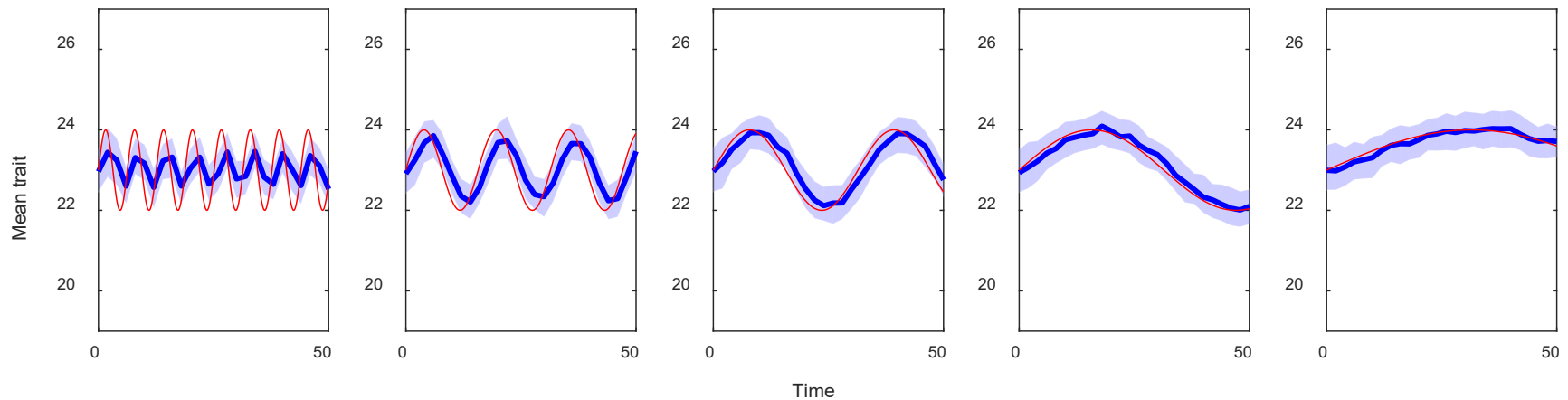


Climate change and scaling up across levels with evolution



John P. DeLong
University of Nebraska – Lincoln

g2p2pop website

Goal: "...predicting how populations and species of vertebrates will respond to a changing climate by integrating knowledge across levels of biological organization."

Four suggestions today:

- Anchor models at points that *must be true*
- Visualize parameters of dynamic models as emergent properties of lower levels – dissect them to access mechanism
- Allow for changes in allele frequencies through evolution
- Try out Gillespie eco-evolutionary models (GEMs) as a tool

LOO

States

Mechanism

Populations

Abundance

Demography

Individuals

Phenotypes

Growth,
development

Cells

Networks

Function

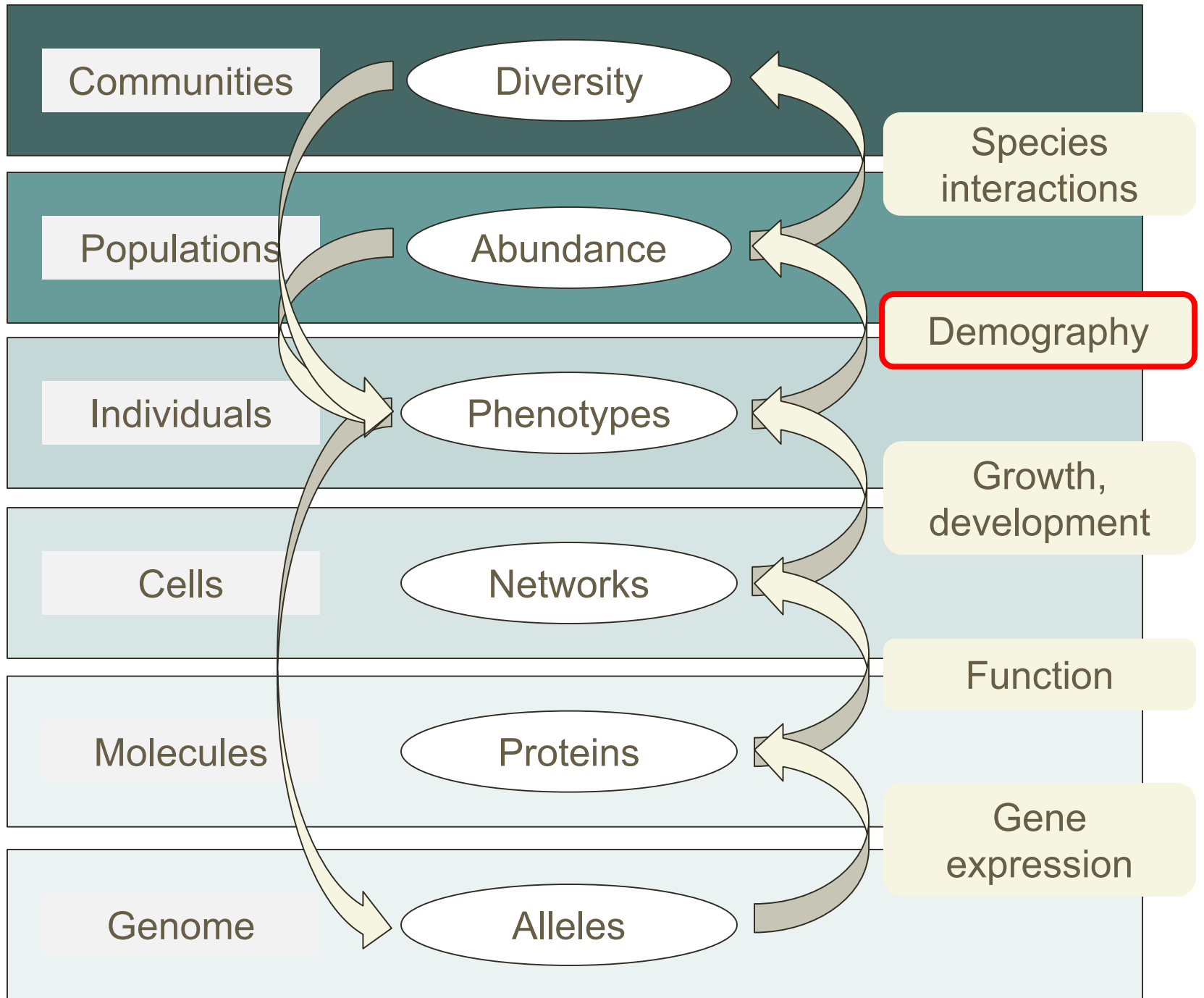
Molecules

Proteins

Gene
expression

Genome

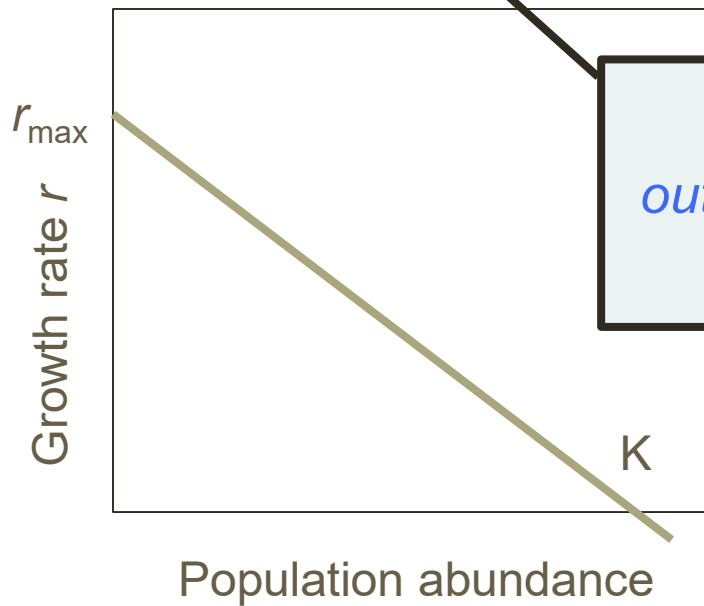
Alleles



Chapter 1. A very simple example.

Logistic model

$$\frac{dR}{dt} = \left[r_{\max} \left(1 - \frac{R}{K} \right) \right] R$$

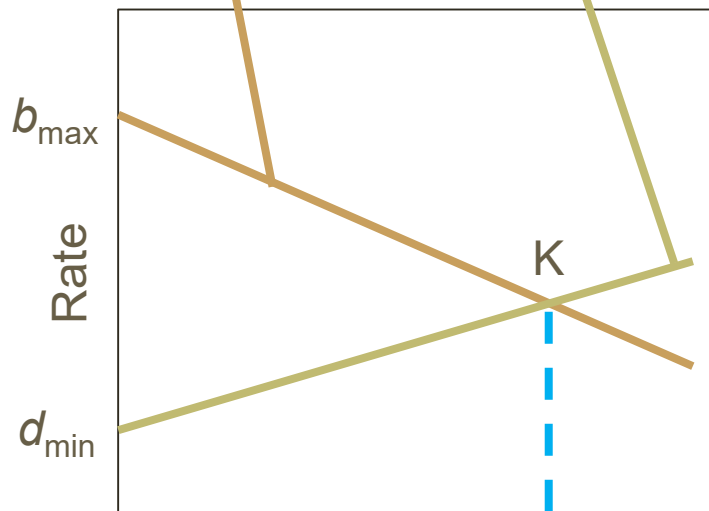


*Parameters are an
outcome of mechanisms
at lower level*

bd logistic

$$\frac{dR}{dt} = [b_{\max}(1 - b_s R) - d_{\min}(1 + d_s R)]R$$

Birth Death



Population abundance

$$r = b_{\max} - d_{\min}$$

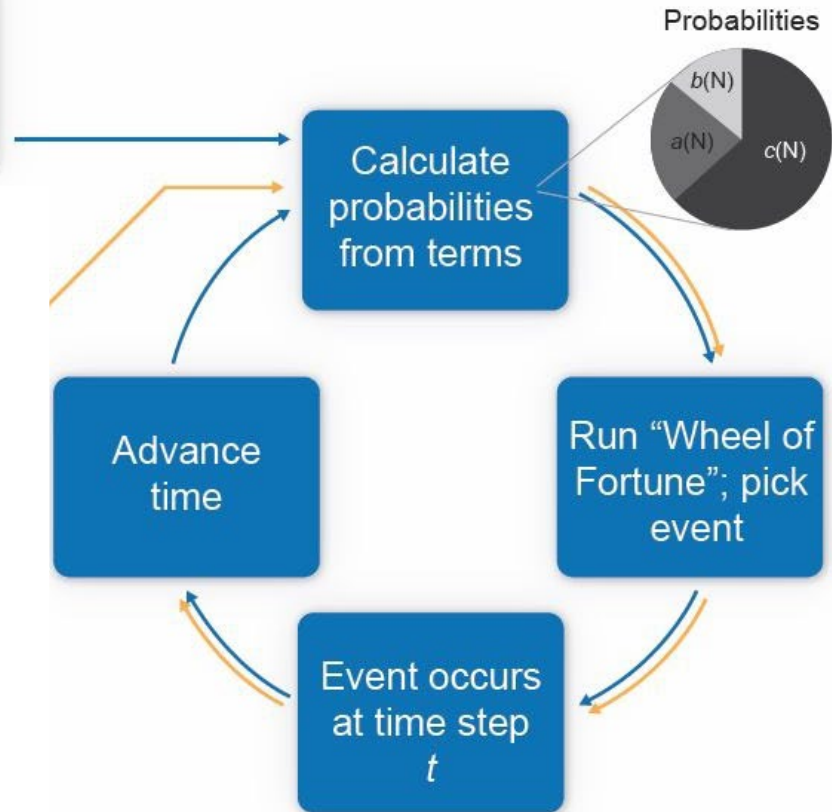
$$K = \frac{b_{\max} - d_{\min}}{b_s + d_s}$$

*Parameters are an
outcome of mechanisms
at lower level*

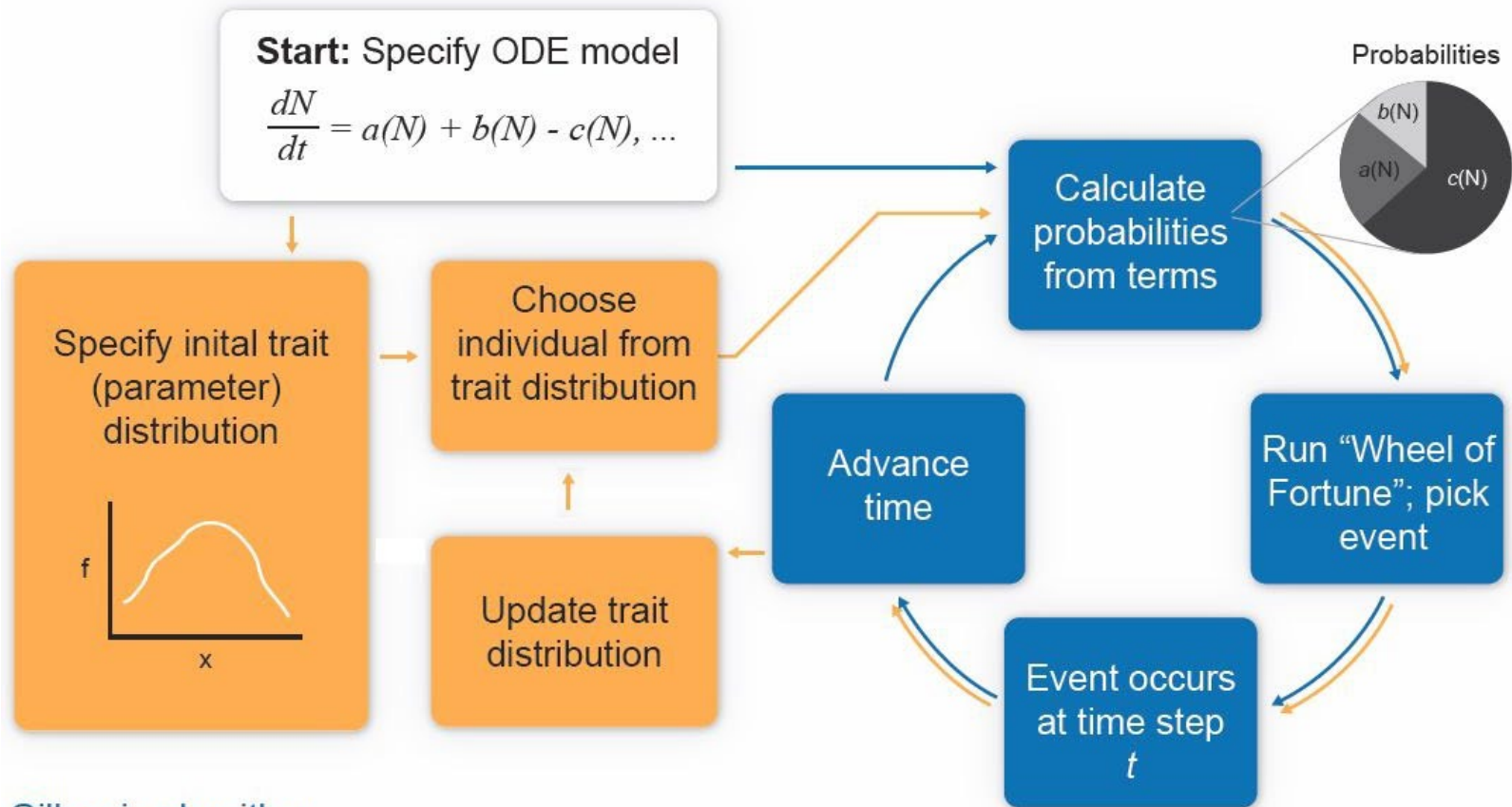
Gillespie algorithm

Start: Specify ODE model

$$\frac{dN}{dt} = a(N) + b(N) - c(N), \dots$$



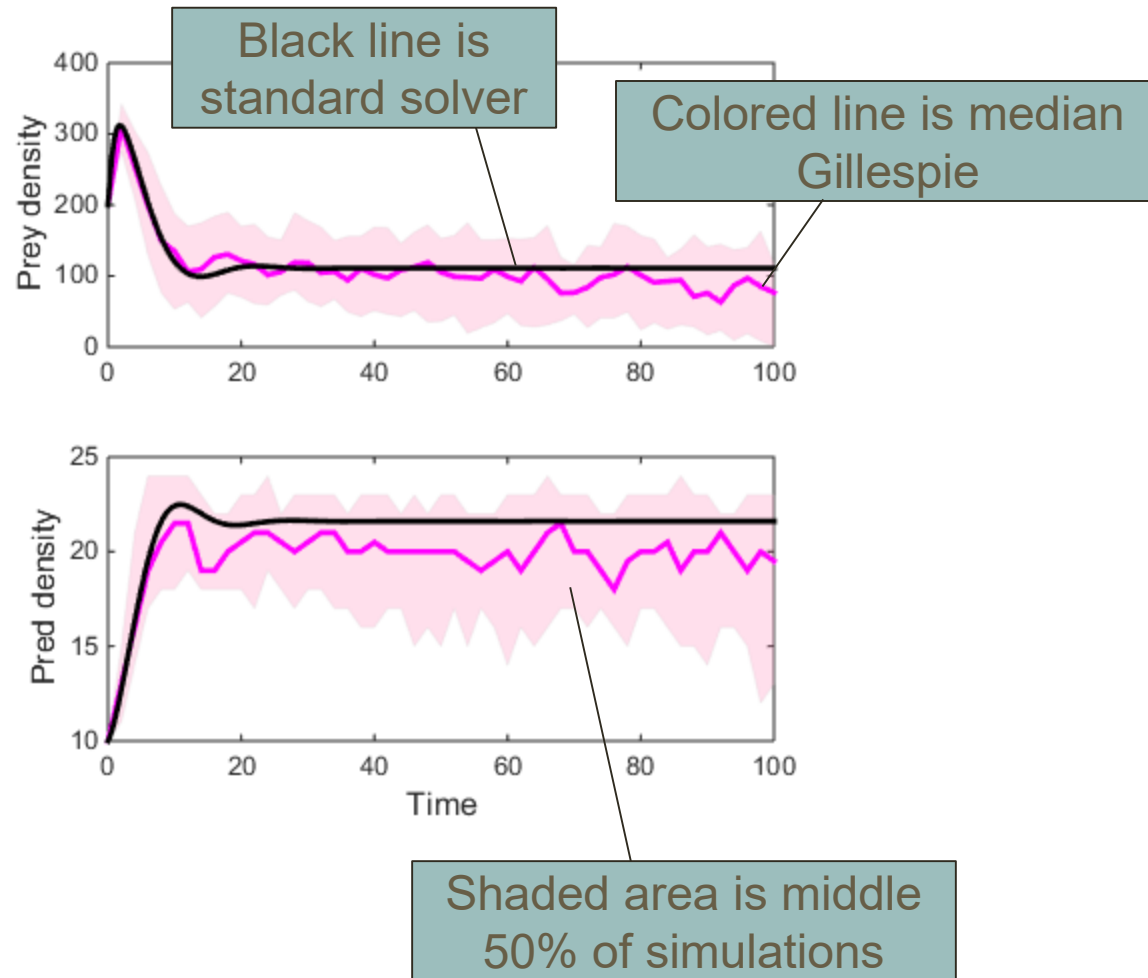
GEMs



Gillespie algorithm

Gillespie eco-evolutionary model

Figure primer

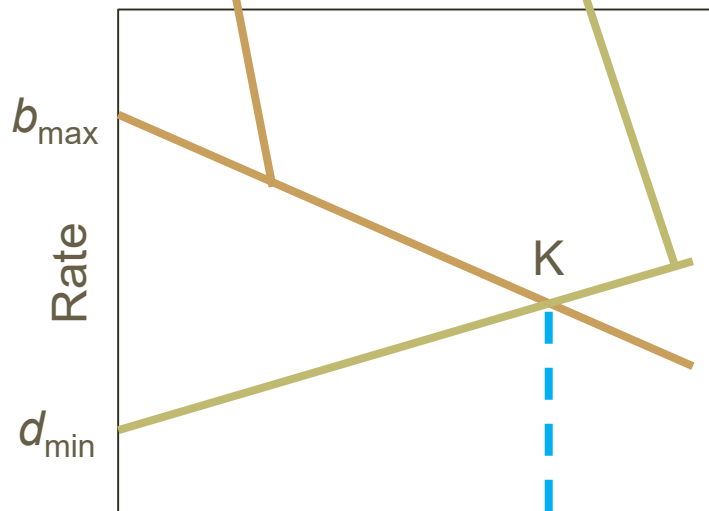


bd logistic

$$\frac{dR}{dt} = [b_{\max}(1 - b_s R) - d_{\min}(1 + d_s R)]R$$

Birth

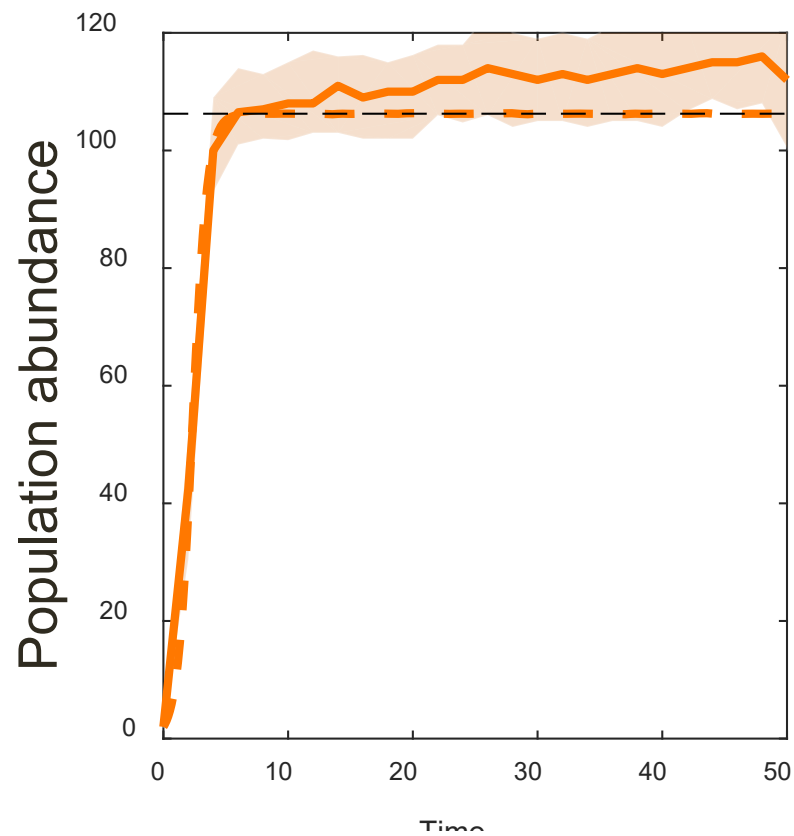
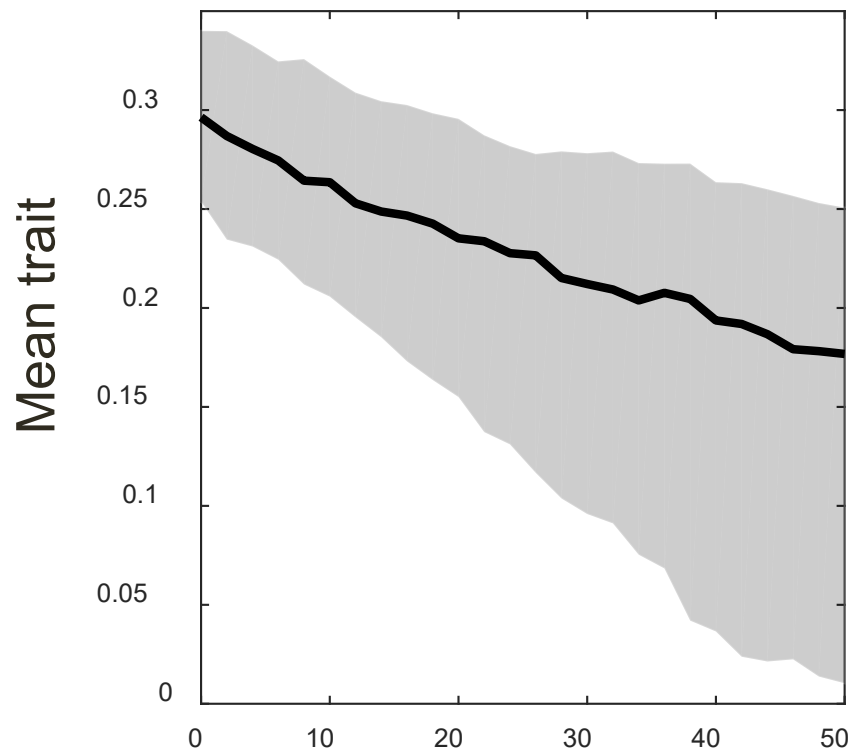
Death



Population abundance

$$r = b_{\max} - d_{\min}$$

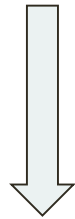
$$K = \frac{b_{\max} - d_{\min}}{b_s + d_s}$$



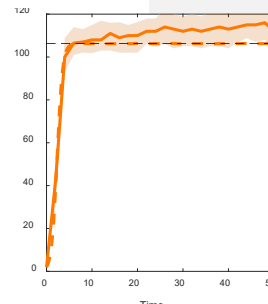
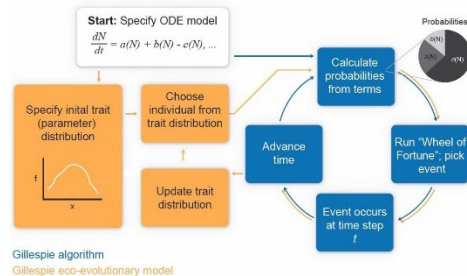
Time

Eco-evolutionary dynamics

$$\frac{dR}{dt} = rR \left(1 - \frac{R}{K}\right)$$



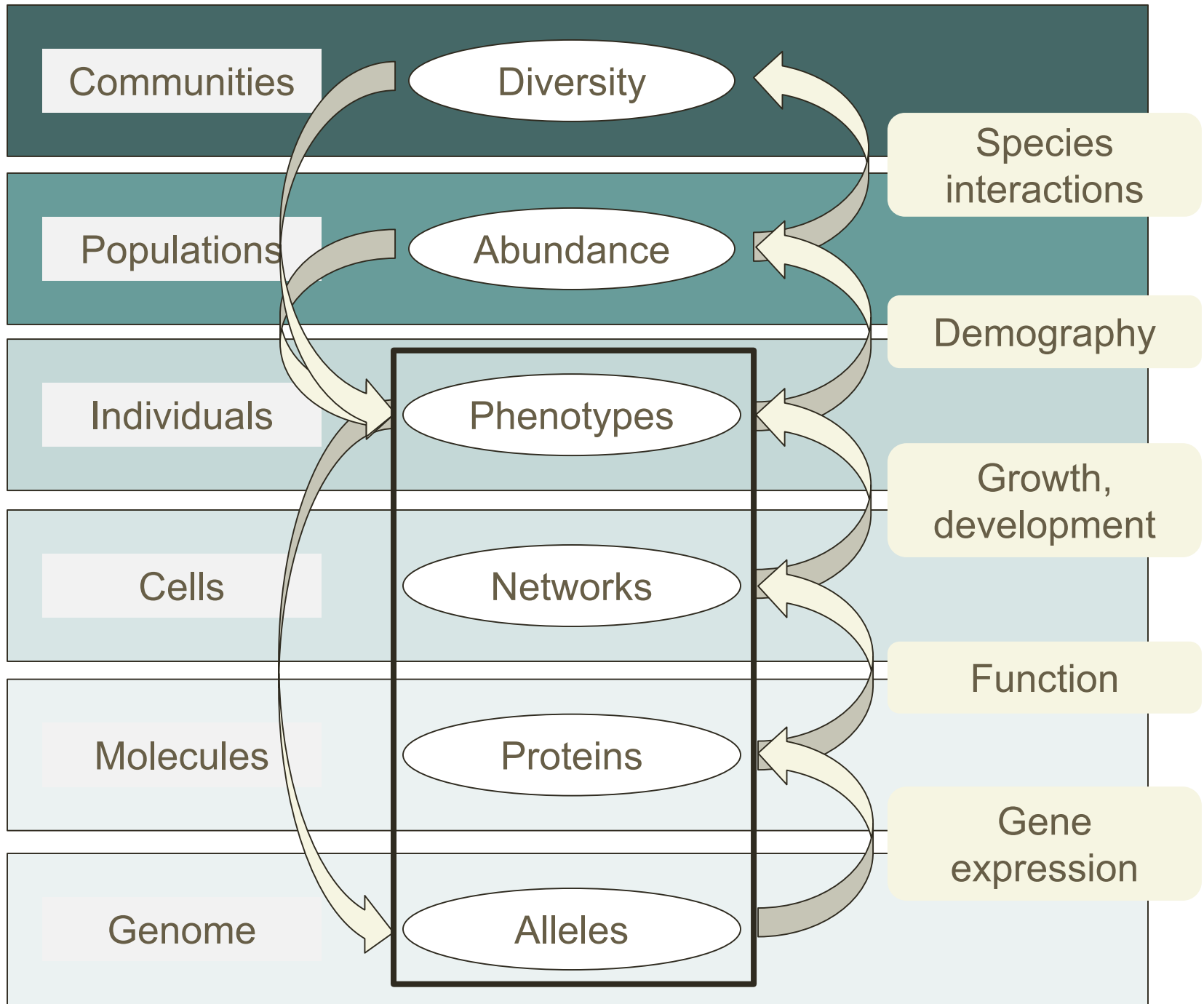
$$[b_{max}(1 - b_s R) - d_{min}(1 + d_s R)]$$



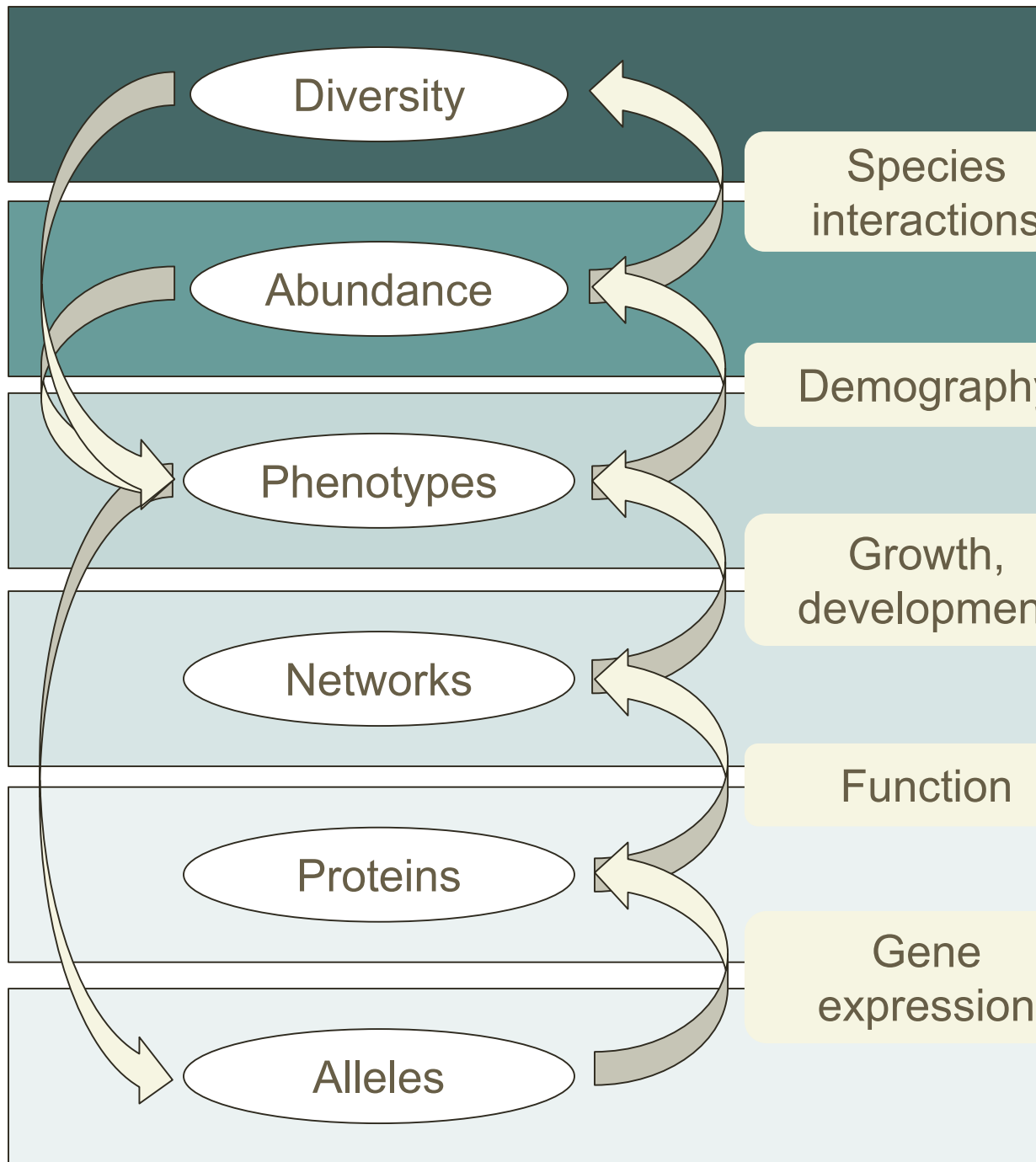
1. Take a population model as a mass-balance constraint

2. Take apart parameters to get at their mechanistic underpinnings

3. Use a GEM to assess real-time evolutionary effects on population outcomes

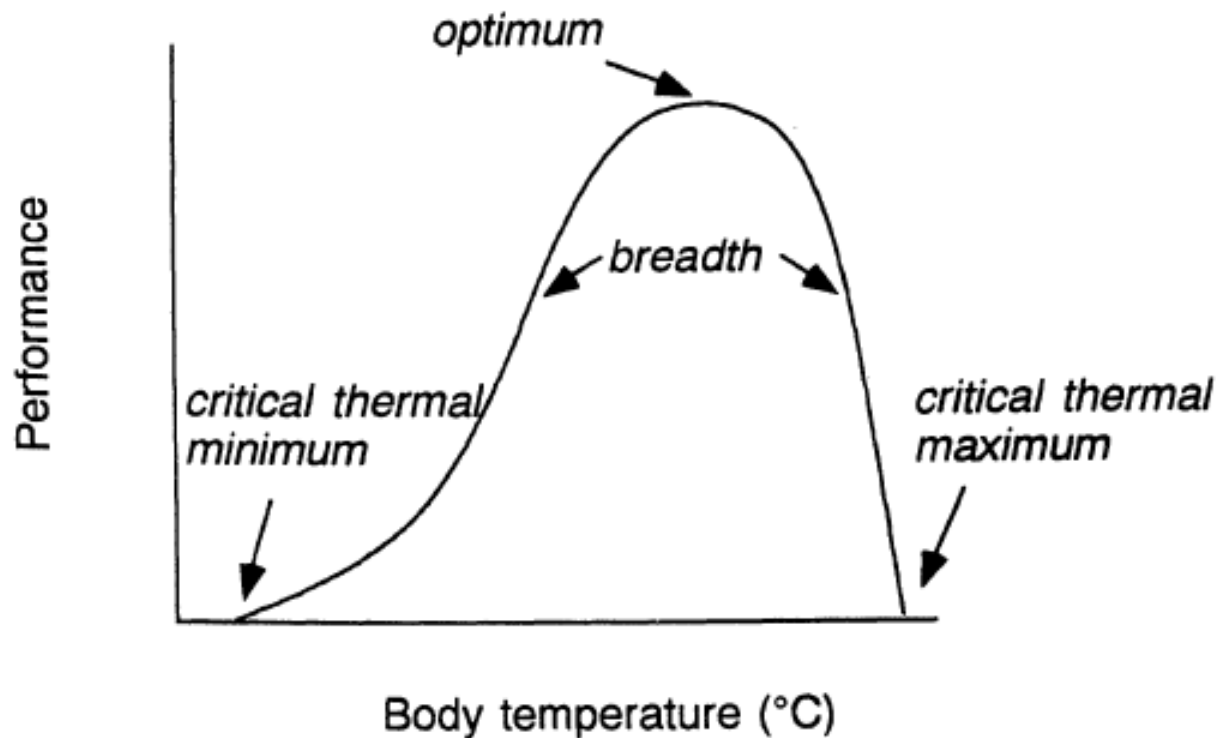


Chapter 2. Evolution of thermal niches.



Temperature

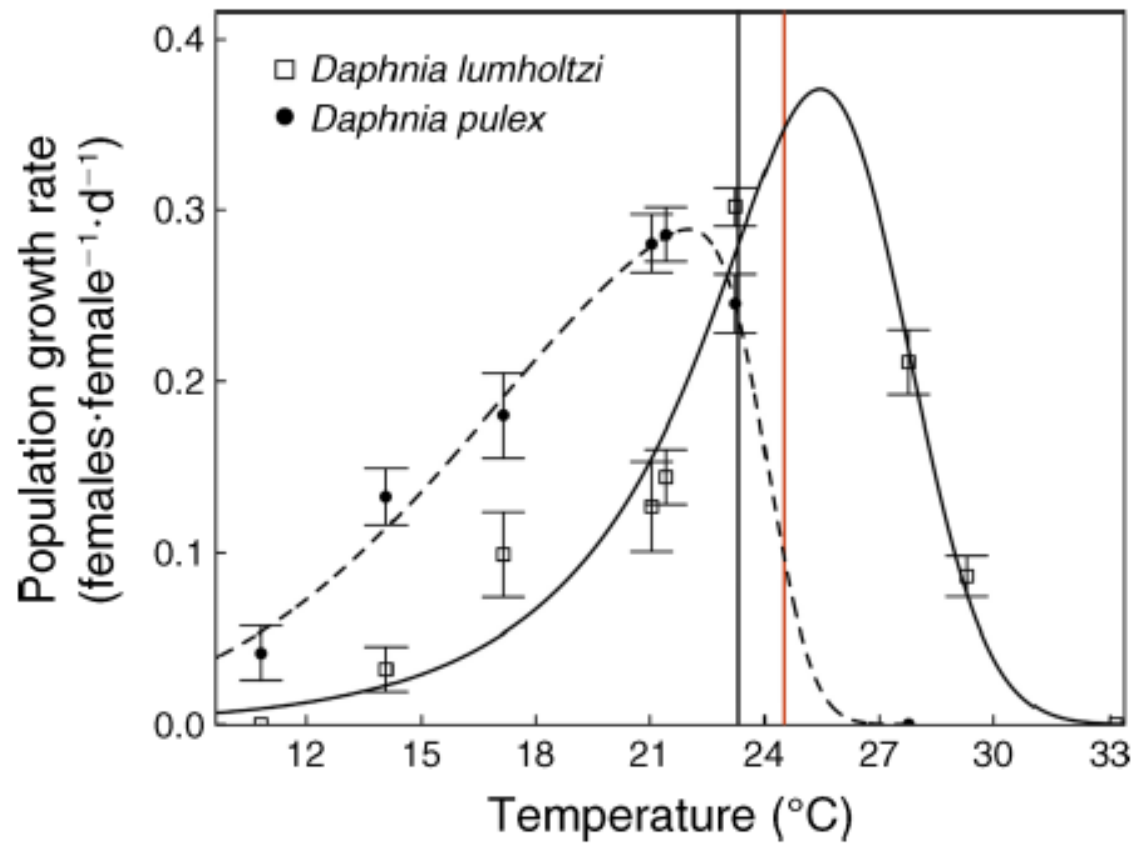
Thermal performance curve (TPC)



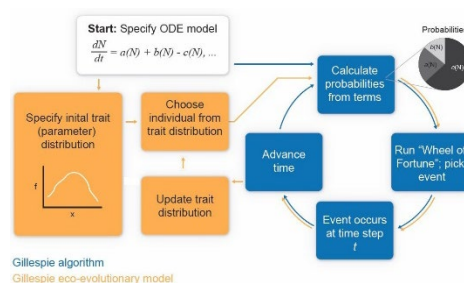
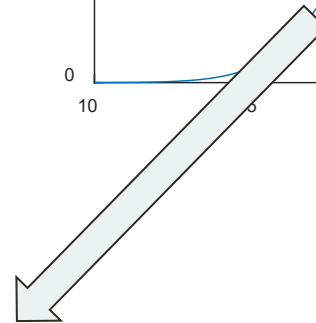
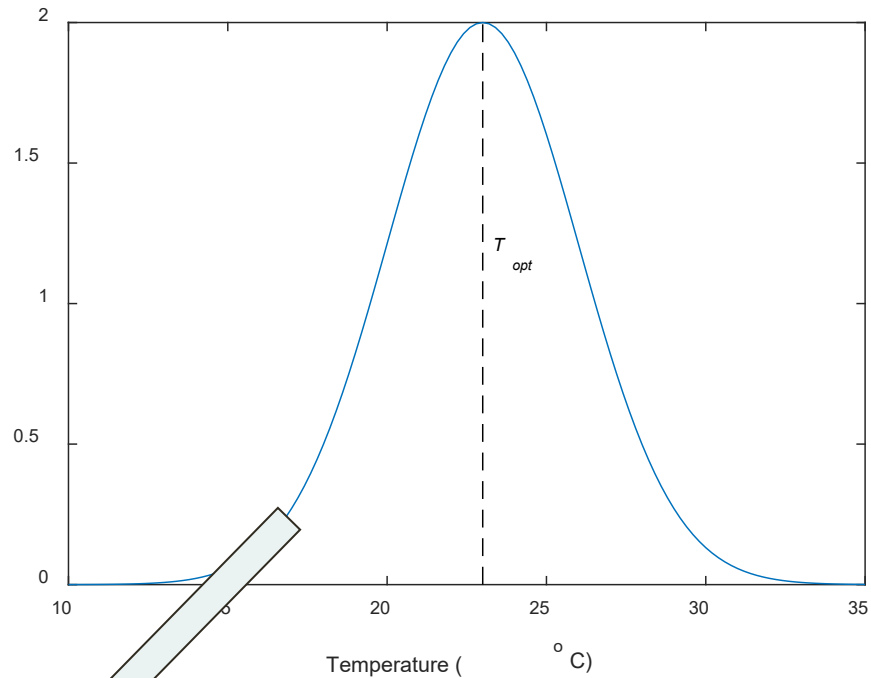
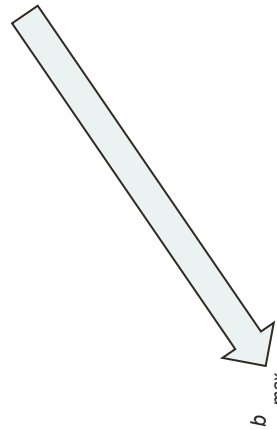
Huey and Kingsolver 1993

Environmental temperature





$$[b_{max}(1 - b_s R) - d_{min}(1 + d_s R)]$$

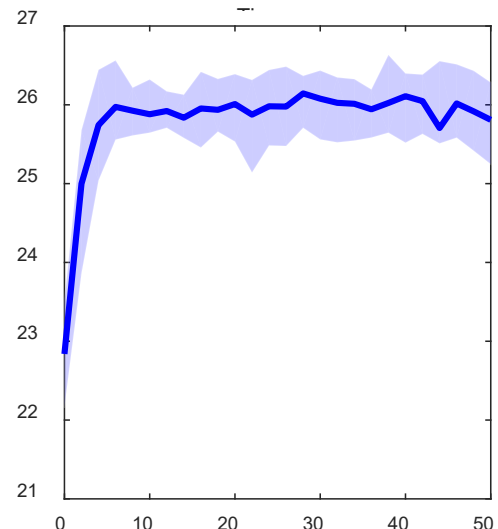
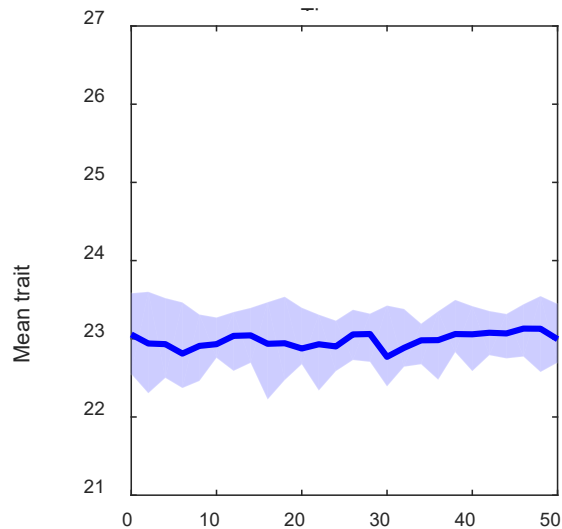
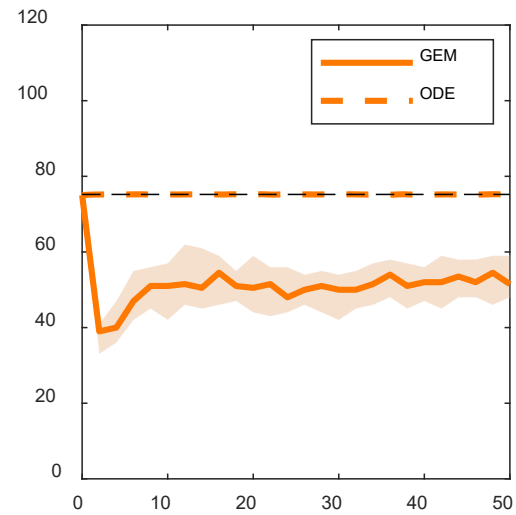
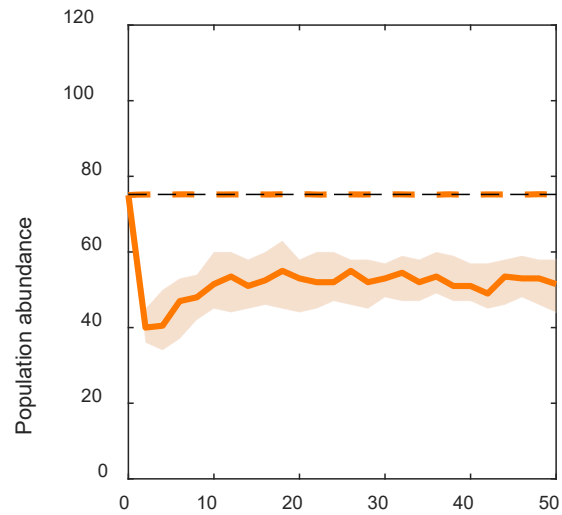


Two things:

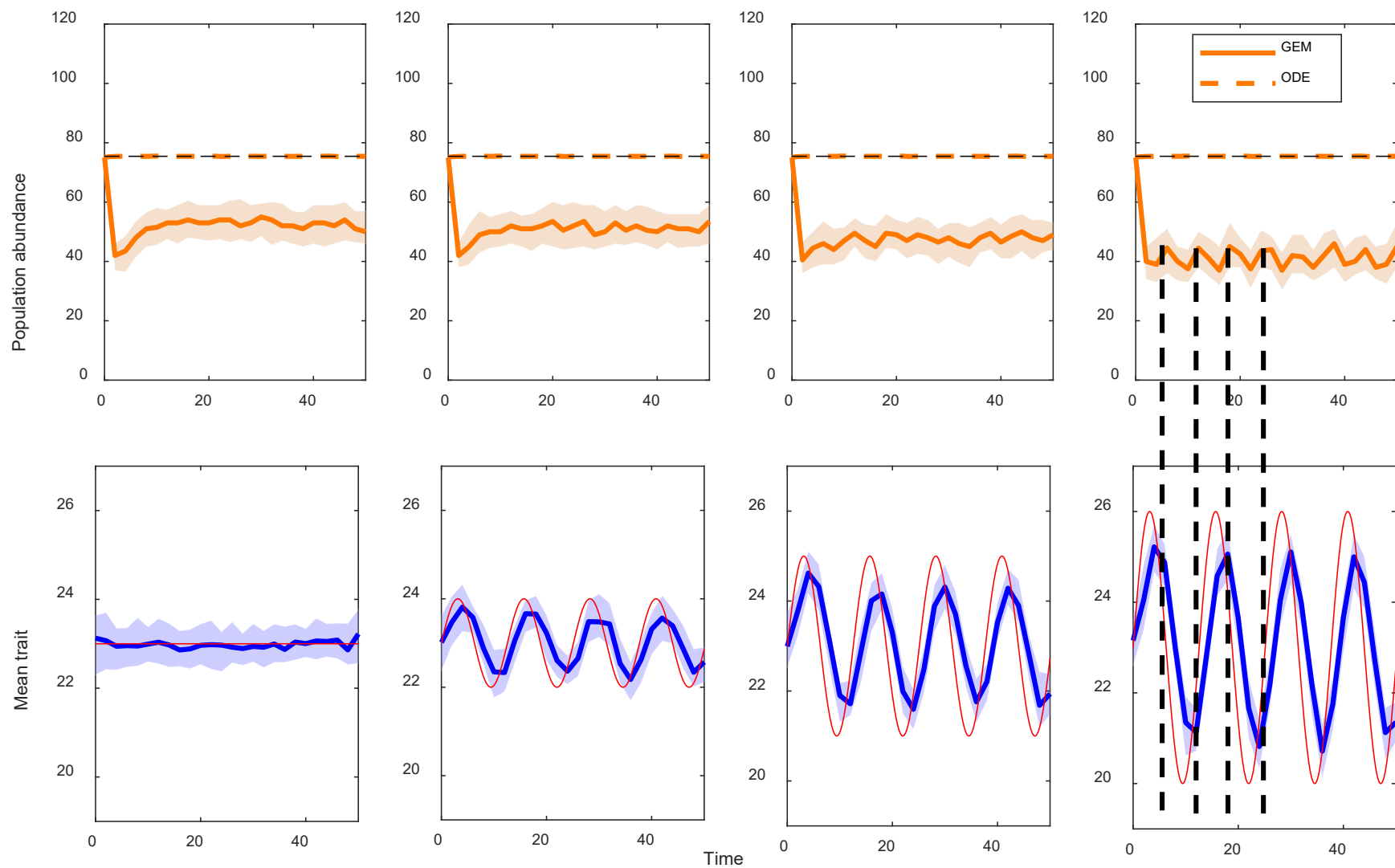
1. Leave temp at 23°C
2. Shift temp to 26°C

Temp = 23°C

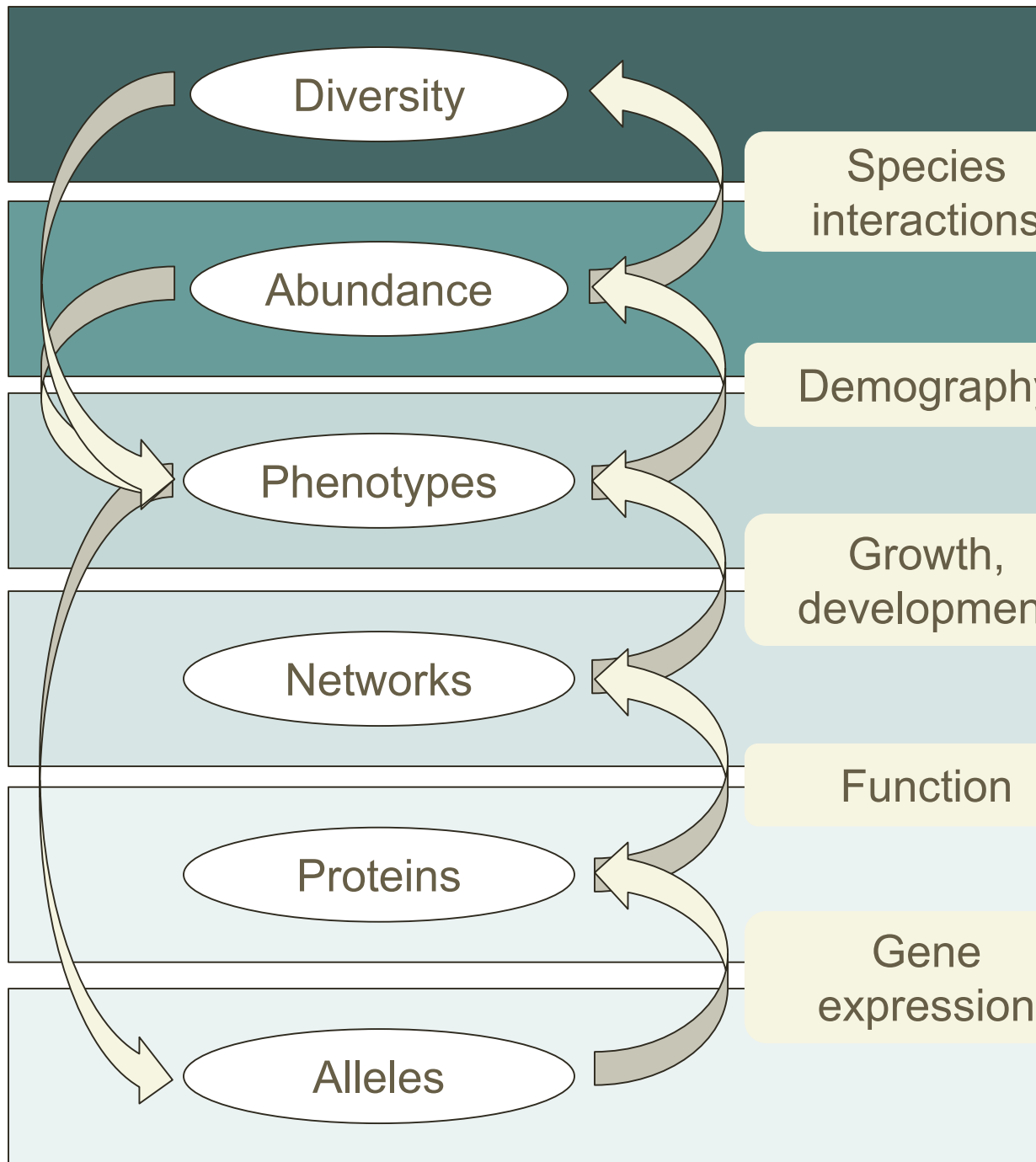
Temp = 26°C



Time

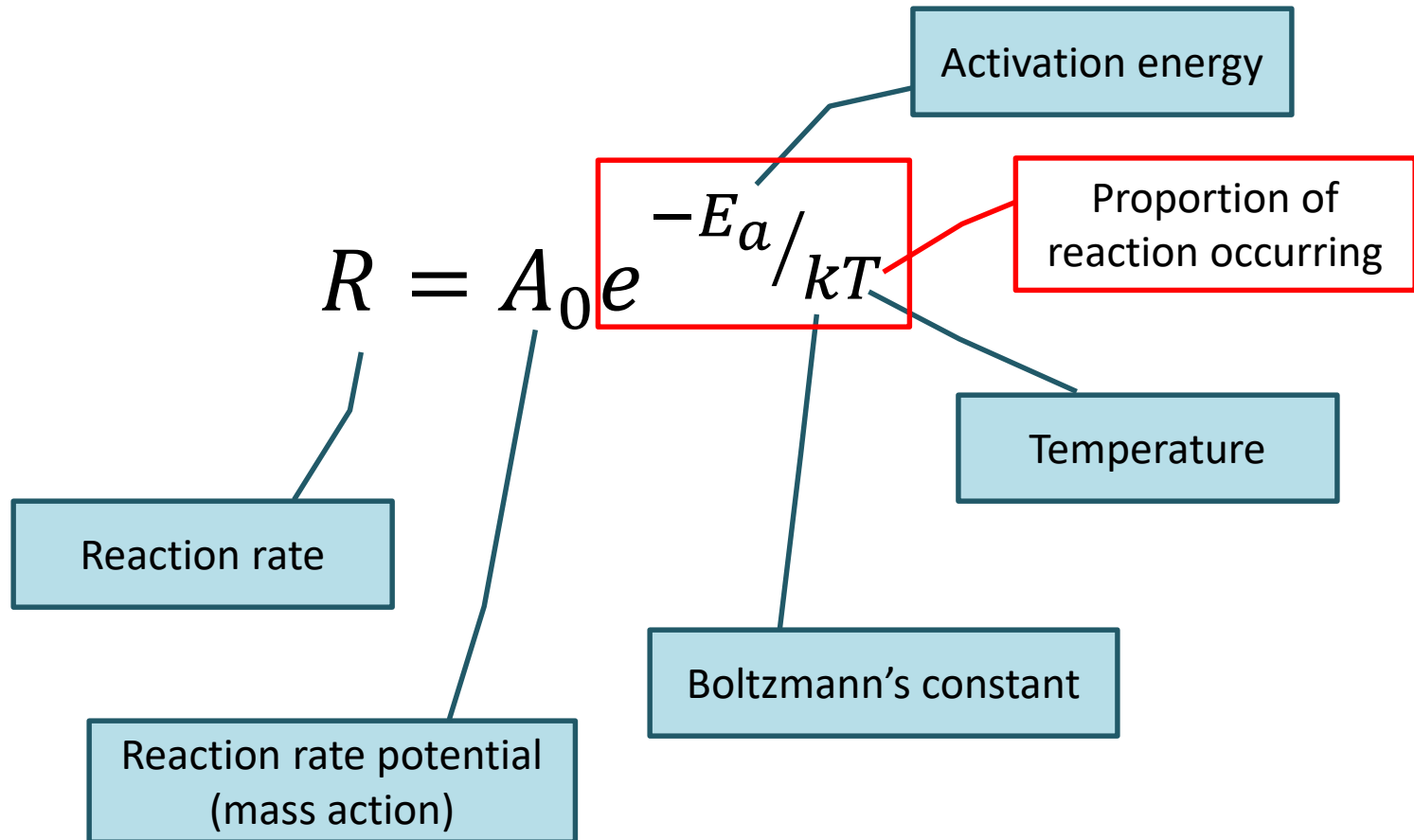


Chapter 3. Mechanisms of temperature dependence.

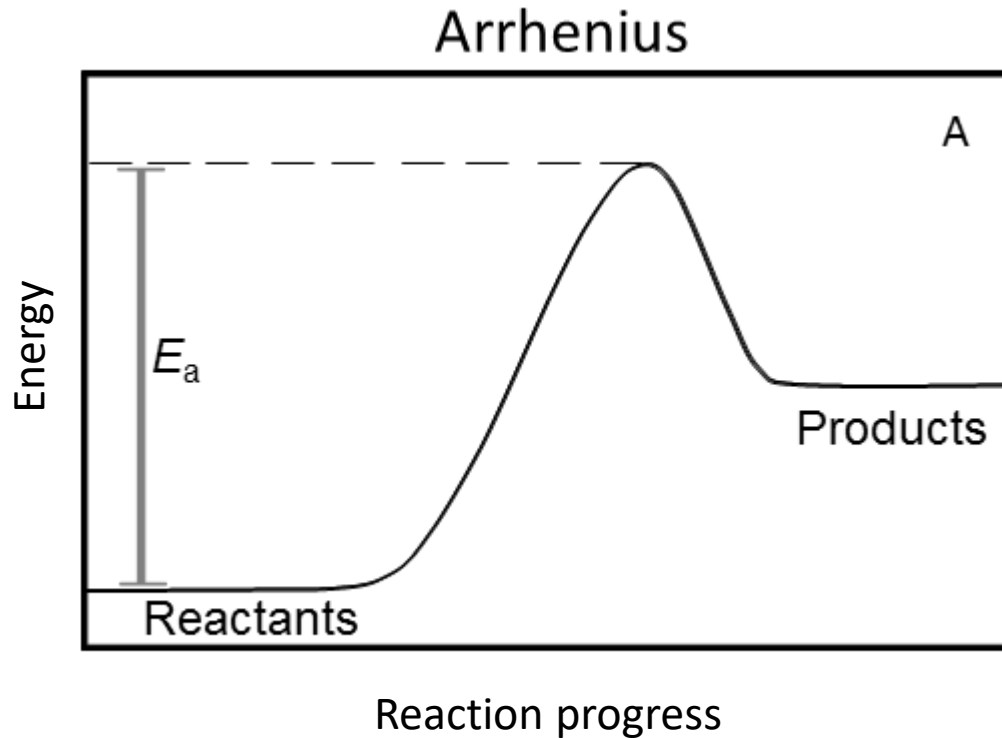


Temperature

Arrhenius equation

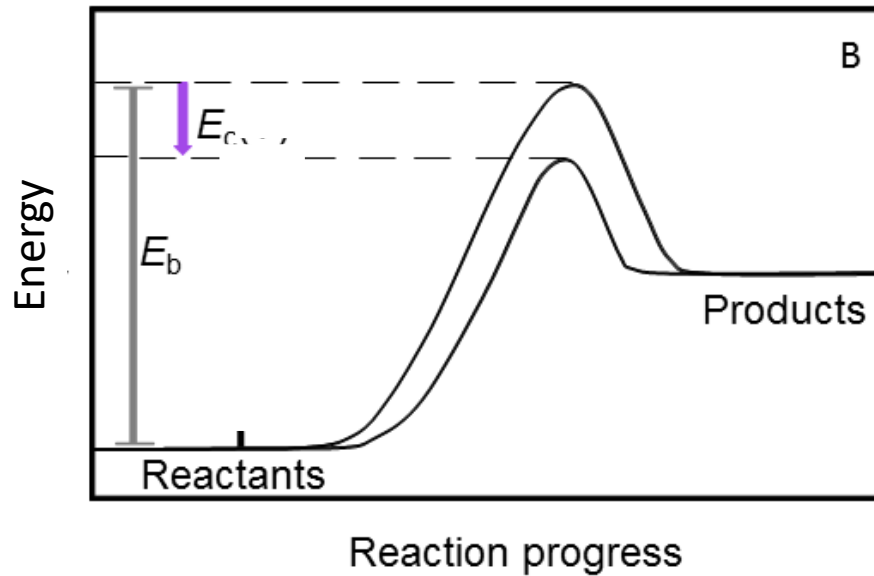


Activation energy



$$R = A_0 e^{-E_a/kT}$$

Enzyme assist

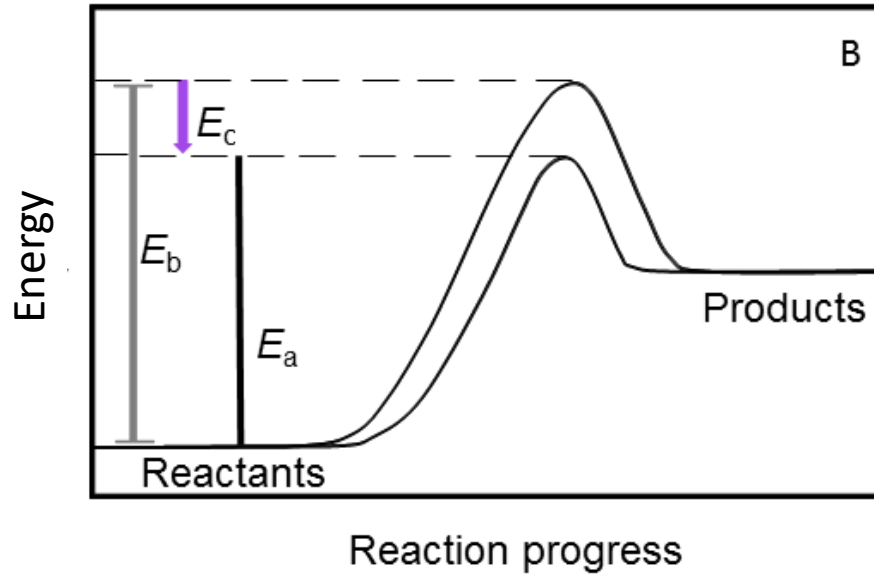


Baseline activation
energy

Contributed energy

$$R = A_0 e^{-(E_b - E_c)/kT}$$

Enzyme assist



Baseline activation
energy

Contributed energy

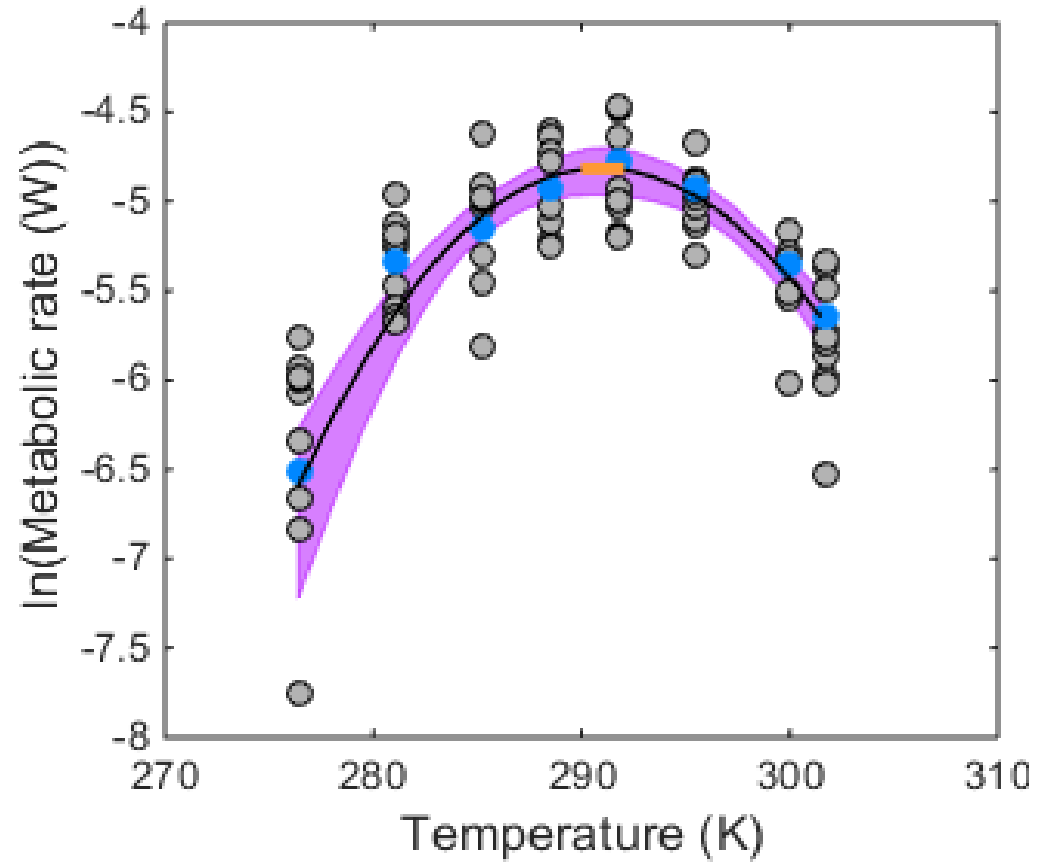
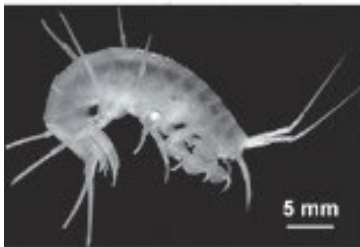
$$R = A_0 e^{-\left(E_b - E_c(T)\right) / kT}$$

Enzyme-assisted Arrhenius (EAAR)

$$R = A_0 e^{-\left(E_b - E_c(T)\right) / kT}$$

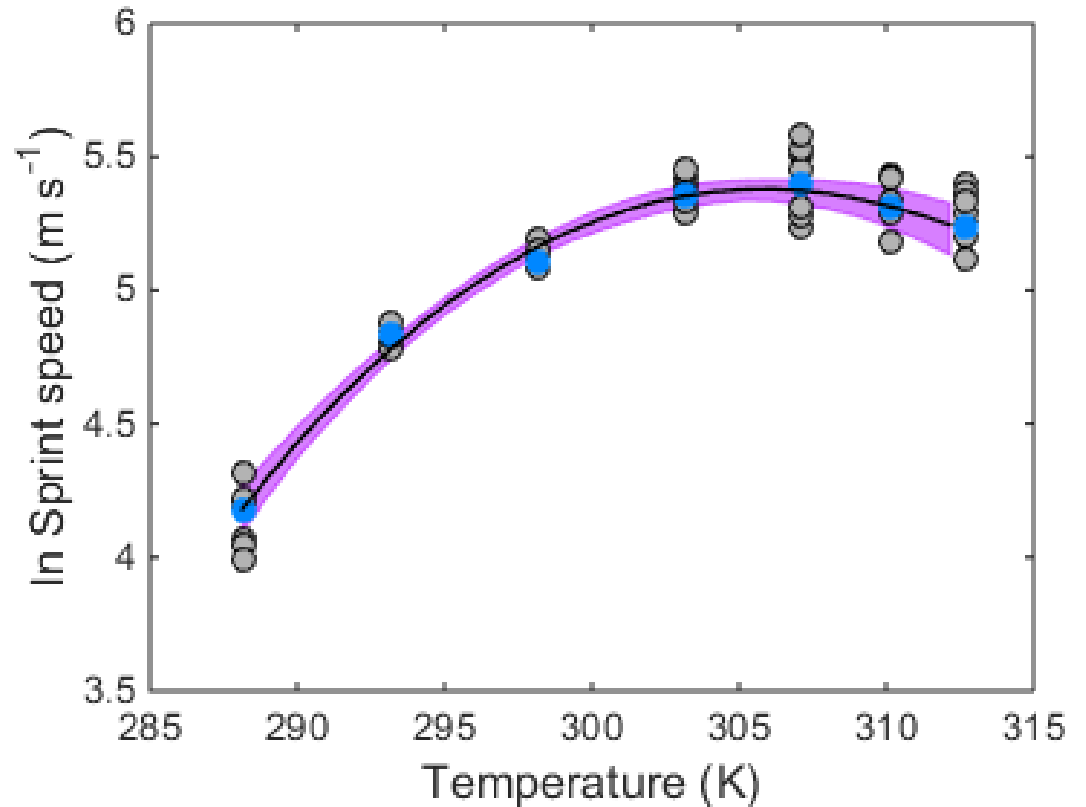
Metabolism

Amphipod
(*Niphargus virei*)



Metabolism-fueled movement

Common house gecko
Hemidactylus frenatus



Huey et al 1989

Climate adaptation experiment

Paramecium caudatum



Drosophila melanogaster



20°C



36°C



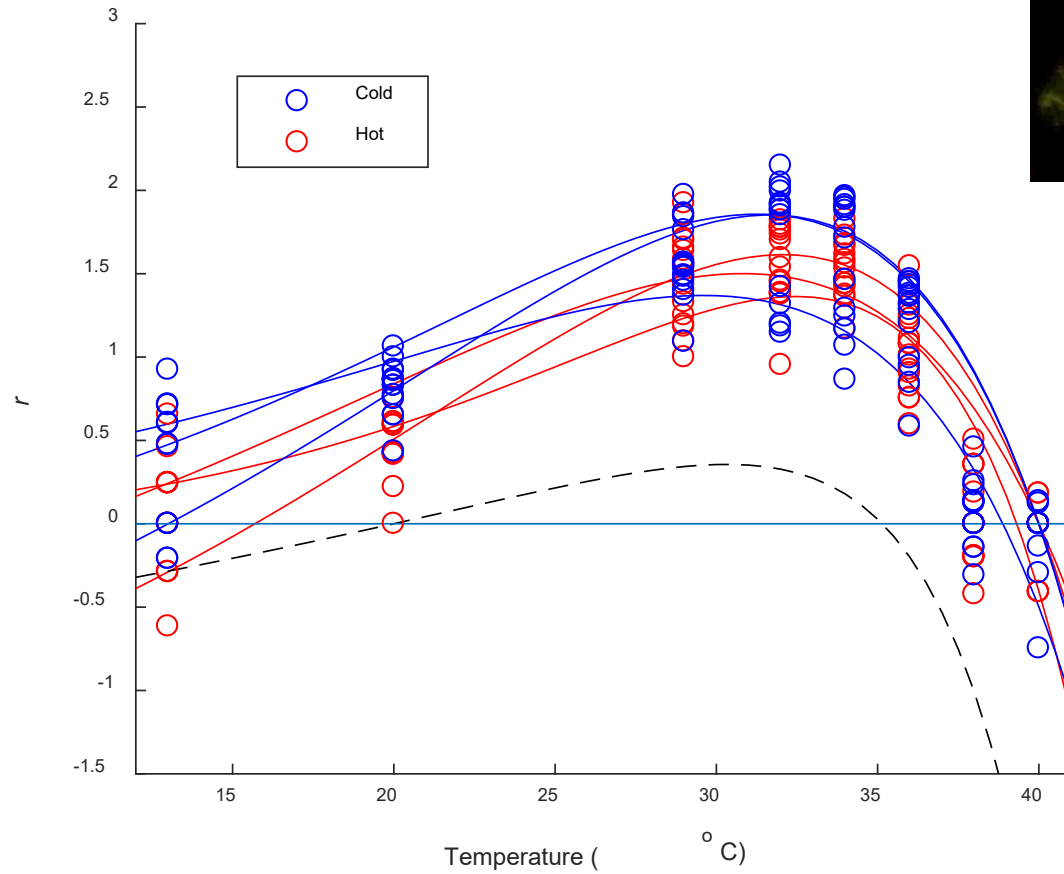
One
ancestral
population



5
different
lakes



After ~350
generations



Communities

Species
interactions

Populations

$$b_{max} = f(T_{opt})$$

Demography

Individuals

$$T_{opt} = f'(E_b, E_c)$$

Growth,
development

Cells

$$E_c = f''(\Delta G)$$
$$\Delta G = f'''(\Delta C_p, \Delta H)$$

Function

Molecules

$$\Delta H = f''''(proteins)$$
$$\Delta C_p = f'''''(proteins)$$

Gene
expression

Genome

$$proteins = f''''''(alleles)$$

Four suggestions today:

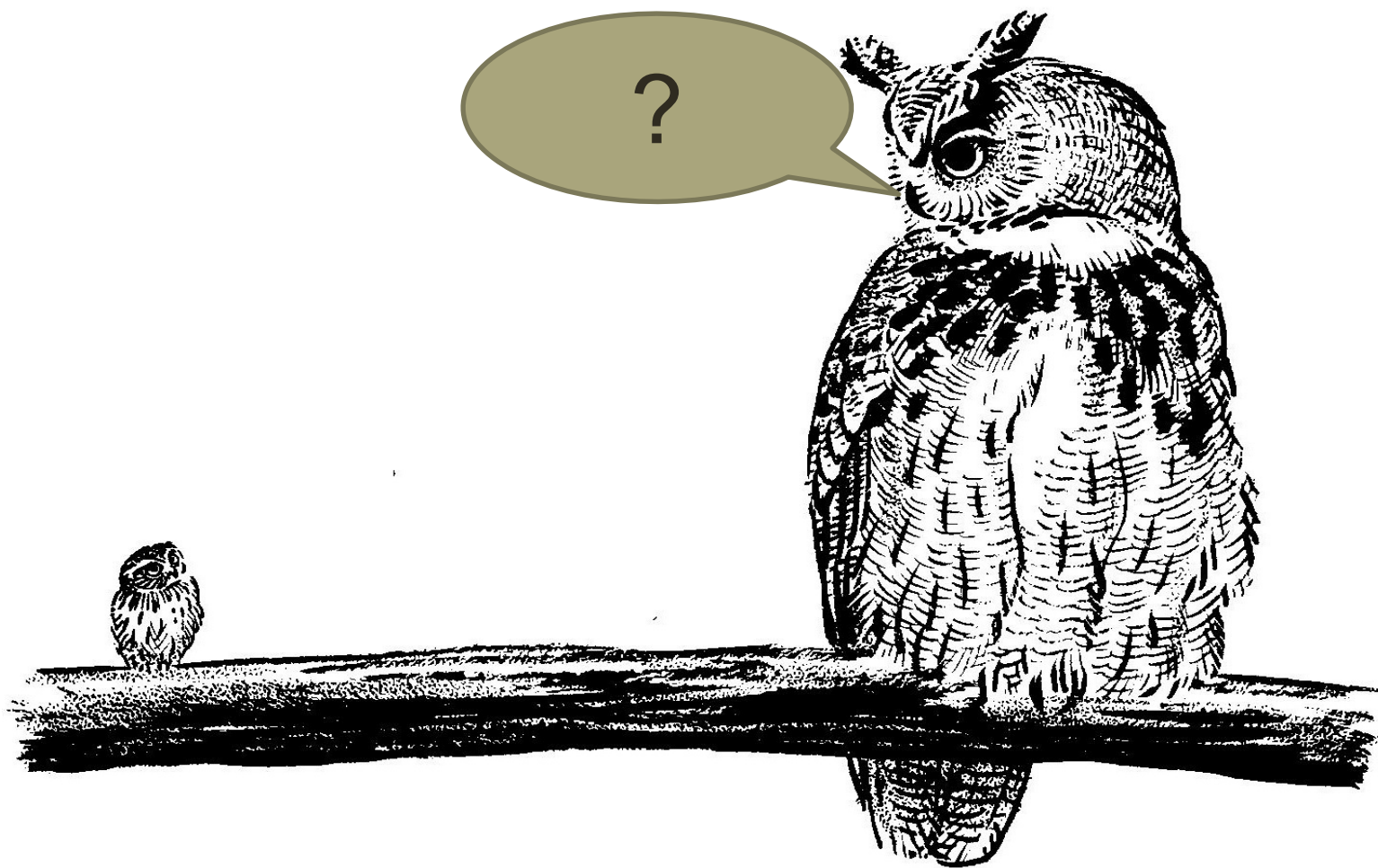
- Anchor models at points that must be true
 - *e.g., population*
- Visualize parameters of dynamic models as emergent properties of lower levels
 - *e.g., r is $b - d$, b is $f(t_{opt})$*
- Allow for changes in allele frequencies through evolution
 - *e.g., climate adaptation changes abundance*
- Try out Gillespie eco-evolutionary models (GEMs) as a tool

Acknowledgements

JAMES S.
MCDONNELL
FOUNDATION

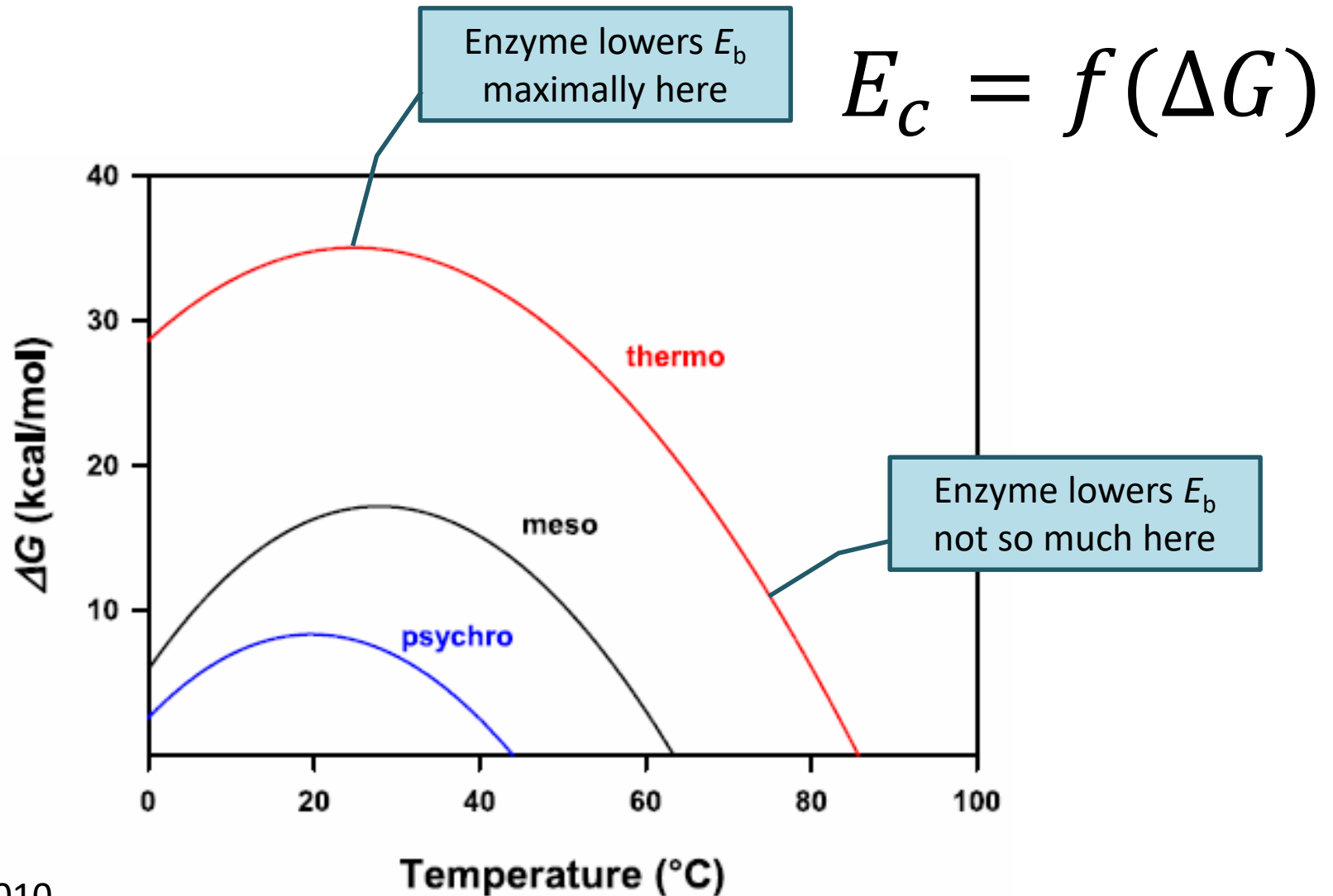


Jean-Philippe Gibert
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Kyle Coblenz
Lyndsie Wszola
Jonathan Belmaker
Miranda Salsbery
Kristi Montooth
Ben Reid
Abby Neyer
Gwen Bachman
Clay Cressler

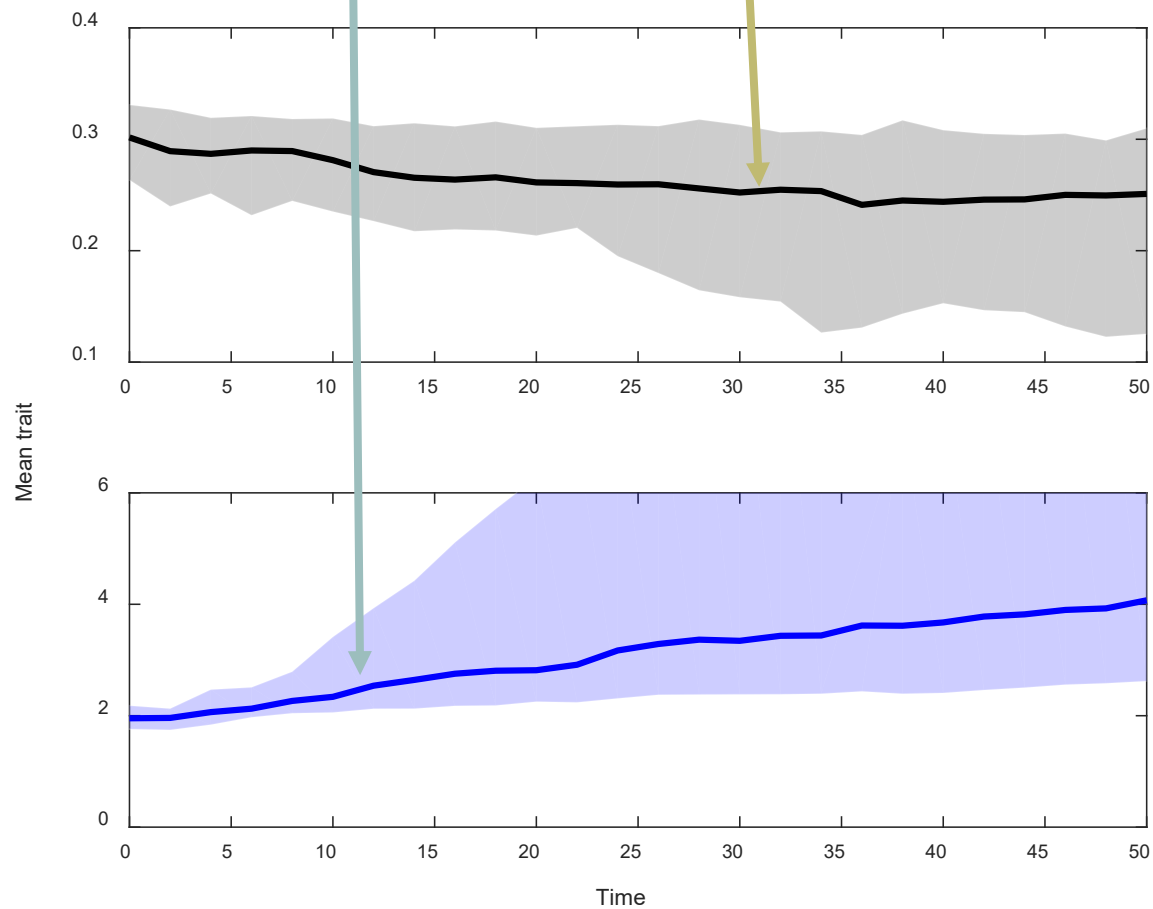


Chapter 4. Scaling up TPCs, empirically and mathematically.

E_c related to protein stability



$$\frac{dR}{dt} = [b_{max}(1 - b_s R) - d_{min}(1 + d_s R)]R$$



Translating Gibbs-Helmholtz

$$\Delta G = \Delta H - T\Delta S$$

These are functions
not constants

Becktel and Shellman 1988

$$\Delta G = \Delta H^0 \left(1 - \frac{T}{T_m} \right) + \Delta C_p \left(T - T_m - T \ln \frac{T}{T_m} \right)$$

‘Robustness’



‘Thermal sensitivity’



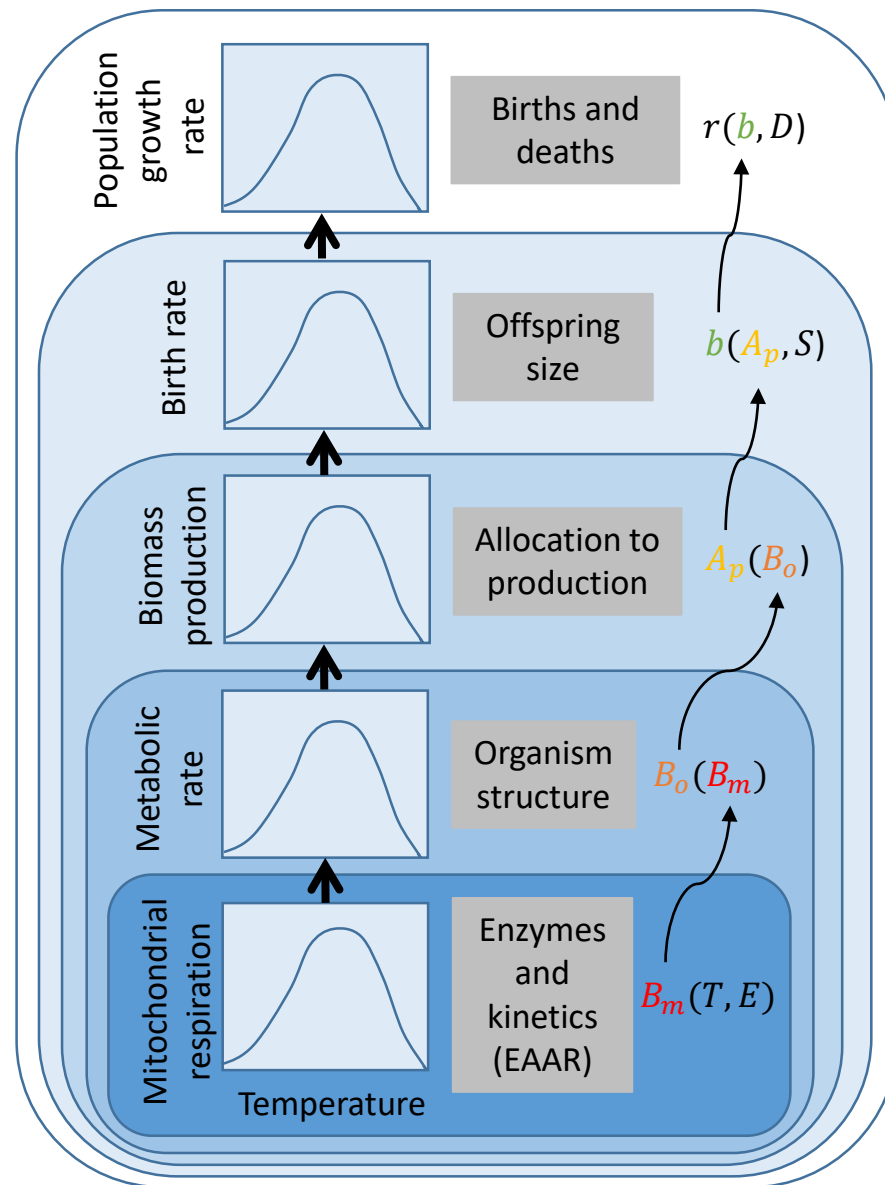
$$\Delta G = \Delta H^0 \left(1 - \frac{T}{T_m} \right) + \Delta C_p \left(T - T_m - T \ln \frac{T}{T_m} \right)$$

$$E_c = E_L \frac{\Delta G}{\Delta G_{max}}$$

$$E_{\Delta H} = E_L \frac{\Delta H}{\Delta G_{max}}$$

$$E_{\Delta C_p} = E_L \frac{\Delta C_p}{\Delta G_{max}}$$

$$E_c = E_{\Delta H} \left(1 - \frac{T}{T_m} \right) + E_{\Delta C_p} \left(T - T_m - T \ln \frac{T}{T_m} \right)$$



Arrhenius factor

