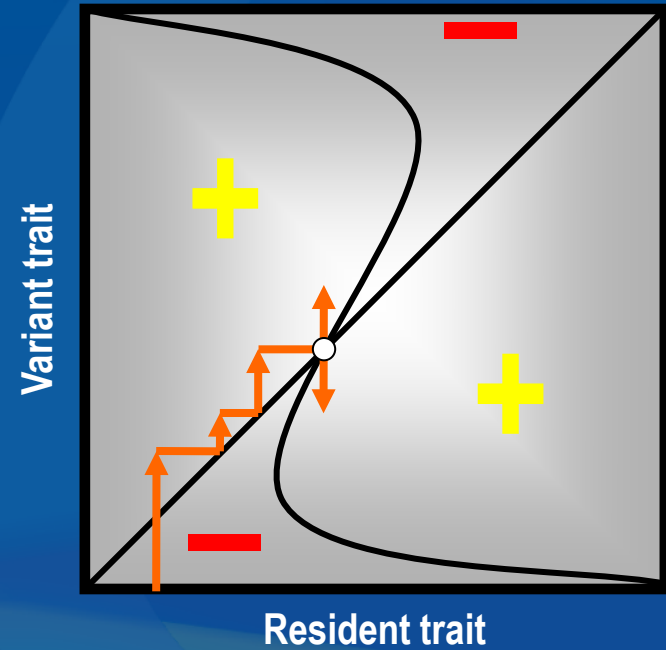
(1) Evolutionary stability, (2) Convergence stability, (3) Invasion potential, (4) Mutual invasibility

# Two Interesting Types of PIPs

## ■ Garden of Eden

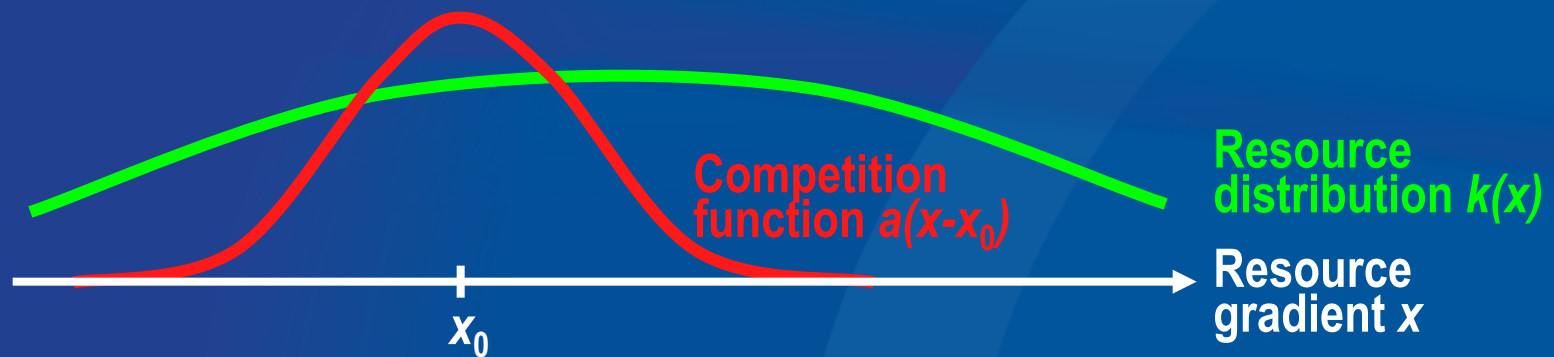


## ■ Branching Point



# Example: Resource Competition

Roughgarden (1976)



Dynamics of population sizes  $n_i$  of strategy  $x_i$

$$\frac{d}{dt} n_i = r n_i \left[ 1 - \frac{1}{k(x_i)} \sum_j \frac{a(x_i - x_j) n_j}{k(x_j)} \right]$$

# Analysis of Example 1/2

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## ■ Invasion Fitness

$$f(x', x) = r \left[ 1 - \frac{1}{k(x')} (a(0)n' + a(x' - x)n) \right]$$



$$n' \rightarrow 0$$



$$n \rightarrow n_{\text{eq}} = k(x)$$

$$f(x', x) = r \left[ 1 - a(x' - x) \frac{k(x)}{k(x')} \right]$$

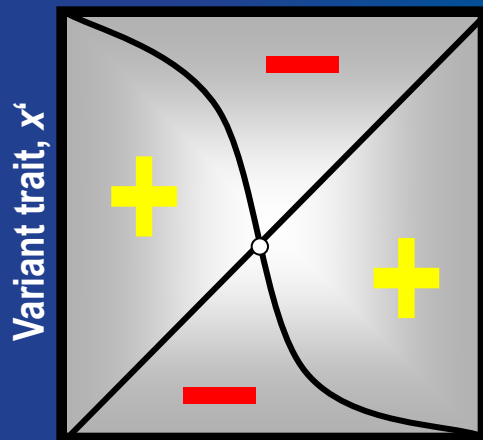
# Analysis of Example 2/2

## ■ Pairwise Invasibility Plots

With  $k = k_0 N(0, \sigma_k)$  and  $a = N(0, \sigma_a)$  we obtain

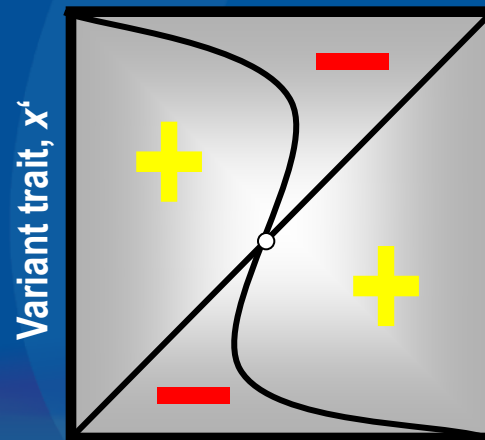
for  $\sigma_a > \sigma_k$

for  $\sigma_a < \sigma_k$



Resident trait, x

Evolutionary stability

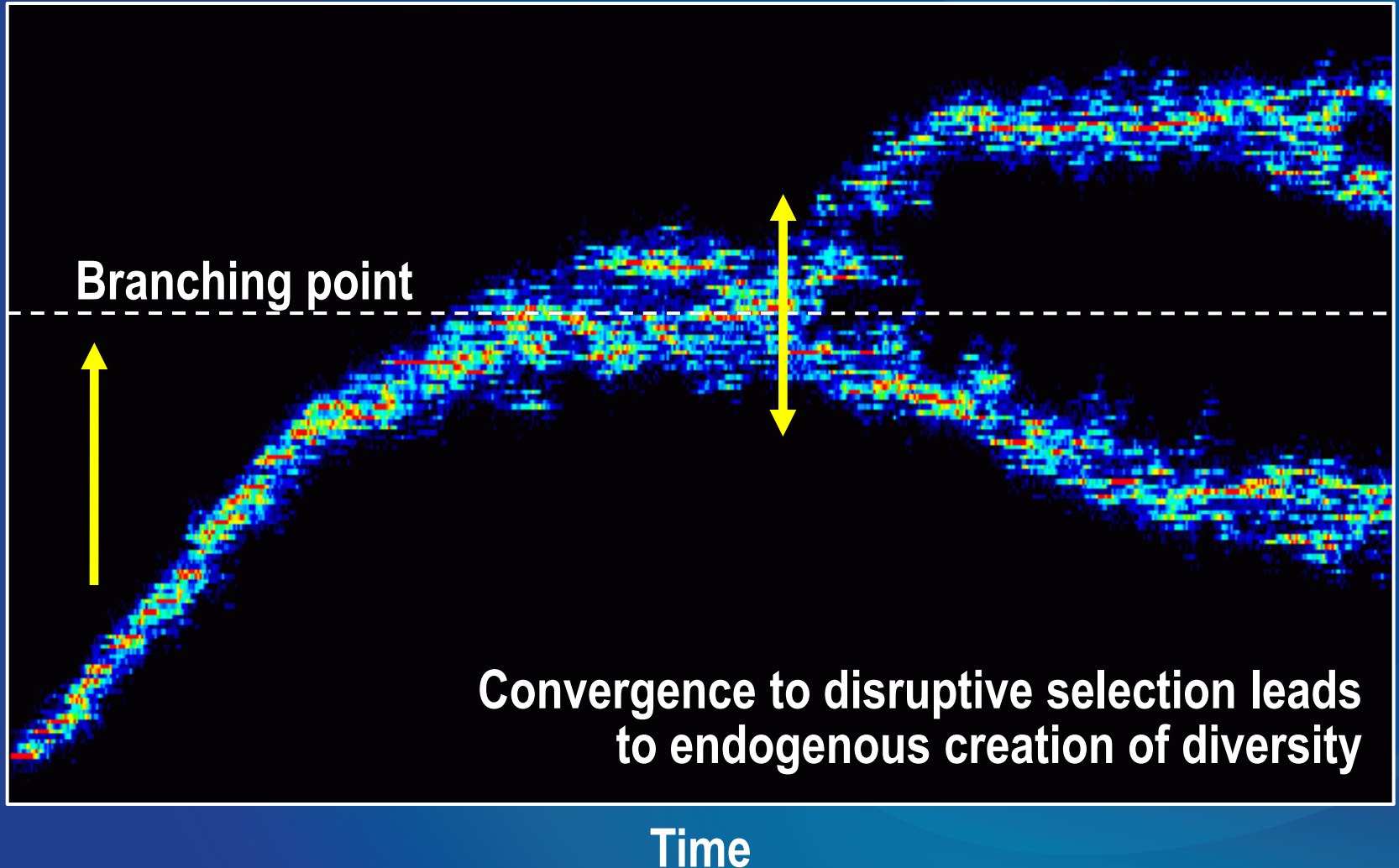


Resident trait, x

Evolutionary branching



# Evolutionary Branching





# Published Analyses 1/2

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- Symmetric intraspecific competition (Doebeli 1996a, 1996b; Metz et al. 1996; Dieckmann and Doebeli 1999)
- Asymmetric intraspecific competition (Kisdi 1999; Doebeli and Dieckmann 2000; Kisdi et al. 2001)
- Interspecific competition (Law et al. 1997; Kisdi and Geritz 2001)
- Resource specialization (Meszéna et al. 1997; Geritz et al. 1998; Day 2000; Kisdi 2001; Schreiber and Tobison 2003; Egas et al. 2004, 2005)
- Ontogenetic niche shifts (Claessen and Dieckmann 2002)
- Mixotrophy (Troost et al. 2005)
- Phenotypic plasticity (Van Dooren and Leimar 2003; Ernande and Dieckmann 2004; Leimar 2005)
- Dispersal evolution (Doebeli and Ruxton 1997; Johst et al. 1999; Parvinen 1999; Mathias et al. 2001; Parvinen and Egas 2004),
- Mutualism (Doebeli and Dieckmann 2000; Law et al. 2001; Ferdy et al. 2002; Ferrière et al. 2002; Day and Young 2004)
- Emergent cooperation (Doebeli et al. 2004)



# Published Analyses 2/2

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- Predator-prey interactions (Brown and Pavlovic 1992; Van der Laan and Hogeweg 1995; Doebeli and Dieckmann 2000; Bowers et al. 2003)
- Cannibalism (Dercole 2003)
- Host-parasite interactions (Boots and Haraguchi 1999; Koella and Doebeli 1999; Regoes et al. 2000; Gudelj et al. 2004)
- Sex-ratio evolution (Metz et al. 1992; Reuter et al. 2004)
- Evolution of selfing (Cheptou and Mathias 2001; De Jong and Geritz 2001)
- Evolution of mating traits (Van Doorn et al. 2001, 2004)
- Evolution of anisogamy (Maire et al. 2001)
- Seed evolution (Geritz et al. 1999; Mathias and Kisdi 2002)
- Microbial cross-feeding (Doebeli 2002)
- Prebiotic evolution (Meszéna and Szathmáry 2001)
- Community assembly (Jansen and Mulder 1999; Bonsall et al. 2004)
- Food-web formation (Loeuille and Loreau 2005; Ito et al. 2009; Brännström et al. 2010)



# Canonical Equation

Dieckmann & Law (1996)

- **Assumptions:** Populations are large, and mutational steps are both rare and small
- Then, the evolutionary rates are given by

$$\frac{d}{dt} x_i = \frac{1}{2} \mu_i(x_i) n_i^*(x) \sigma_i^2(x_i) \frac{\partial}{\partial x'_i} f_i(x'_i, x) \Big|_{x'_i=x_i}$$

↑  
evolutionary  
rate in species  $i$

↑  
mutation  
probability

↑  
equilibrium  
population  
size

↑  
mutation  
variance-covariance

↑  
local  
selection  
gradient

↑  
invasion  
fitness



# Synthetic Perspective

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- The common form of the canonical equation of adaptive dynamics and of Lande's equation of quantitative genetics is reassuring
- In first approximation, evolutionary rates are proportional to the gradient of a frequency-dependent fitness
- Beyond this commonality, the two dynamics differ, describing different kinds of evolutionary processes

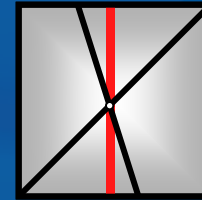


# Reminder: Four Independent Properties

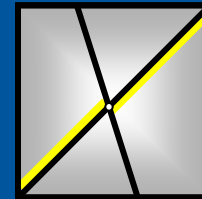
Geritz et al. (1997)

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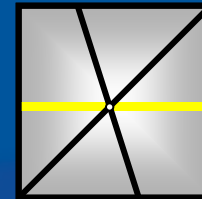
■ Evolutionary stability



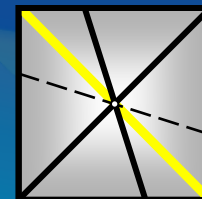
■ Convergence stability



■ Invasion potential



■ Mutual invasibility





# Fitness, Gradient, Hessians

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## ■ Invasion fitness

$$f(x', x)$$

$$f(x, x) = 0 \quad \text{at ecological equilibrium}$$

## ■ Selection gradient

$$g(x) = \frac{\partial}{\partial x'} f(x', x) \Big|_{x'=x}$$

$$g(x^*) = 0 \quad \text{at evolutionary equilibrium}$$

## ■ Hessians of invasion fitness

$$h_{mm} = \frac{\partial^2}{\partial x'^2} f(x', x) \Big|_{x'=x=x^*}$$

$$h_{rm} = \frac{\partial^2}{\partial x' \partial x} f(x', x) \Big|_{x'=x=x^*}$$

$$h_{rr} = \frac{\partial^2}{\partial x^2} f(x', x) \Big|_{x'=x=x^*}$$

$$h_{mr} = \frac{\partial^2}{\partial x \partial x'} f(x', x) \Big|_{x'=x=x^*}$$



# Conditions for Evolutionary Branching

- Evolutionary stability: no

$$h_{mm} < 0$$

- Convergence stability: yes

$$h_{mm} - h_{rr} < 0 \quad (\text{or } h_{mm} + h_{mr} < 0)$$

- Invasion potential

$$h_{rr} > 0$$

- Mutual invasibility

$$h_{mm} + h_{rr} > 0$$

These conditions apply to one-dimensional traits.

For higher-dimensional traits, they involve matrices and are a bit more complex.





# Summary:

## Main Tools of Adaptive Dynamics Theory

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- Individual-based modelling
- Invasion fitness
- Pairwise invasibility plots
- Canonical equation
- Evolutionary branching conditions



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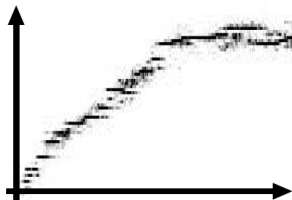


# **Interlude: Mathematical Connections**

# Four Types of Evolutionary Dynamics

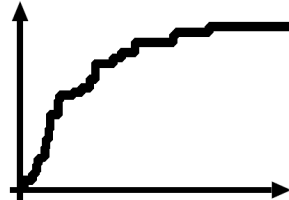
Polymorphic  
stochastic

Individual-  
based  
evolutionary  
dynamics



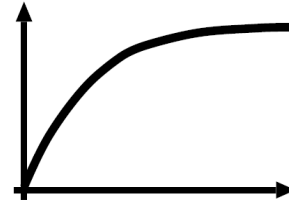
Monomorphic  
stochastic

Evolutionary  
random  
walk



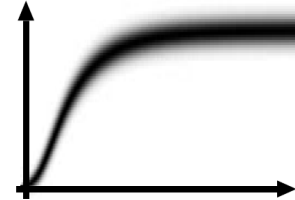
Monomorphic  
deterministic

Gradient-  
ascent on  
adaptive  
landscape



Polymorphic  
deterministic

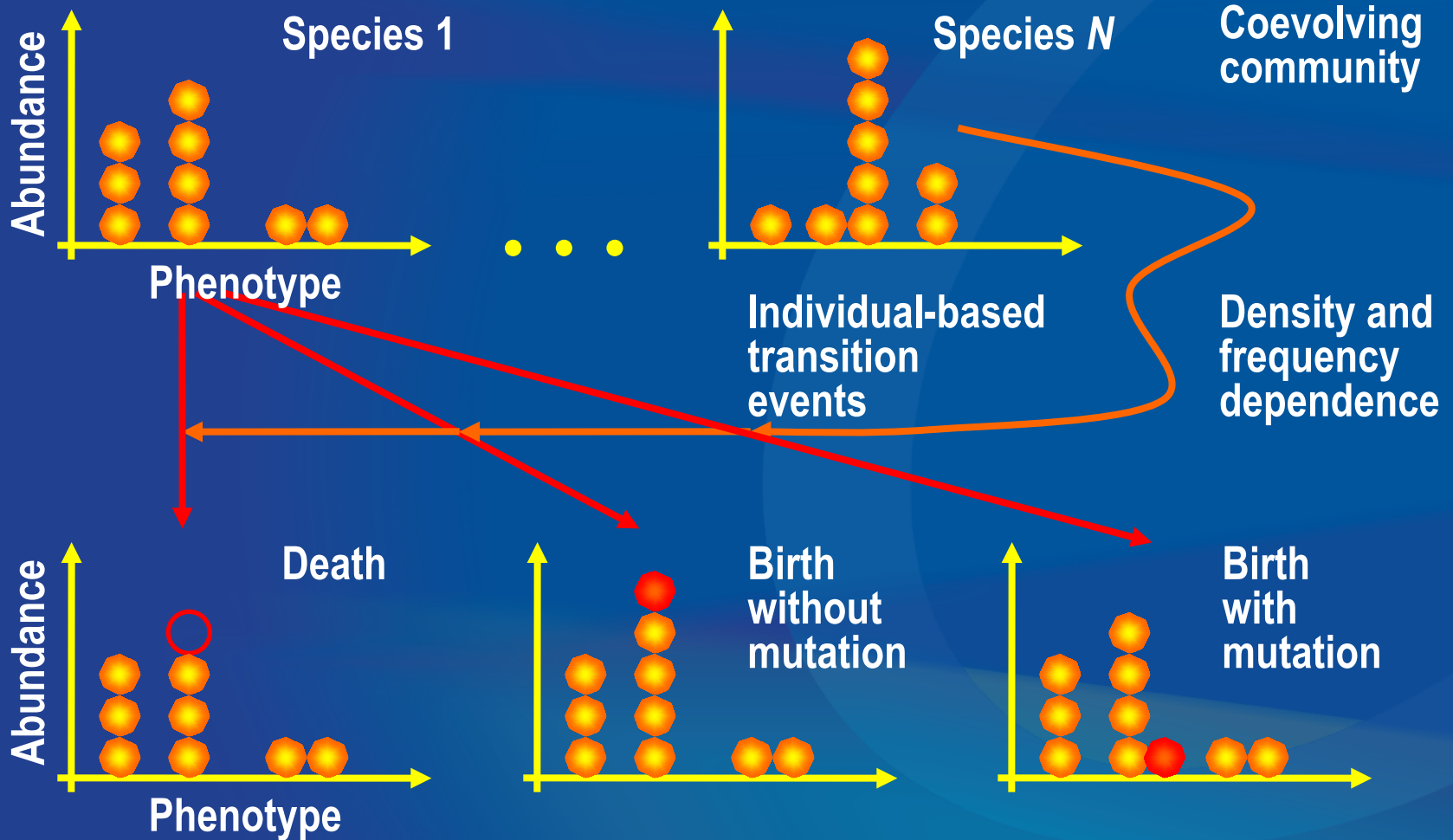
Reaction-  
diffusion  
dynamics





# Birth-Death-Mutation Processes

Dieckmann (1994)





# Individual-based Evolutionary Dynamics

Polymorphic and Stochastic

Dieckmann (1994)

## ■ State variables

$$p_i = \sum_{k=1}^{n_i} \delta_{x_{ik}}$$

Phenotypic distribution  
in species  $i$

$$p = (p_1, \dots, p_N)$$

Phenotypic distribution  
of entire community

## ■ Parameters

$$b_i(x_i, p)$$

Per capita birth rate  
in species  $i$

$$d_i(x_i, p)$$

Per capita death rate  
in species  $i$

$$\mu_i(x_i)$$

Mutation probability  
in species  $i$

$$M_i(x'_i, x_i)$$

Mutation distribution  
in species  $i$

$$f_i(x_i, p) = b_i(x_i, p) - d_i(x_i, p)$$

Fitness-generating function  
in species  $i$



# Individual-based Evolutionary Dynamics

Polymorphic and Stochastic

Dieckmann (1994)

## ■ Master equation

$$\frac{d}{dt}P(p) = \int [w(p | p')P(p') - w(p' | p)P(p)] dp'$$

$$w(p' | p) = \sum_{i=1}^N \int [d_i(x'_i, p) p_i(x'_i) \Delta(p_i - \delta_{x'_i} - p_i) \prod_{j \neq i} \Delta(p'_j - p_j) + b_i(x_i, p) p_i(x_i) B_i(x'_i, x_i) dx_i \Delta(p_i + \delta_{x'_i} - p_i) \prod_{j \neq i} \Delta(p'_j - p_j)] dx'_i$$

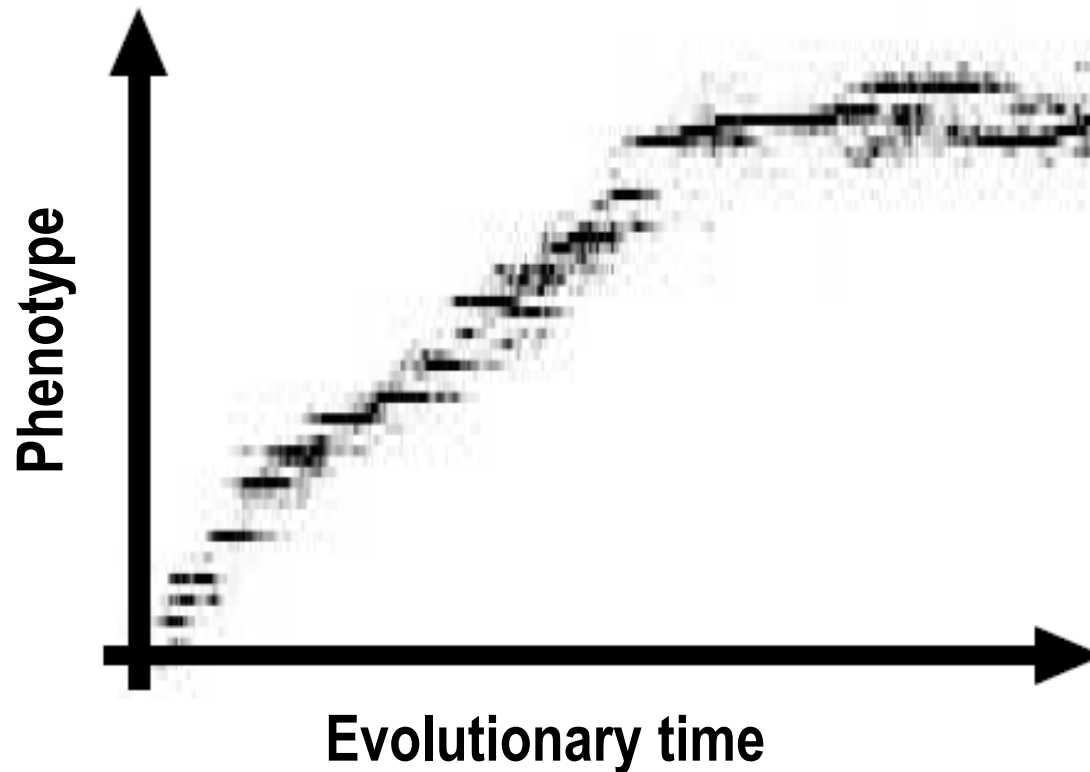
Measure-valued stochastic process in the space of atomic distributions (Dirac measures)

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# Individual-based Evolutionary Dynamics

Polymorphic and Stochastic

Dieckmann (1994)



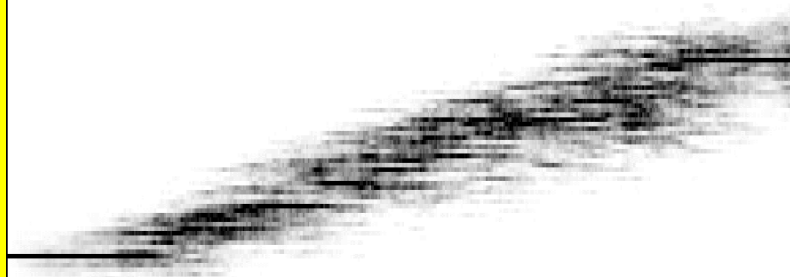


# Effect of Mutation Probability

**Large: 10%**

**Dynamic mutation-selection  
equilibrium**

Phenotype



**“Moving cloud”**

Evolutionary time

**Small: 0.1%**

**Mutation-limited  
evolutionary dynamics**



**“Staircase”**

Evolutionary time





# Probability for a Trait Substitution

Dieckmann & Law (1996)  
Geritz et al. (2002)

## Mutation

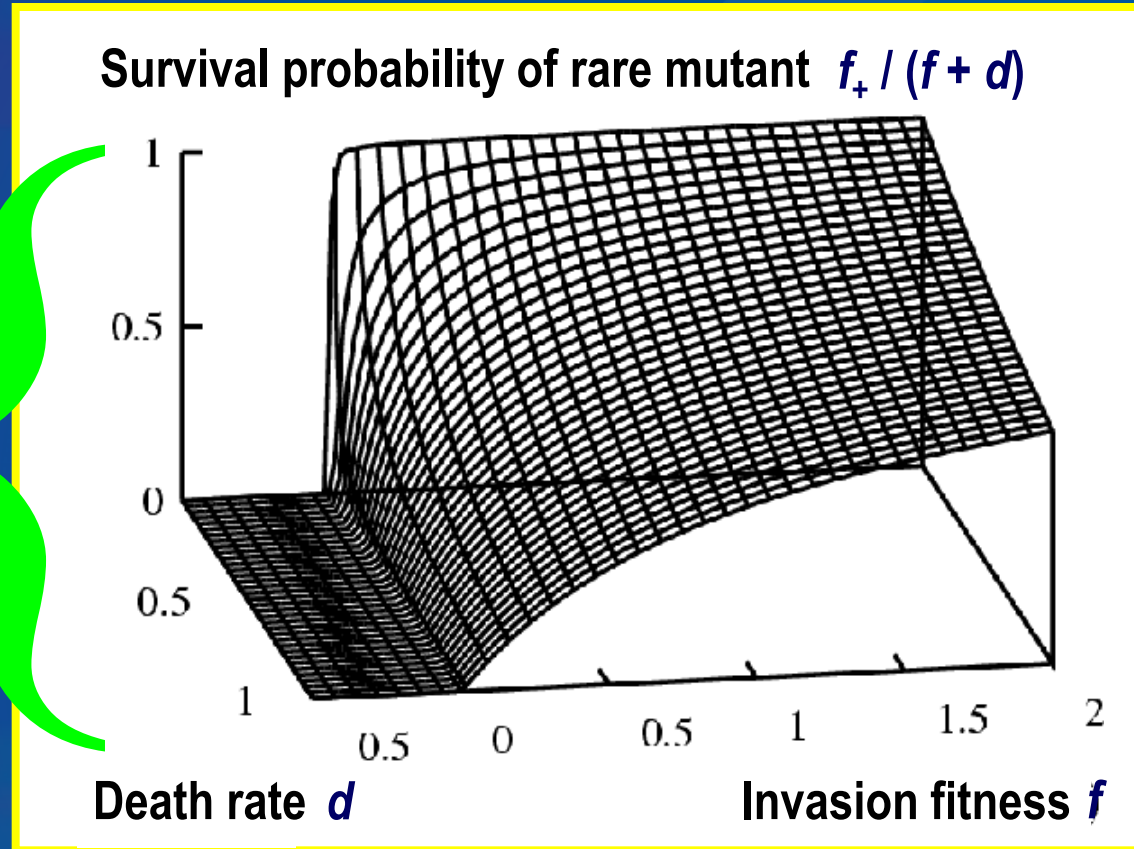
Population dynamics

## Invasion

Branching process theory

## Fixation

Invasion implies fixation



Invasion probabilities can alternatively be based on the Moran process, on diffusion approximations, or on graph topologies



# Evolutionary Random Walk

Monomorphic and Stochastic

Dieckmann & Law (1996)

## ■ Master equation

$$\frac{d}{dt} P(x) = \int [w(x | x') P(x') - w(x' | x) P(x)] dx'$$

$$w(x' | x) = \sum_{i=1}^N w_i(x'_i, x) \prod_{j \neq i} \delta(x'_j - x_j)$$

$$w_i(x'_i, x) = \mu_i(x_i) b_i(x_i, x) n_i^*(x) M_i(x'_i, x_i) \cdot (f_i(x'_i, x))_+ / b_i(x'_i, x)$$

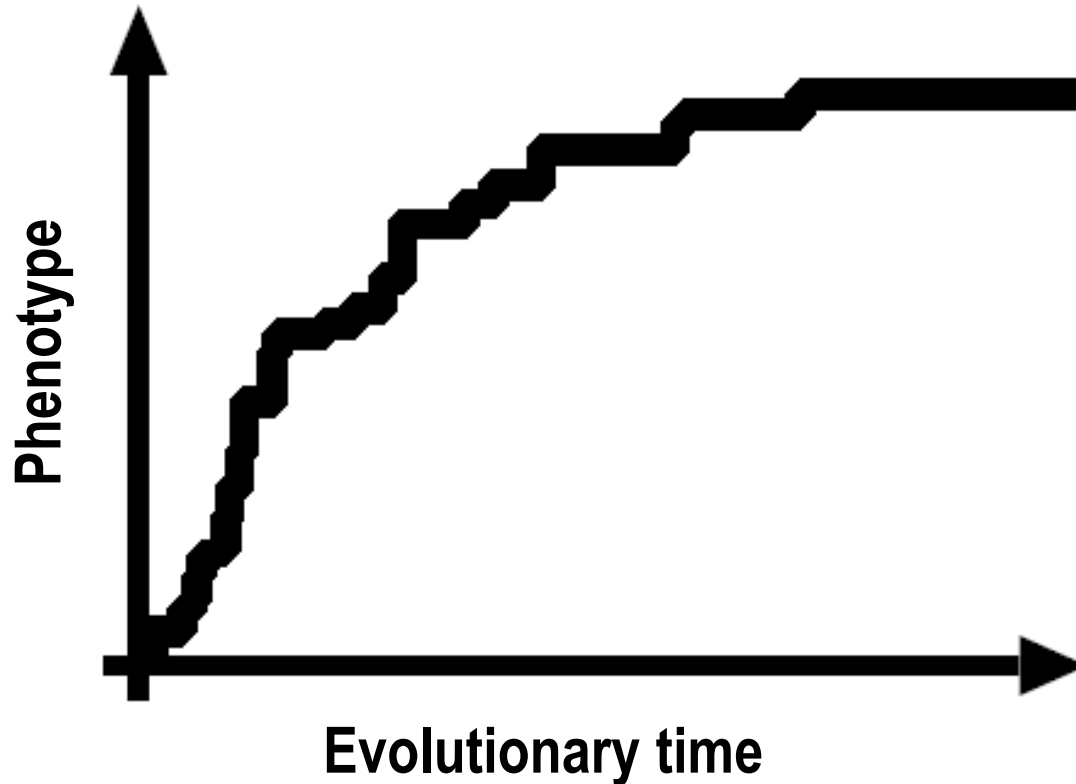
Real-valued stochastic process in trait space



# Evolutionary Random Walk

Monomorphic and Stochastic

Dieckmann & Law (1996)



# Gradient-Ascent on Adaptive Landscapes

Monomorphic and Deterministic

Dieckmann & Law (1996)

## ■ Canonical equation of adaptive dynamics

$$\frac{d}{dt} x_i = \frac{1}{2} \mu_i(x_i) n_i^*(x) \sigma_i^2(x_i) \left. \frac{\partial}{\partial x'_i} f_i(x'_i, x) \right|_{x'_i=x_i}$$

↑  
evolutionary  
rate in species  $i$

↑  
mutation  
probability

↑  
equilibrium  
population  
size

↑  
mutation  
variance-covariance

↑  
local  
selection  
gradient

↑  
fitness

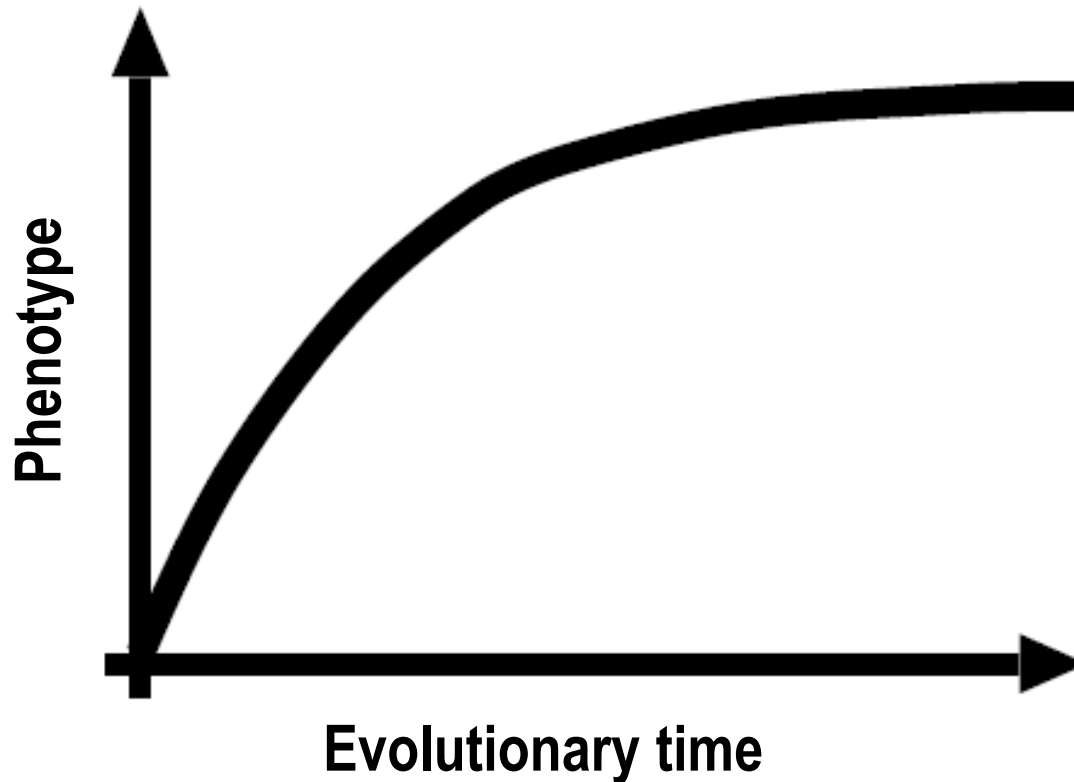




# Gradient-Ascent on Adaptive Landscapes

Monomorphic and Deterministic

Dieckmann & Law (1996)





# Reaction-Diffusion Dynamics

Kimura (1965)

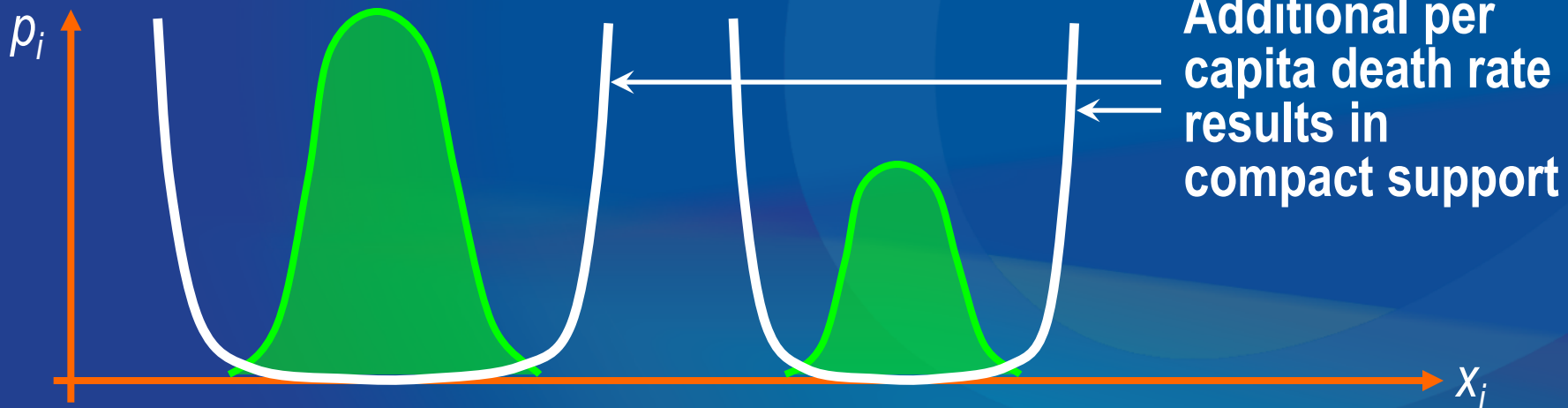
Polymorphic and Deterministic

Dieckmann (unpublished)

## ■ Kimura limit

$$\frac{d}{dt} p_i(x_i) = f_i(x_i, p) p_i(x_i) + \frac{1}{2} \mu_i(x_i) \sigma_i^2(x_i) * \frac{\partial^2}{\partial x_i^2} b_i(x_i, p) p_i(x_i)$$

## ■ Finite-size corrections

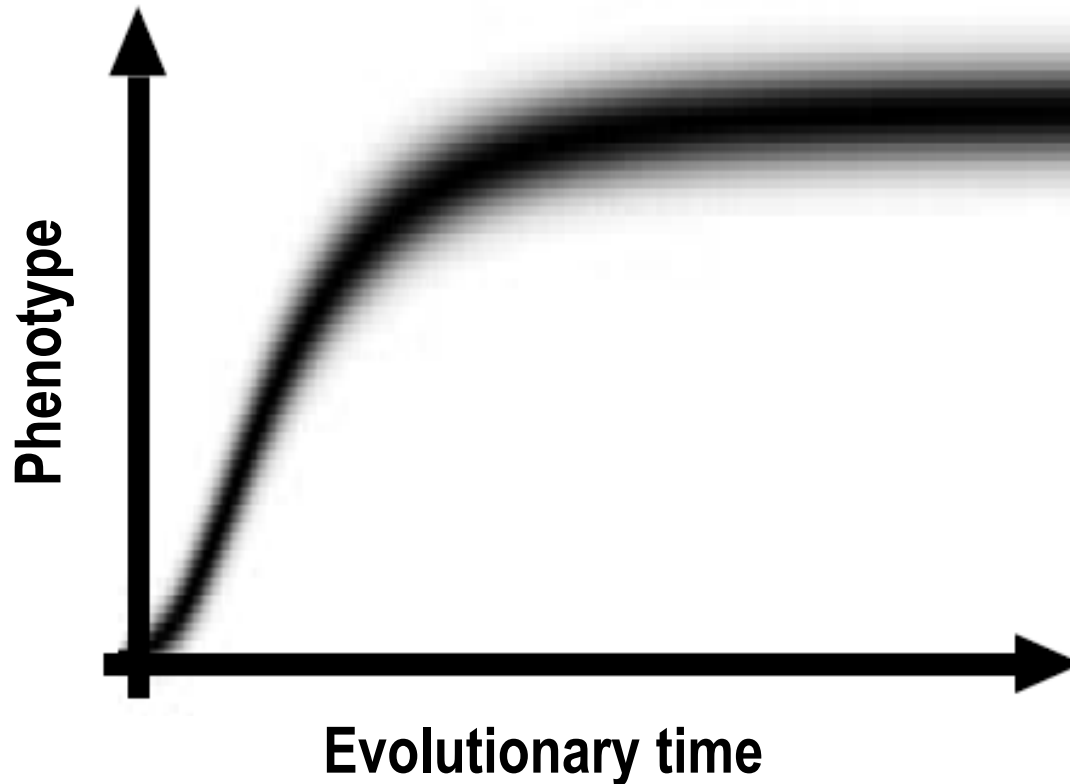




# Reaction-Diffusion Dynamics

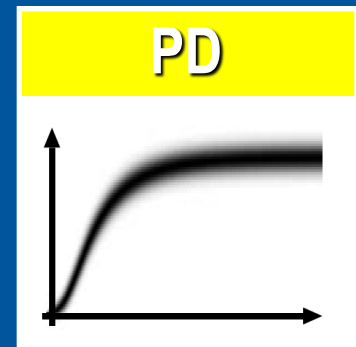
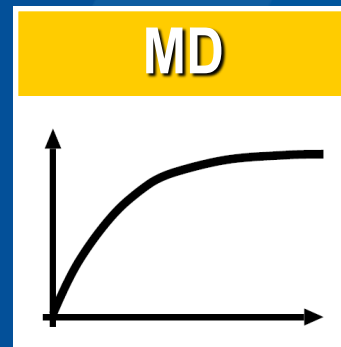
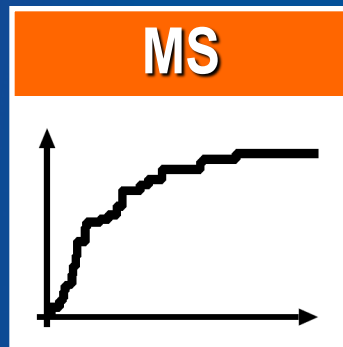
Polymorphic and Deterministic

Kimura (1965)



# Summary of Connections

large population size  
small mutation probability      small mutation variance      fixed standing variance



large population size  
large mutation probability





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# Eco-genetic models

# Eco-Genetic Models

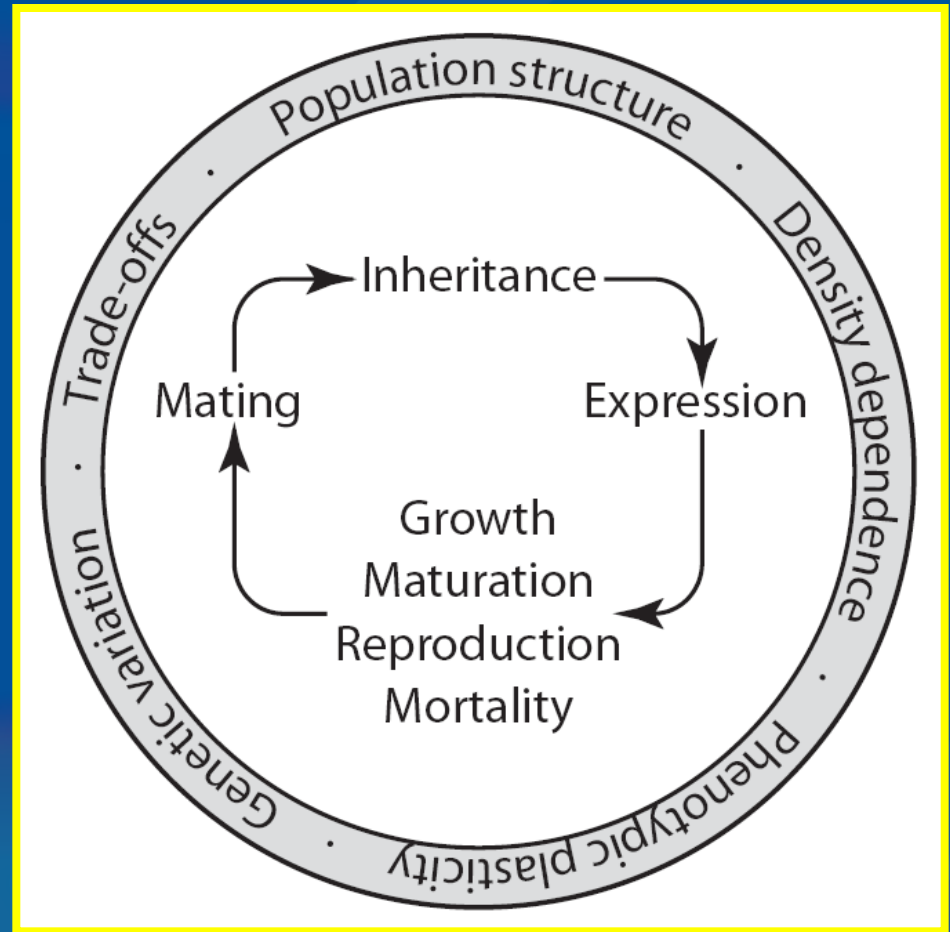
Eco-genetic models are process-based and designed to incorporate

**Ecological detail**

together with

**Genetic detail**

in the context of a stock's life cycle





# Population Structure

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- Individuals differ in physiological states and heritable traits; population structure thus arises from both
- Relevant physiological states may include
  - (1) Age
  - (2) Length
  - (3) Weight
  - (4) Condition
  - (5) Maturity status
  - (6) Sex
  - (7) Stock component



# Phenotypic Plasticity

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- Phenotypes are not determined by genotypes alone, but also depend on environmental conditions
- Systematic effects of the latter kind are known as phenotypic plasticity
- Broader and narrower definitions exist: in a broader sense, learning and growth variation induced by resource availability are considered plastic responses
- The level and kind of phenotypic plasticity can itself evolve, and thus may be fine-tuned by selection



# Genetic Variation

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- Without genetic variation, selection pressures cannot elicit selection responses
- Without a correlation between the phenotypes of parents and their offspring (heritability), the effects of selection cannot be transmitted between generations
- Heritabilities are defined as the ratio between (additive) genetic variance and total phenotypic variance
- Genetic variances (and covariances) are insufficient for predicting selection responses when selection is strong, since genotypic distributions are then usually not normal



# Purposes of Eco-Genetic Models

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- **Evaluate** hypotheses advanced for explaining observed data
- **Understand** and quantify anthropogenic selection pressures
- **Forecast** the direction, speed, and outcome of evolutionary changes
- **Predict** the differential evolutionary vulnerability of species and populations
- **Investigate** the consequences of alternative management scenarios



# Modelling Frameworks: Summary

	(1)	(2)
Game-theoretical models	No	No
Quantitative genetics models	No	Yes
Adaptive dynamics models	Yes	No
Eco-genetic models	Yes	Yes

- (1) Ecological complexity, including both frequency- and density-dependent selection
- (2) Genetic variation, enabling predicting the speed of evolution