



*The Abdus Salam
International Centre for Theoretical Physics*



2022-1

Workshop on Theoretical Ecology and Global Change

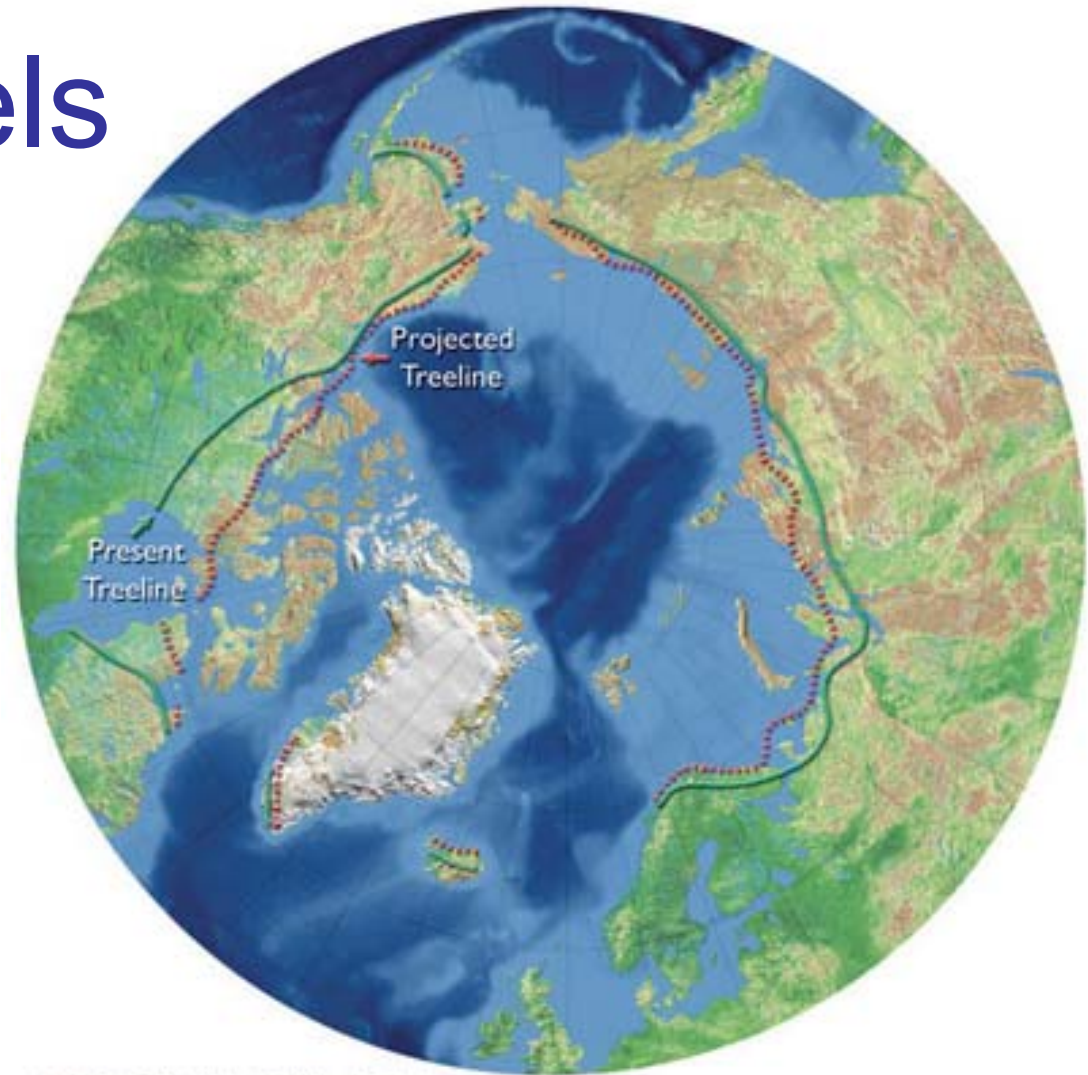
2 - 18 March 2009

Range Models

Lauren Buckley
*The University of North Carolina at Chapel Hill
USA*

Range models

1. Theory



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Lauren Buckley

buckley@bio.unc.edu

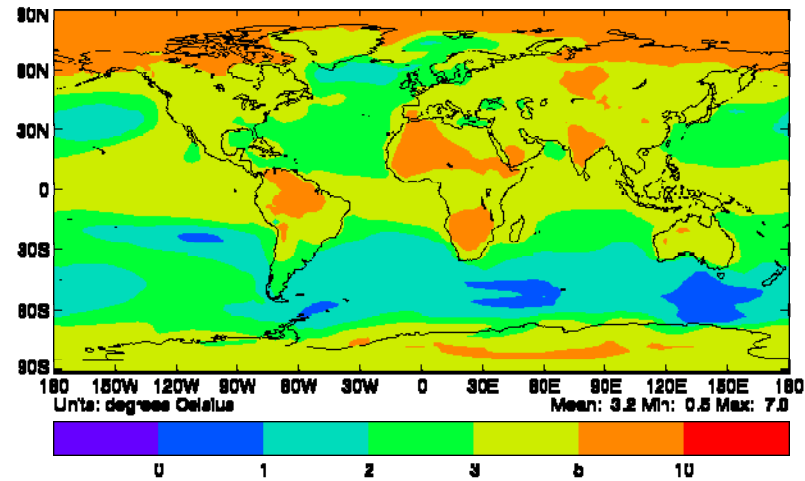
<http://www.bio.unc.edu/Faculty/Buckley/Lab/>



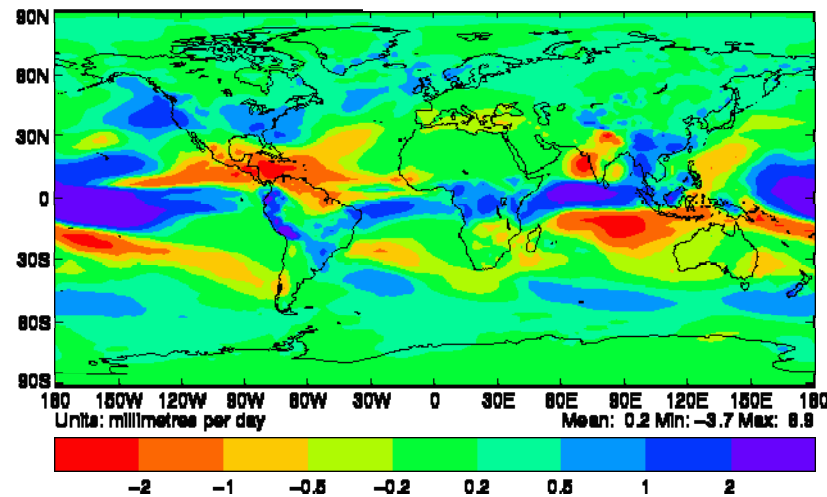
THE UNIVERSITY
of NORTH CAROLINA
at CHAPEL HILL

Climate-induced range shifts and extinctions?

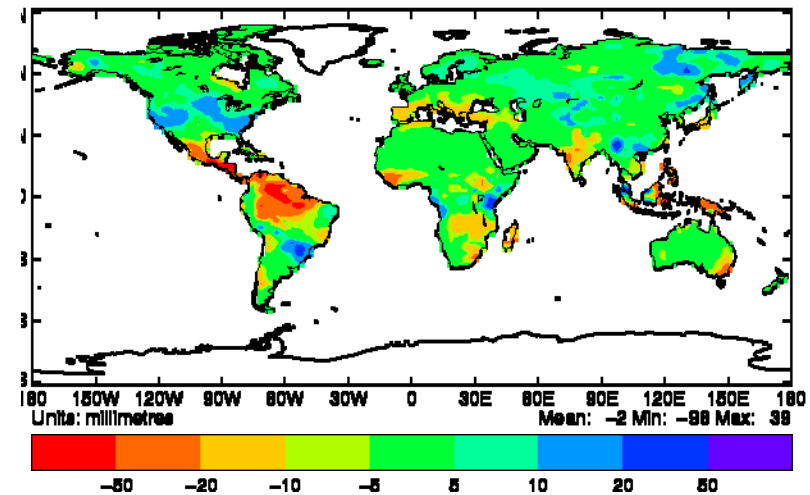
Temperature (°C)



Precipitation (mm day⁻¹)



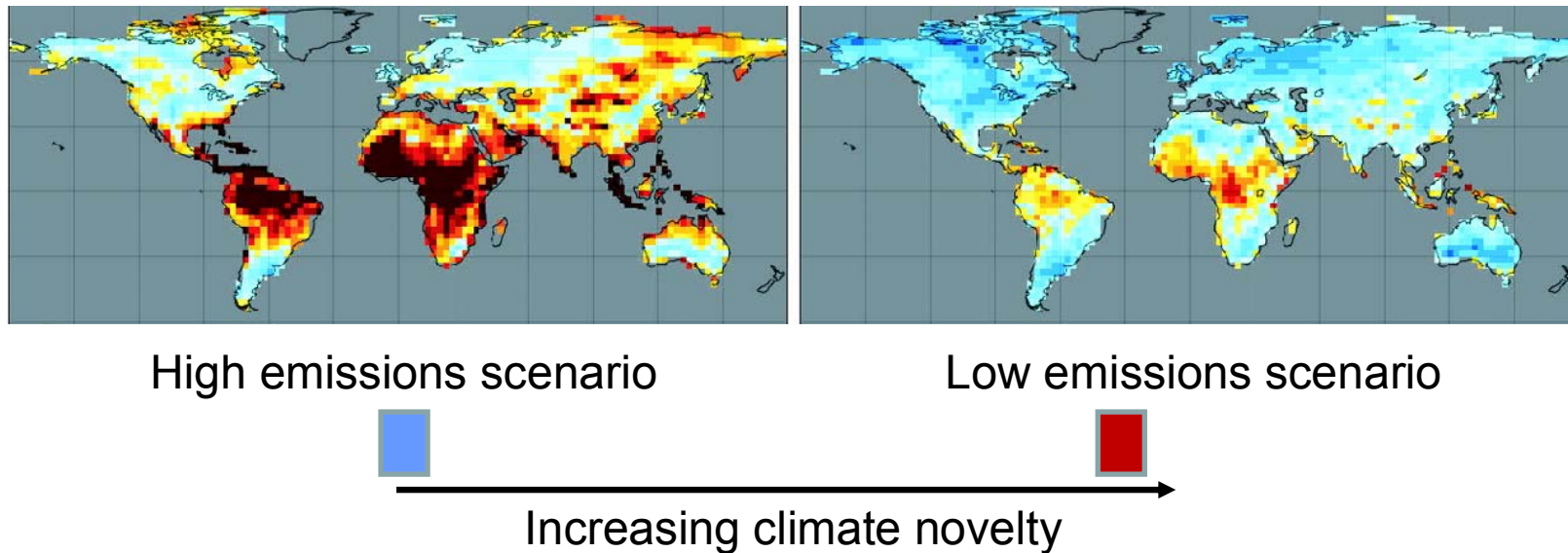
Soil moisture (mm)



1960-1990 to 2070-2100
(Hadley Center)

Climate-induced range shifts and extinctions?

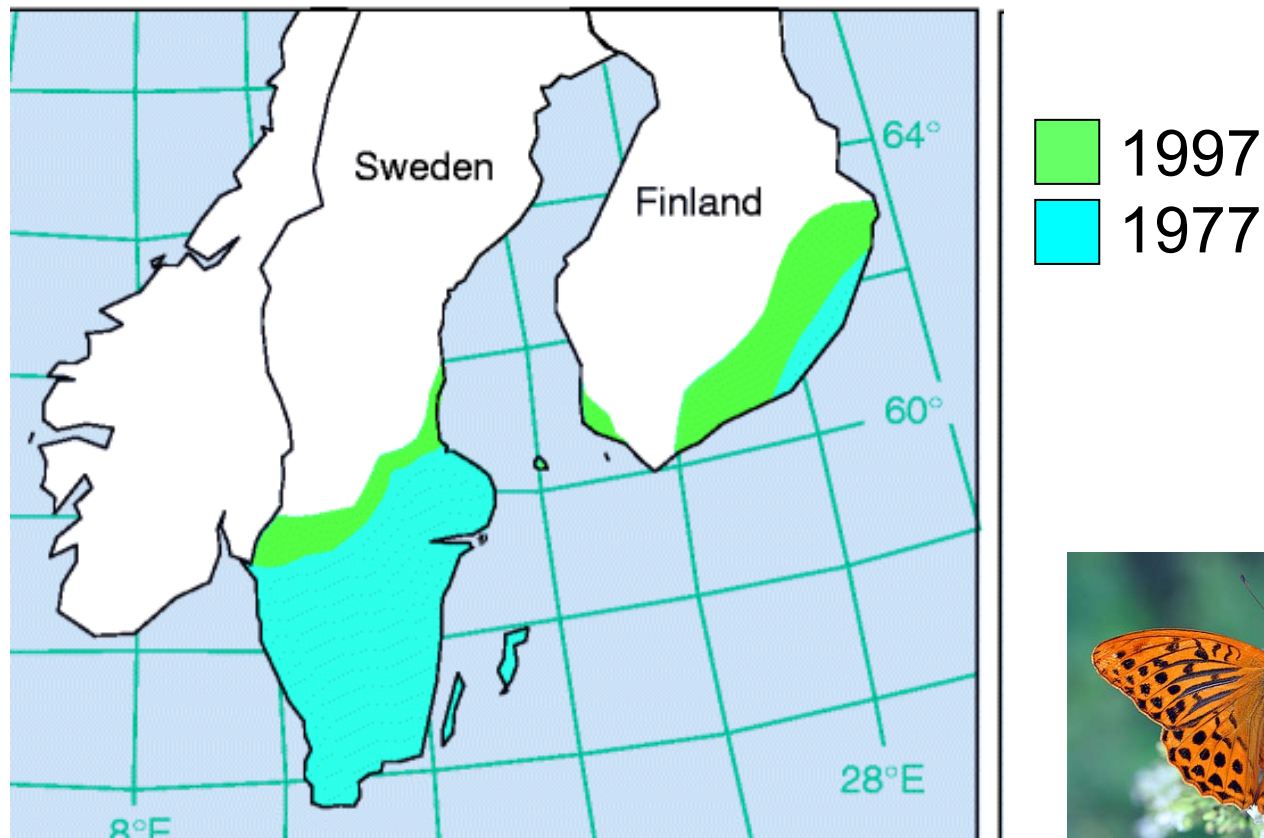
Novel Climates by 2100



(Williams *et al.* 2007)

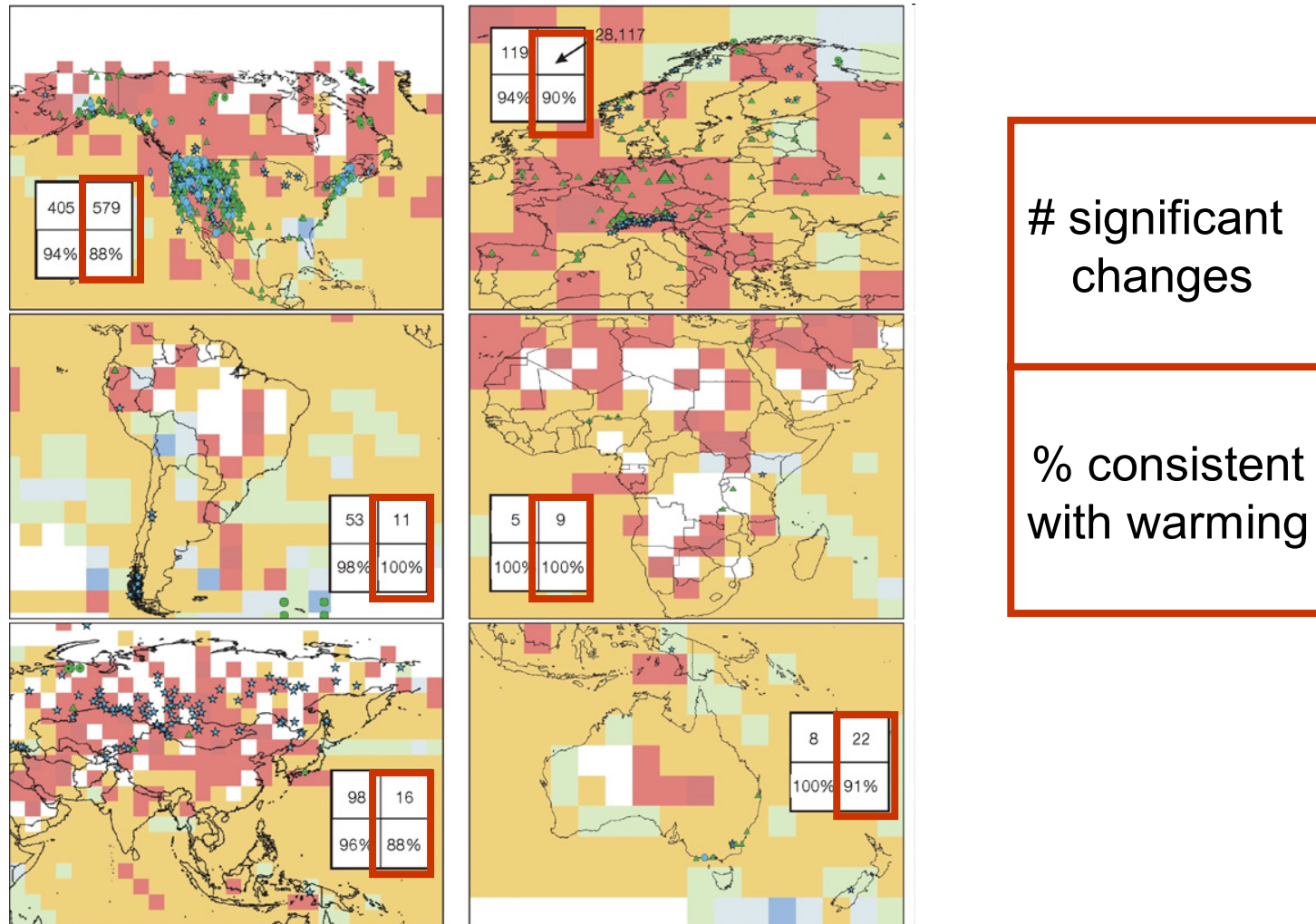
Recent range shifts

35 European non-migratory butterflies:
63% N range shift of 30-240 km during last century



Argynnis paphia
(Parmesan et al 1999)

Recent biological responses to warming



(Rosenzweig et al. 2008)

Climate-induced range shifts and extinctions?

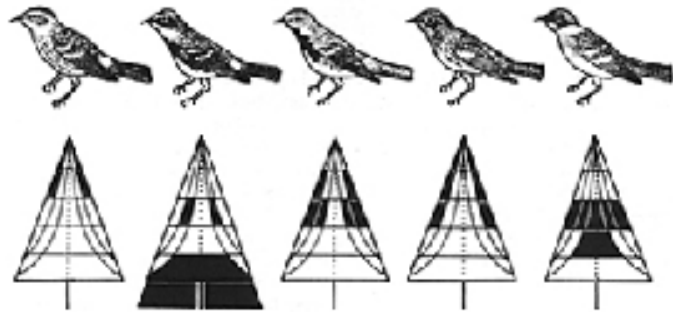
SOURCE: "Extinction Risk from Climate Change" Nature, January 8th, 2004

35%

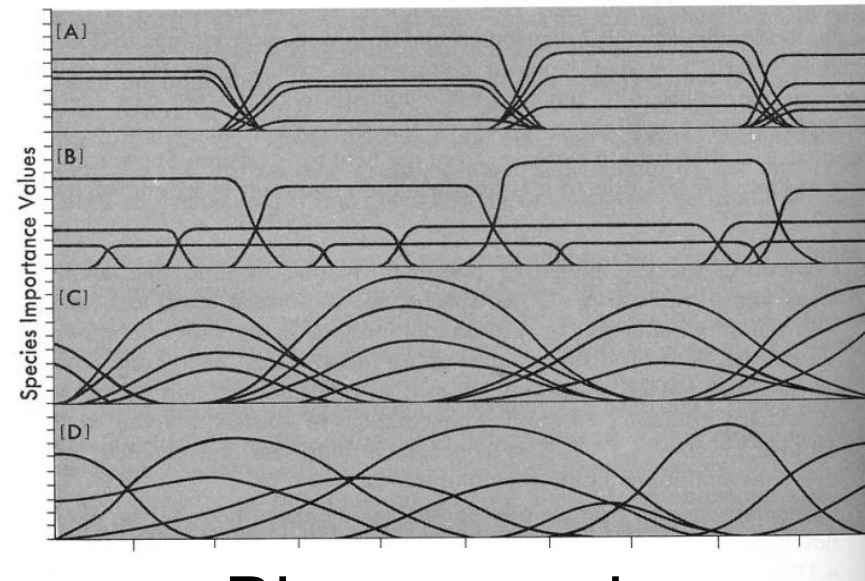
OF CURRENT
SPECIES
COULD GO
EXTINCT BY
2050 AS A
RESULT OF
CLIMATE
CHANGE.

(SEED on Thomas et al. 2004)

What sets a species' range boundary?

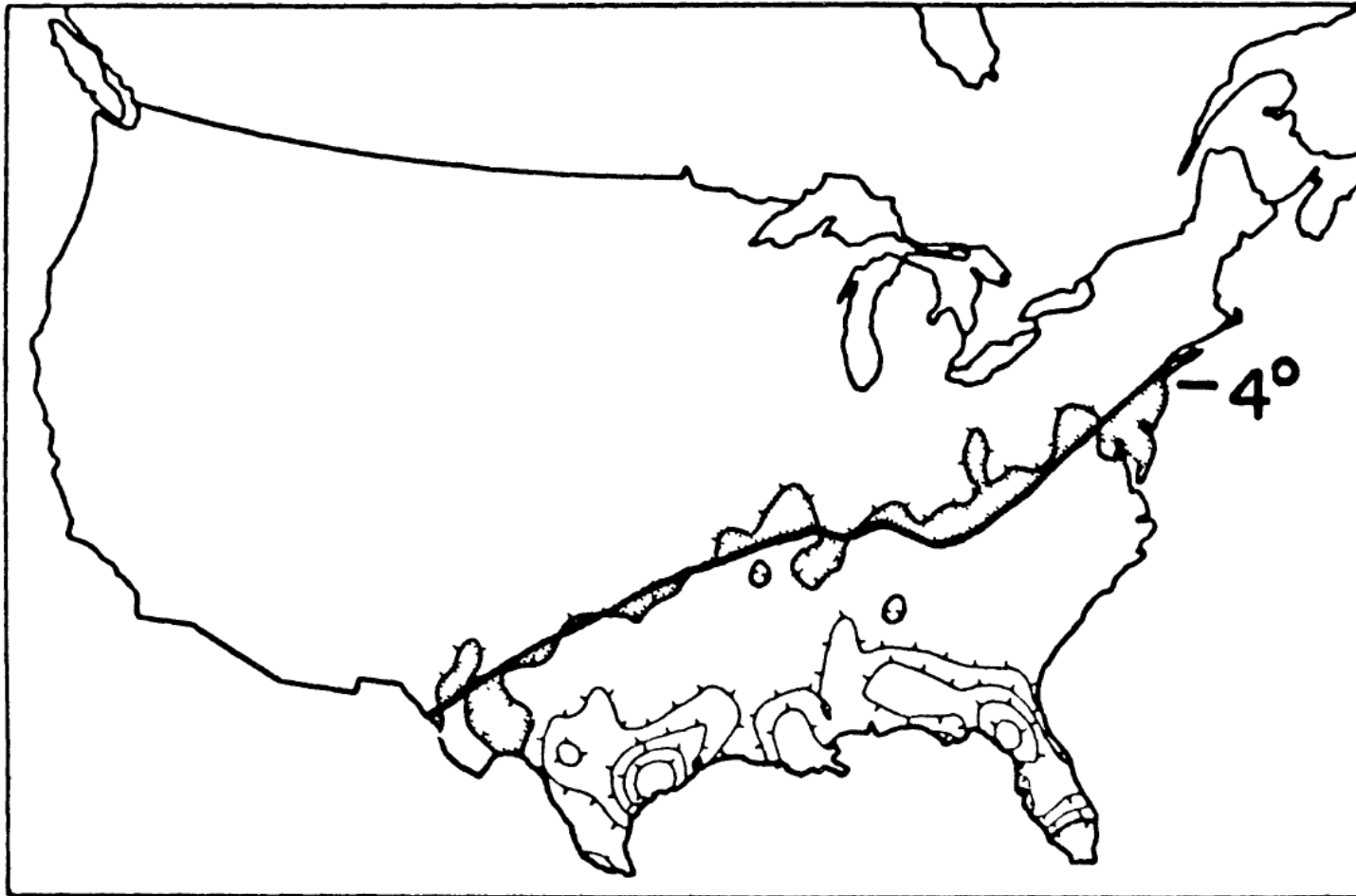


Community ecology

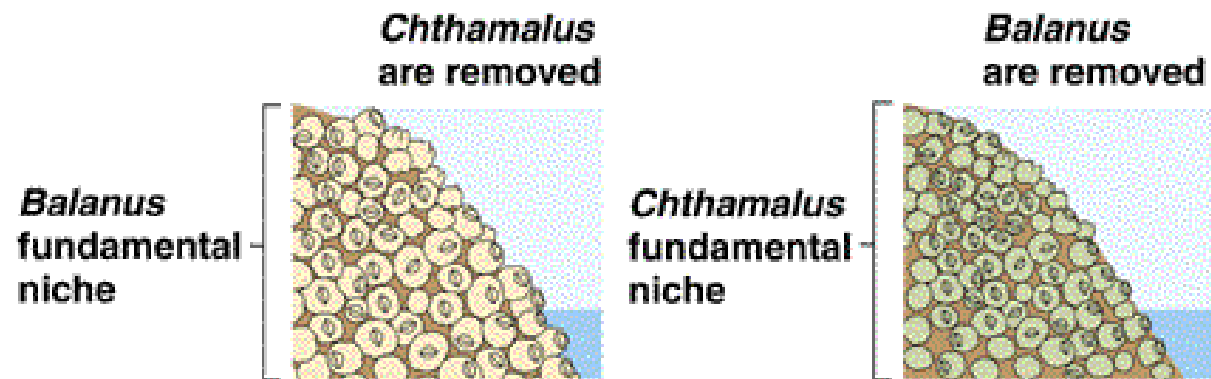
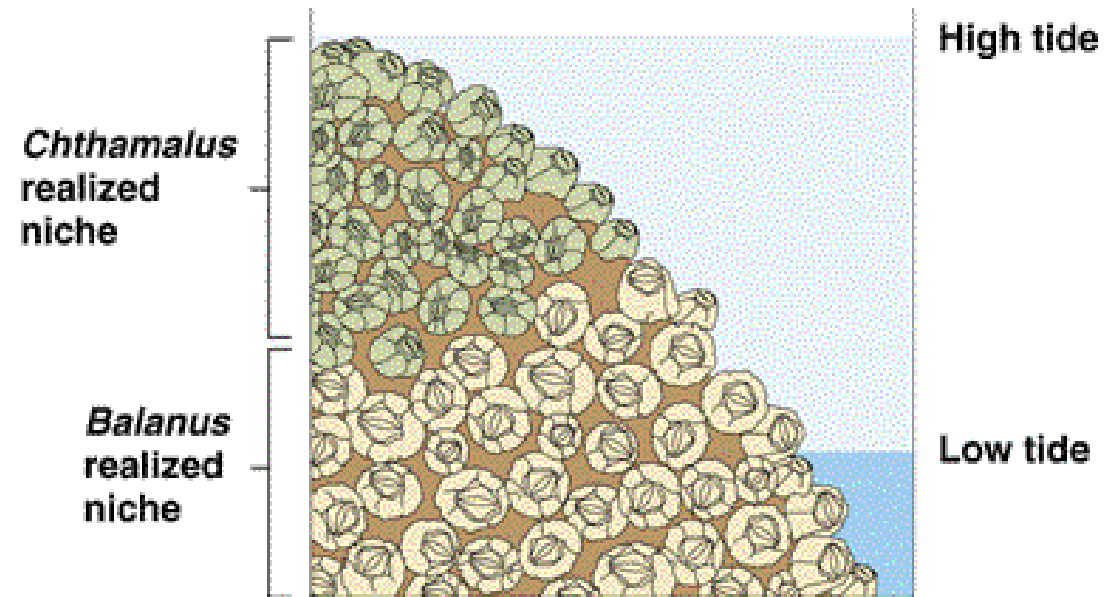


Biogeography

SCALE



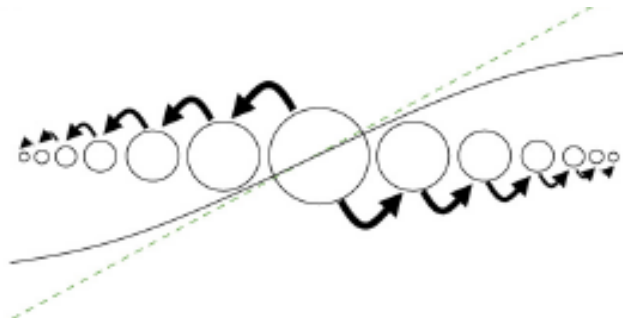
Eastern Phoebe
metabolic constraint
(Root 1988)



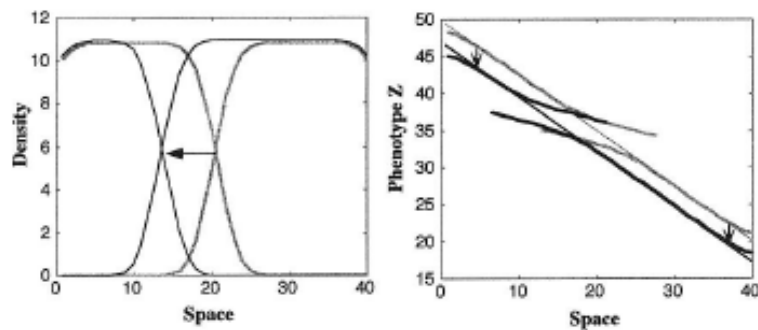
(Connell 1961)

Theory

Gene flow along selection gradient
(Kirkpatrick & Barton 1997)



Incorporate species interactions
(Case & Taper 2000)



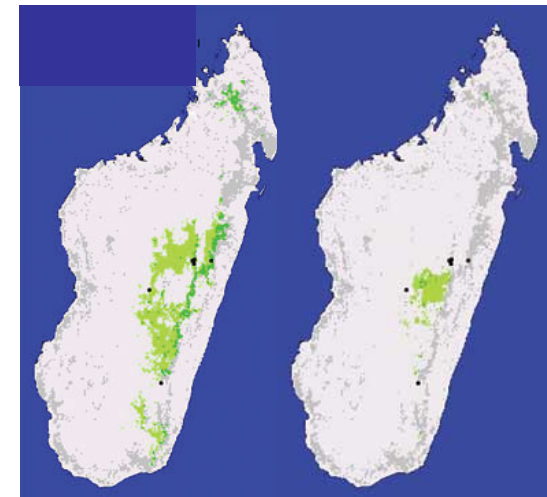
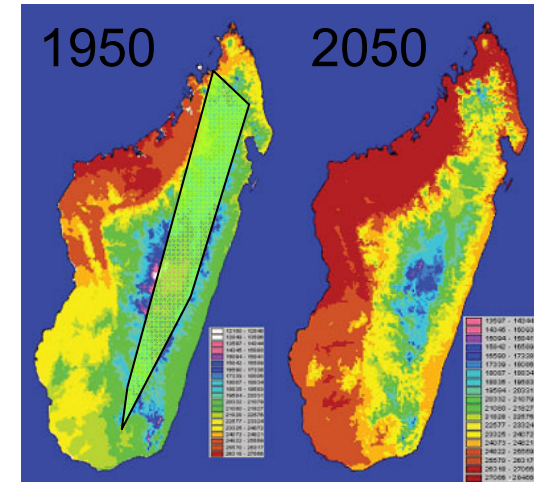
Practice: Correlative models

specimen localities



 Climatic envelope

Temperature



(Lees 2002)

Range models – Theory

Single populations

Multiple populations

Patch models

Genetic models

SPACE

TIME (no genotypic variation)

Trait evolution in response to climate change

Single populations

$$N_{t+1} = N_t + b - d + i - e$$

b- births

d- deaths

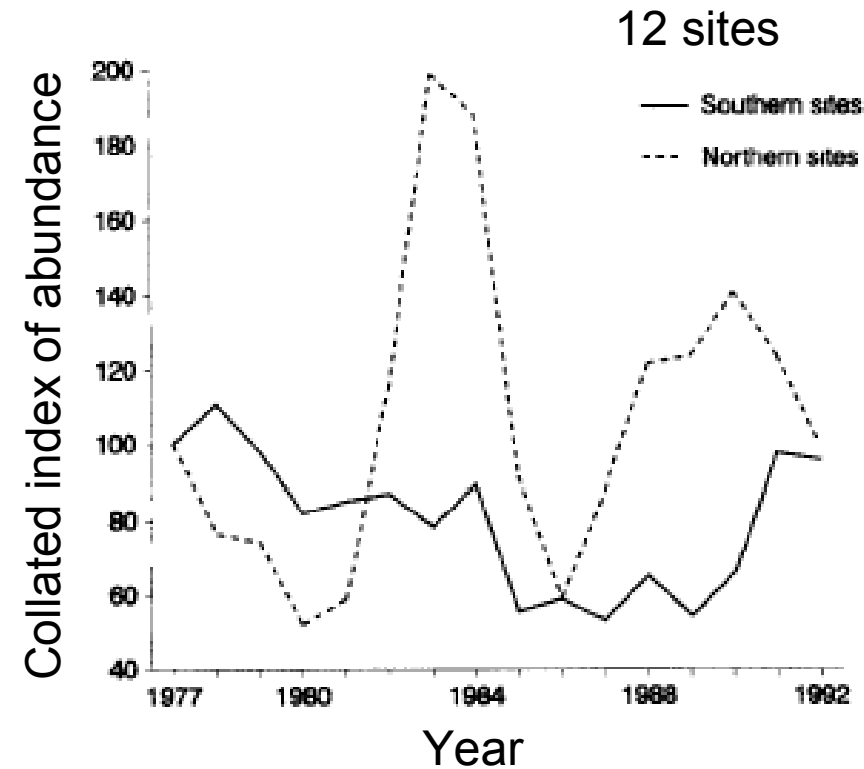
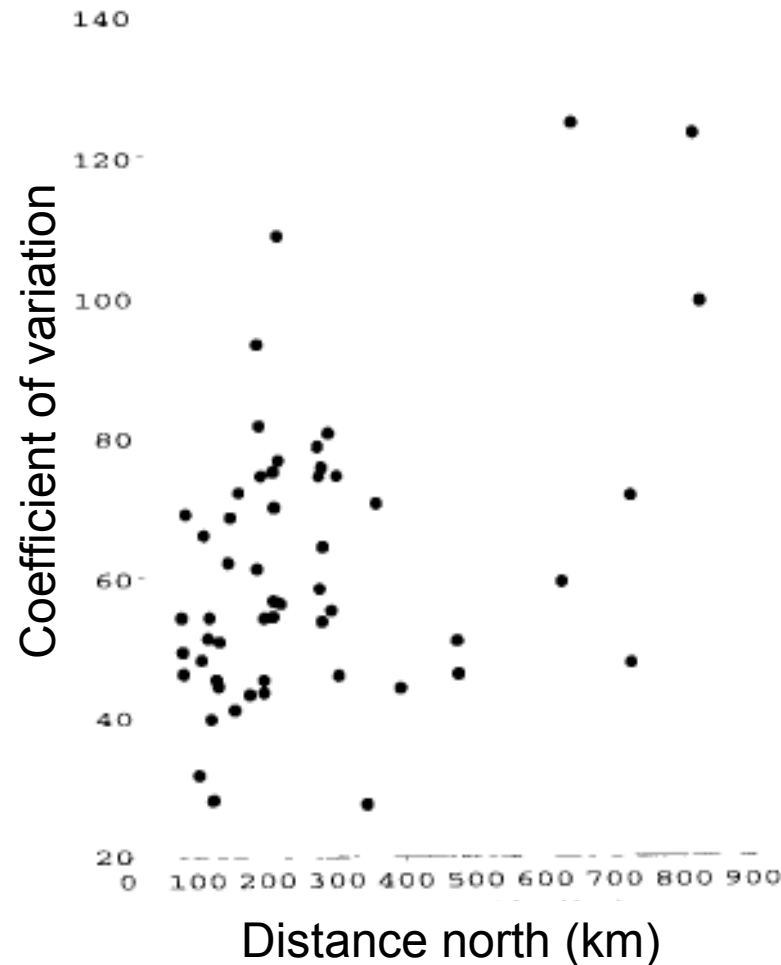
i- immigrants

e- emigrants

Species occur when $b-d+i-e > 0$

Hard to detect how parameters changes causes range edge

Population fluctuations at northern range



Less density regulation?
Abiotic factors?



24 species of British butterflies
1976-1992
(Thomas et al. 1994)

Multiple populations

Environmental heterogeneity, no evolution

$$\frac{dn(x)}{dt} = \overset{\text{colonization}}{n(x)[k(x)-n(x)]c(x)} - \overset{\text{extinction}}{e(x)n(x)}$$

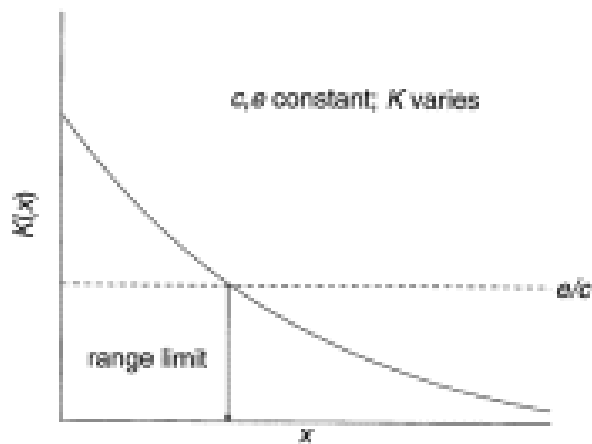
$$n^*(x) = k(x) - e(x)/c(x)$$

absent where

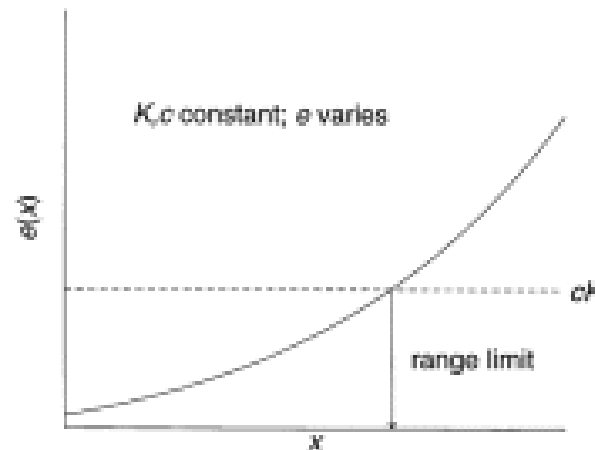
$$k(x) < e(x)/c(x)$$

$k(x)$ - fraction of suitable patches
 $e(x)$ - extinction rate (per occupied patch)
 $c(x)$ - colonization rate of suitable, empty patches (per occupied patch)

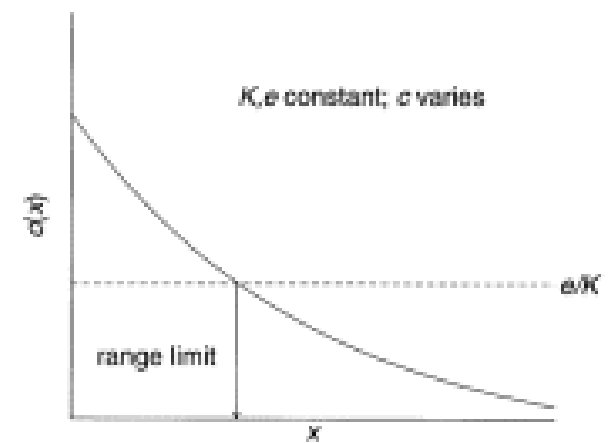
Gradient in habitat availability



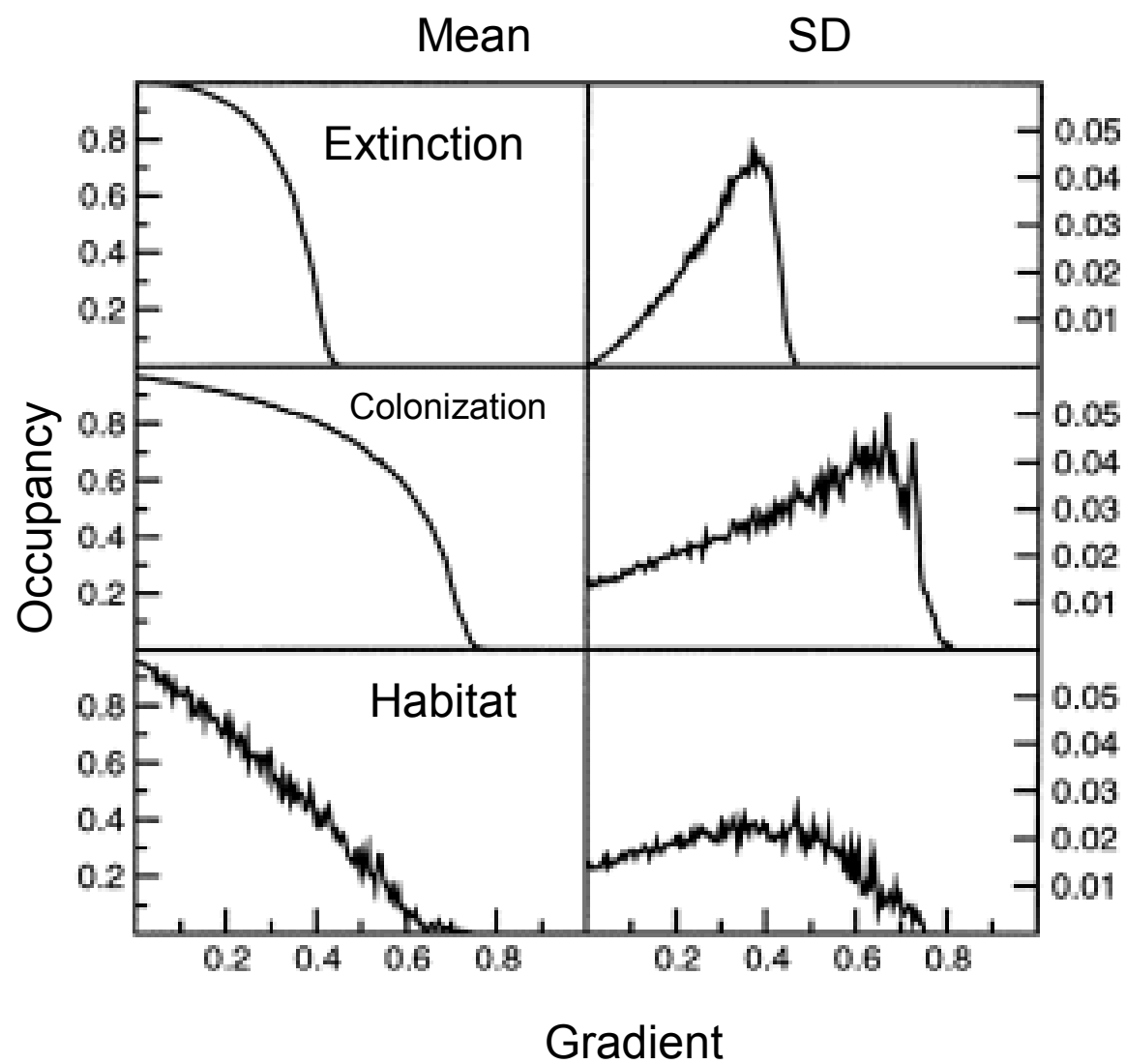
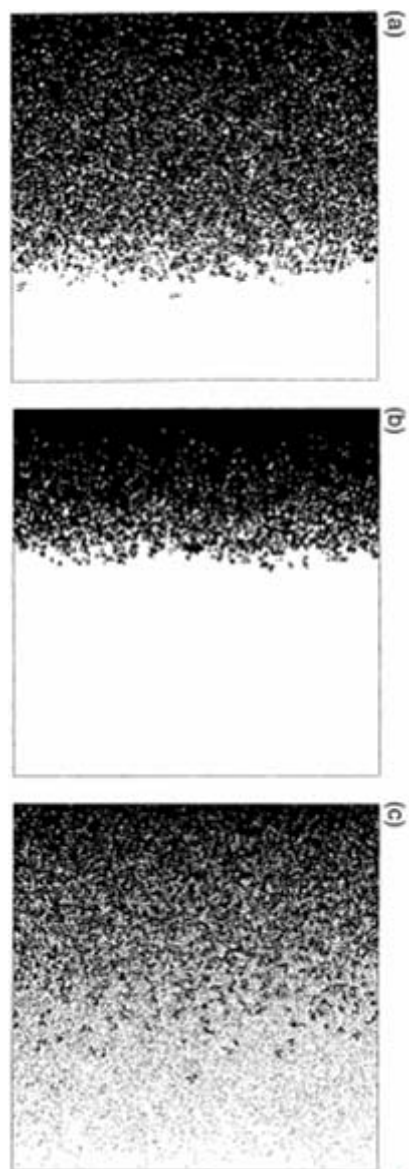
extinction rate



colonization success



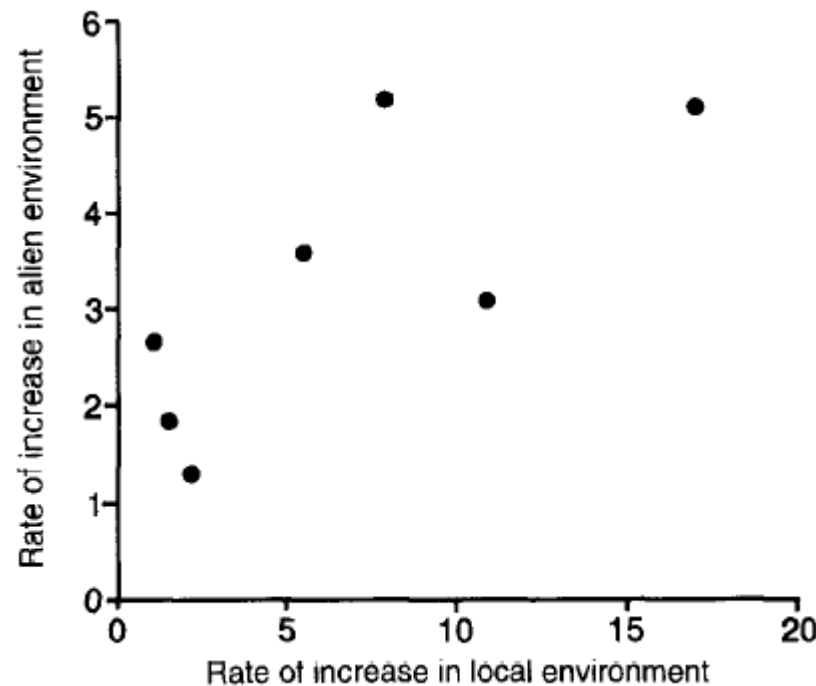
Levins metapopulation models
 (Holt and Keitt 2000)



Multiple populations

Environmental heterogeneity with evolution

What prevents traits evolving to enable an organism to expand its range boundaries? (Mayr 1962)



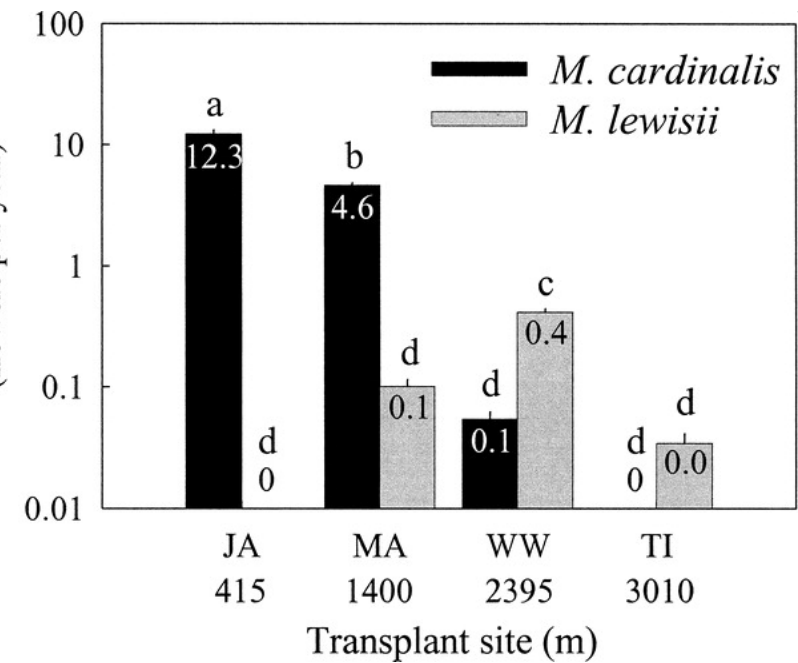
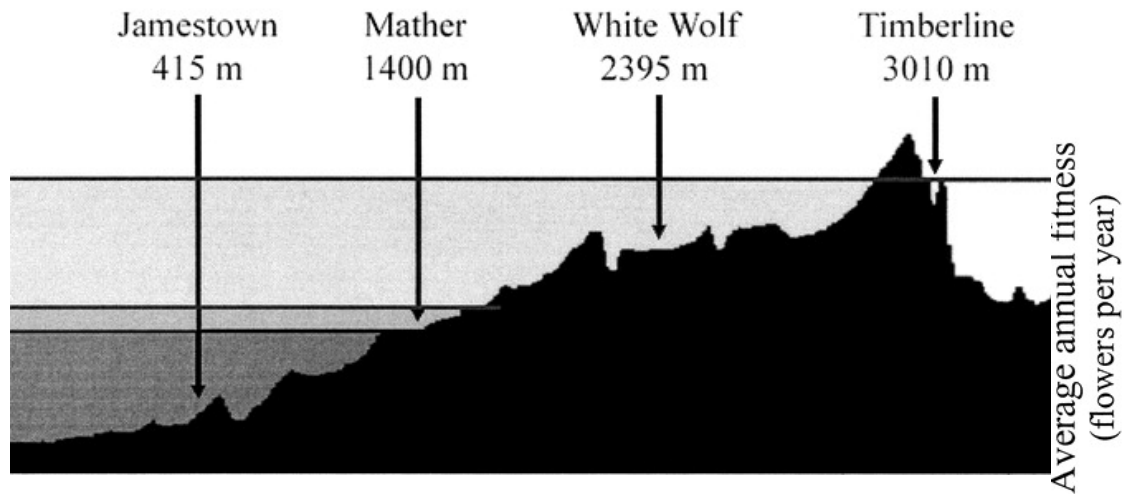
Species from poor environments-
Weak performance in other environments



Phlox drummondii
(Schmidt and Levin 1985)

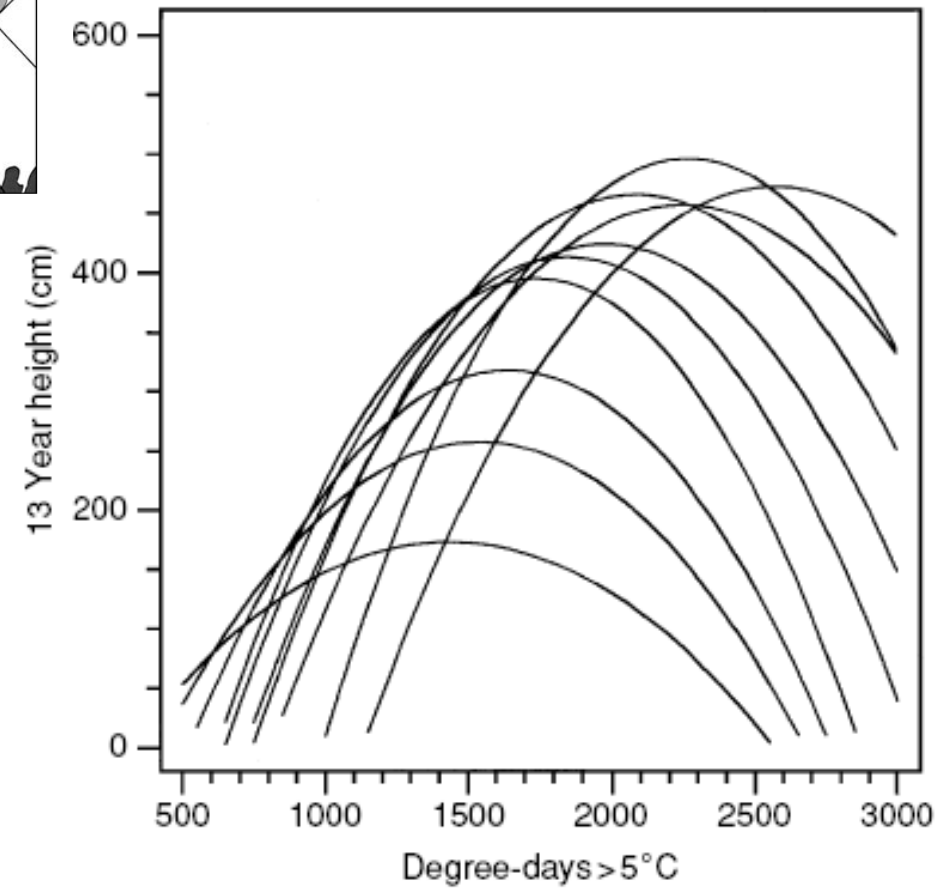
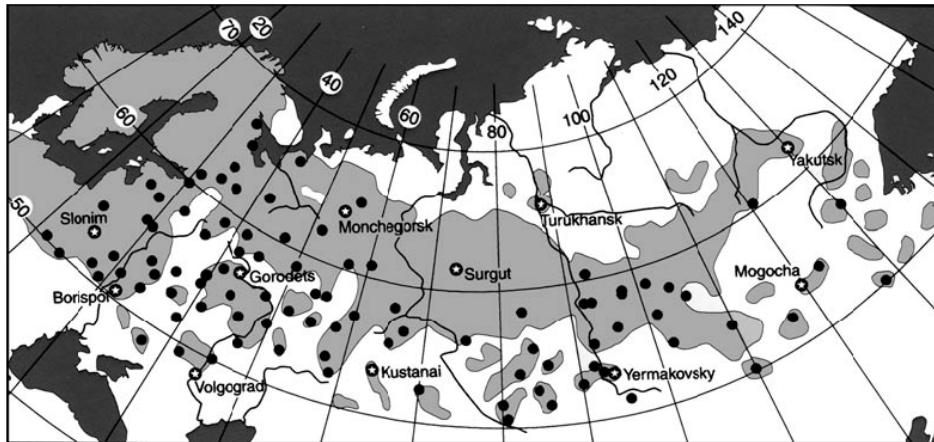


M. lewisii
 Sympatry
 M. cardinalis



greatest average fitness at elevations central within the range

(Angert and Schemske 2005)



Pinus sylvestris
(Rehfeldt et al. 2002)

What prevents traits evolving to enable an organism to expand its range boundaries? (Mayr 1962)

LOW HERITABILITIES (proportion phenotypic var due to genetic var)

1. Low n : low genetic variation
2. Directional selection in marginal environments
3. Environmental variability in marginal environments

$$\text{Var}(\textit{Phenotype}) = \text{Var}(\textit{Genotype}) + \text{Var}(\textit{Environment}) + 2 \text{Cov}(\textit{G}, \textit{E})$$

What prevents traits evolving to enable an organism to expand its range boundaries? (Mayr 1962)

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GENETIC INTERACTIONS

4. Require evolution of multiple traits
5. Genetic trade-offs between fitness in favorable and stressful envi

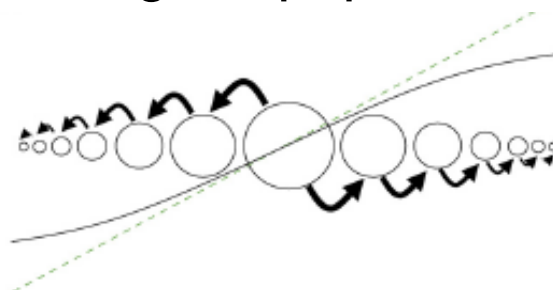
What prevents traits evolving to enable an organism to expand its range boundaries? (Mayr 1962)

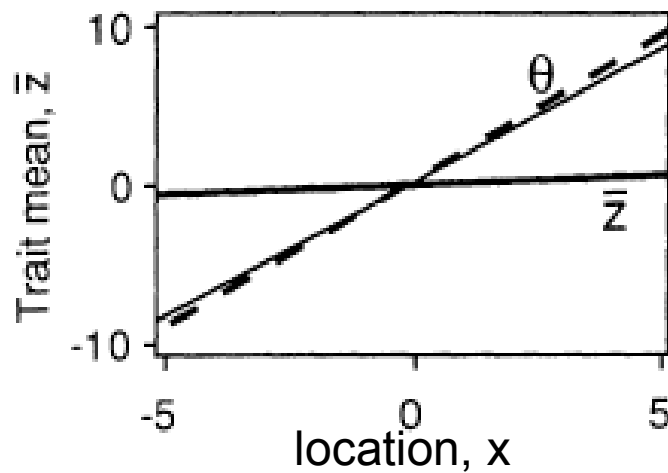
LOW HERITABILITIES (proportion phenotypic var due to genetic var)

1. Low n : low genetic variation
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3. Environmental variability in marginal environments

GENETIC INTERACTIONS

4. Require evolution of multiple traits
5. Genetic trade-offs between fitness in favorable and stressful envi
6. Deleterious mutations under stressful conditions
7. Genotypes favored in marginal populations swamped by gene flow from center





Z- trait mean

θ- trait optimum

b- rate of environmental change in space

$$\theta = bx$$

$$\frac{\partial \bar{z}}{\partial t} = \underbrace{\frac{\sigma^2}{2} \frac{\partial^2 \bar{z}}{\partial x^2} + \sigma^2 \frac{\partial \ln(n)}{\partial x} \frac{\partial \bar{z}}{\partial x}}_{\text{Gene flow}} + \underbrace{G \frac{\partial \bar{r}}{\partial \bar{z}}}_{\text{Directional selection}},$$

diffusion asymmetrical

Fitness decreases with distance from optimum

$$r(z) = r_{\max} - \frac{(\theta(x) - z)^2}{2V_s}$$

σ^2 - variance in offspring dispersal

n - population size

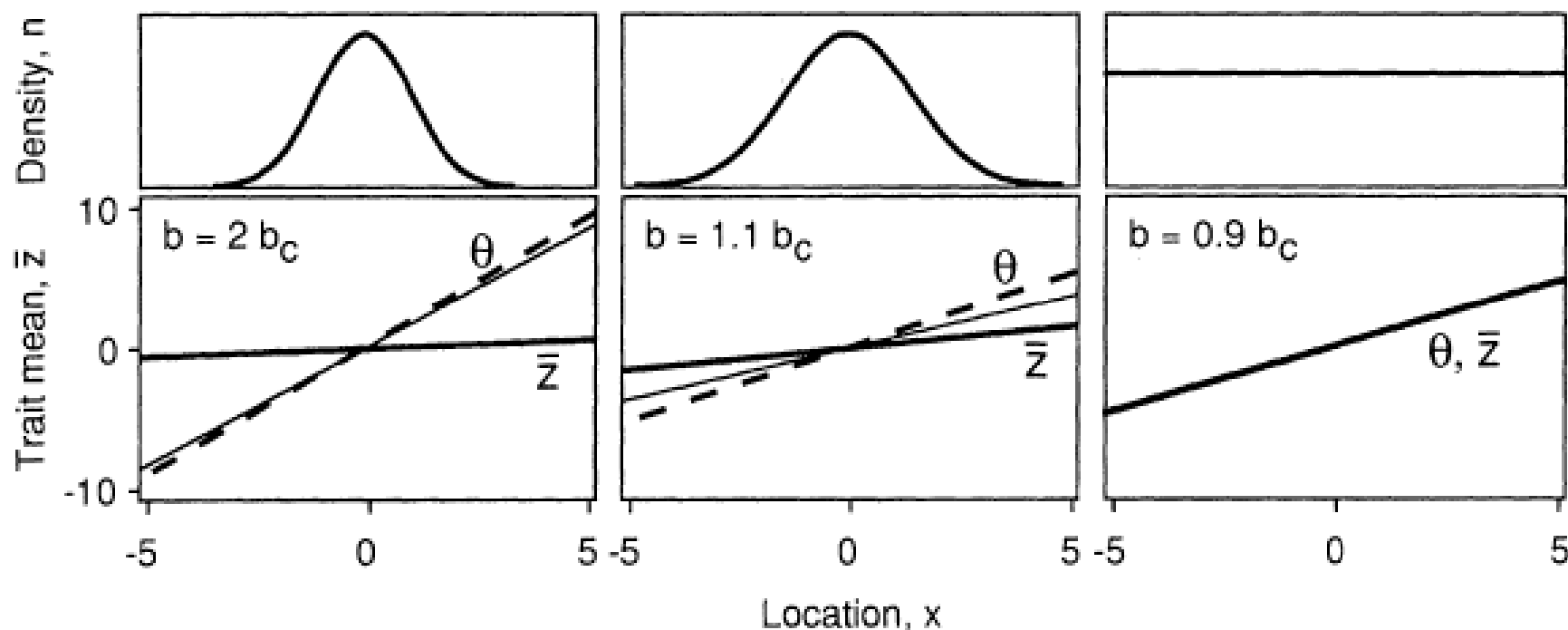
G - additive genetic variance

V_s - strength of stabilizing selection

(Kirkpatrick & Barton 1997)

Equilibrium population density increasing function of local fitness

$$n(x) = ke^{\bar{r}\gamma}$$

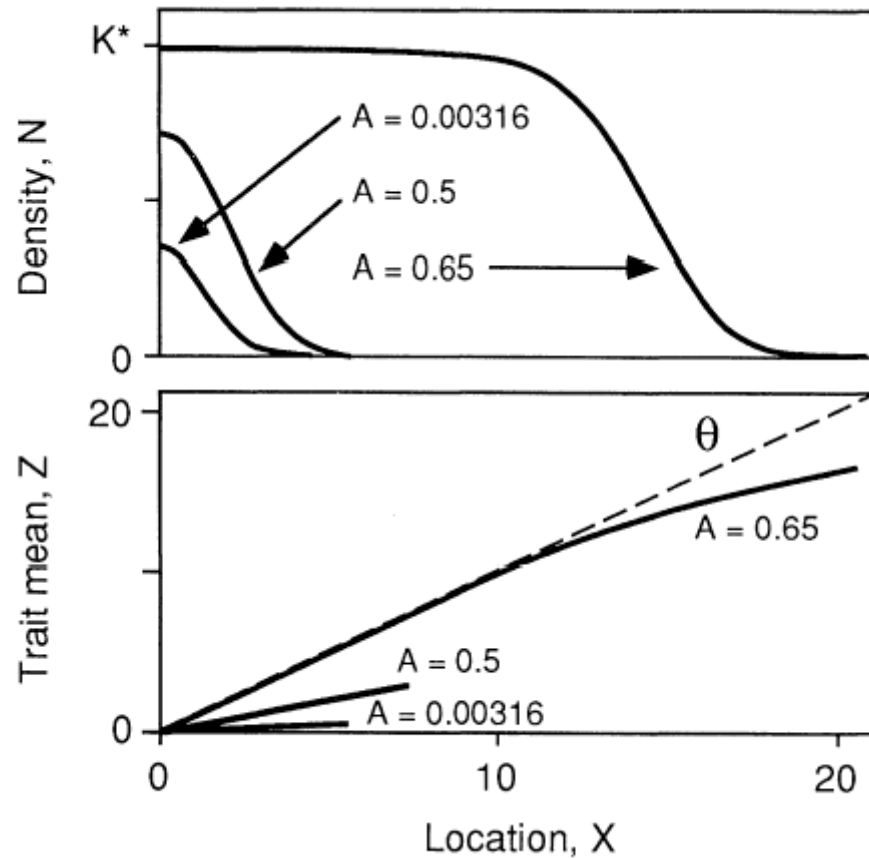


σ^2 - variance in offspring dispersal
 n - population size, k - carrying cap
 G - additive genetic variance
 V_s - strength of stabilizing selection
 γ - strength of density regulation

$$\frac{\partial \bar{z}}{\partial t} = \frac{\sigma^2}{2} \frac{\partial^2 \bar{z}}{\partial x^2} + \frac{(bx - \bar{z})}{V_s} \left[G - \sigma^2 \gamma \frac{\partial \bar{z}}{\partial x} \left(b - \frac{\partial \bar{z}}{\partial x} \right) \right]$$

(Kirkpatrick & Barton 1997)

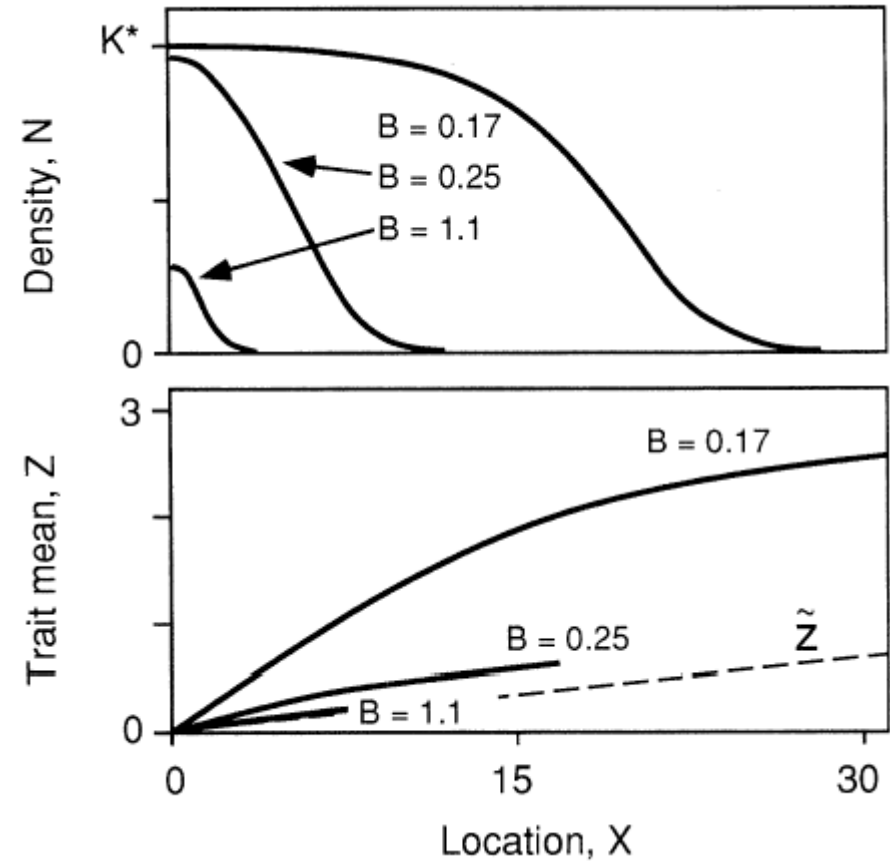
A- potential for local adaptation

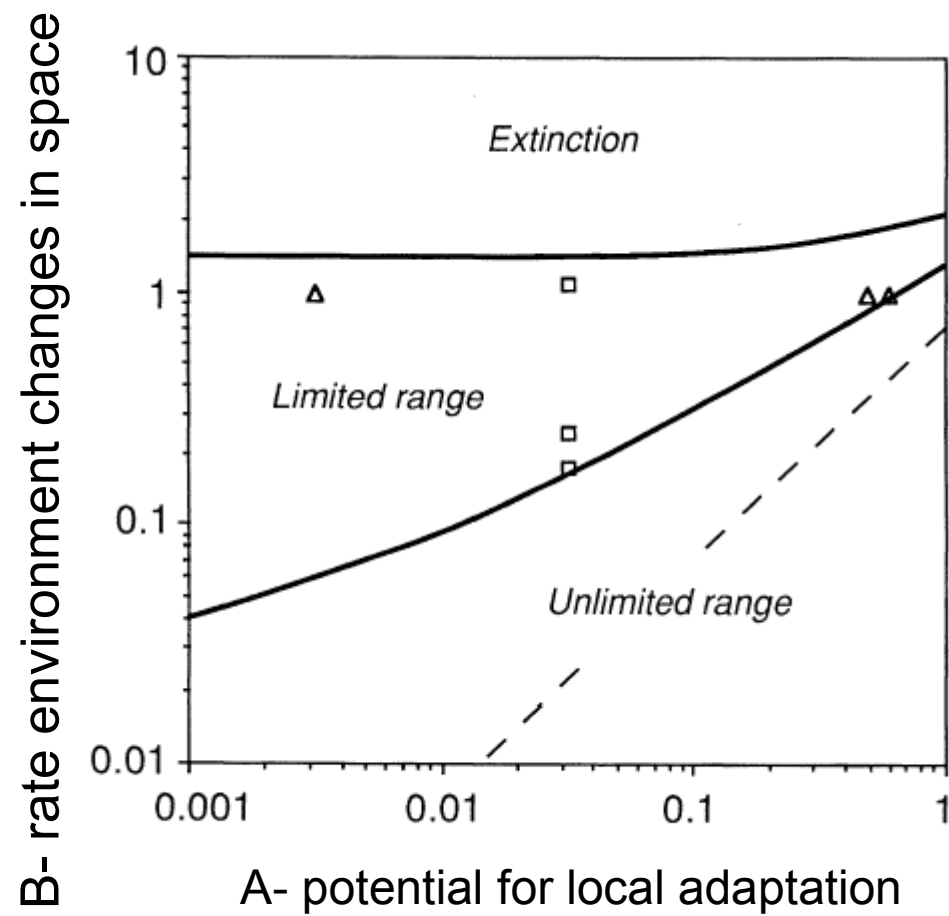


$$A = \frac{G}{2V_s r^*} = \frac{h^2 I_s}{2r^*},$$

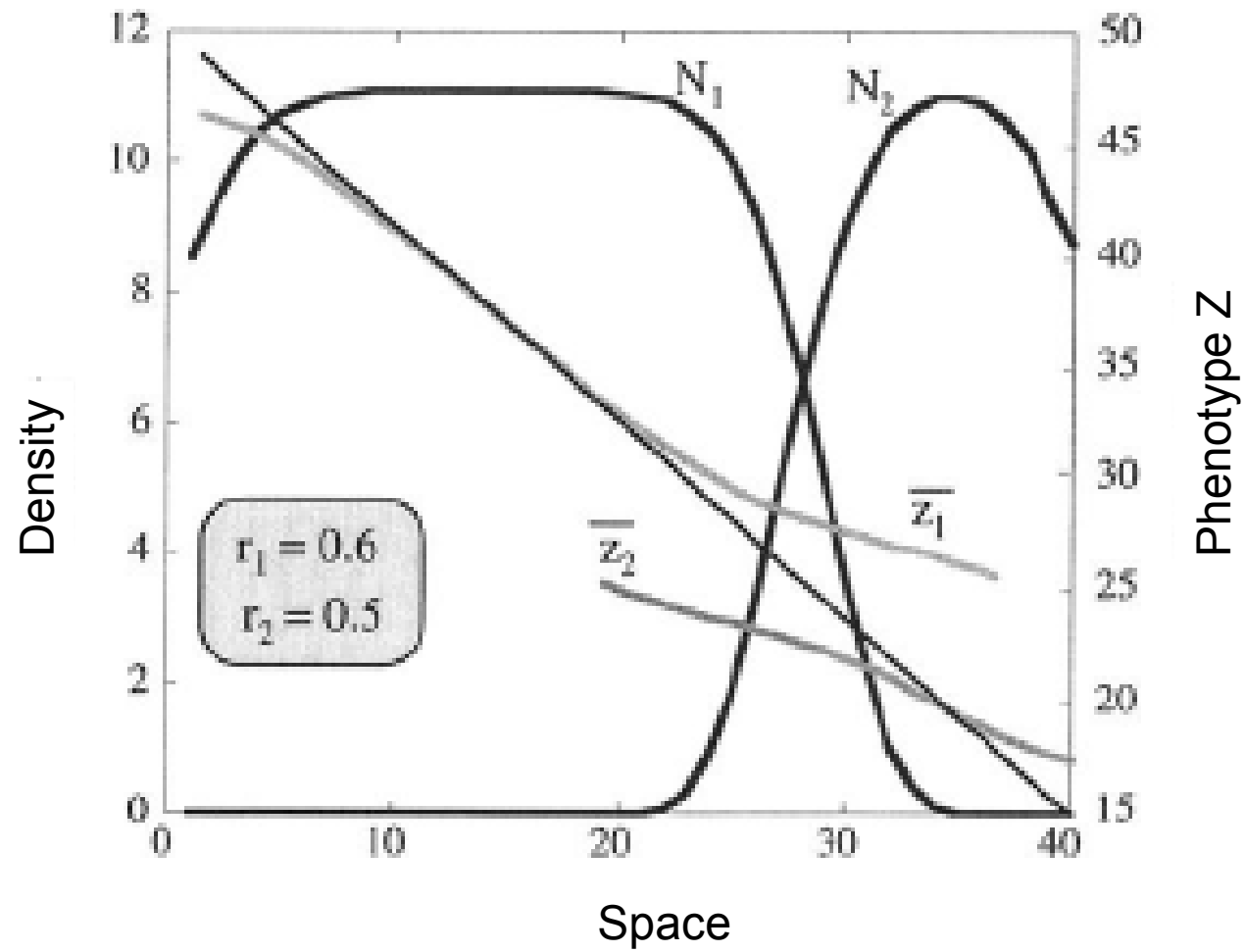
$$B = \frac{b\sigma}{r^* \sqrt{2V_s}},$$

B- rate environment changes in space



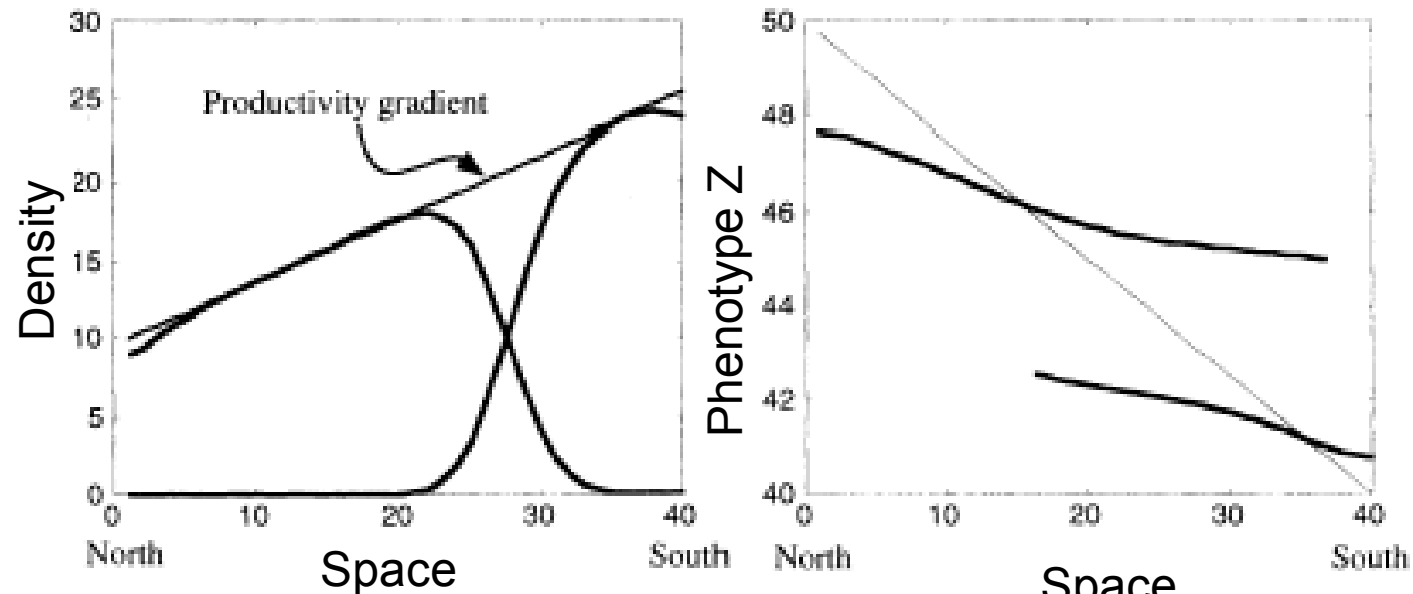


Species interactions

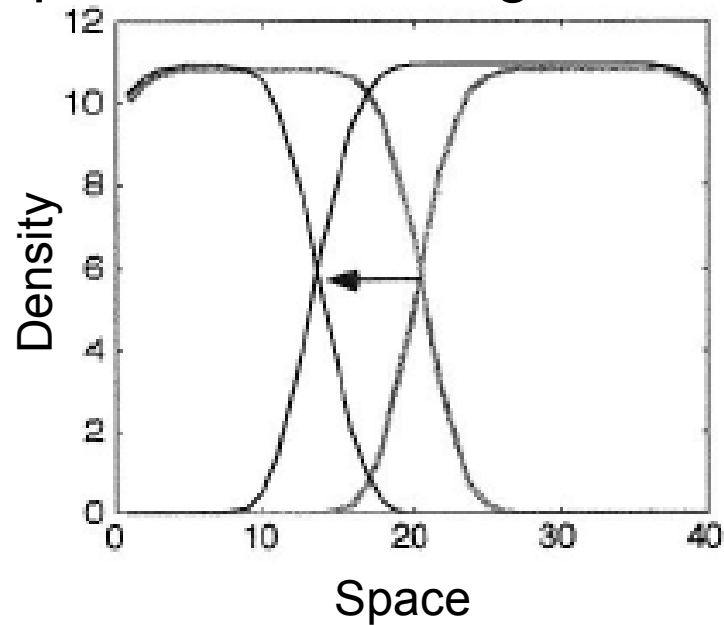


(Case & Taper 2000)

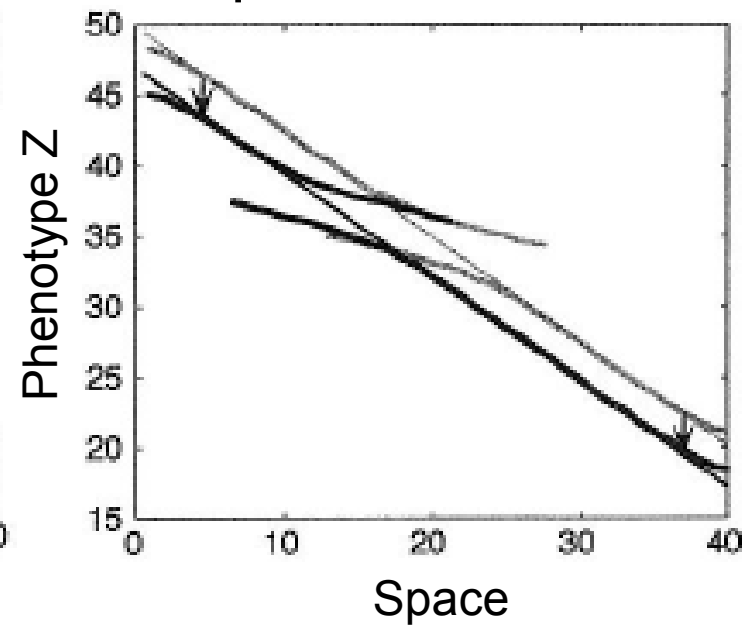
Productivity gradient



Rapid climate change



Space optimum shifts down



Range models – Theory

Single populations

Multiple populations

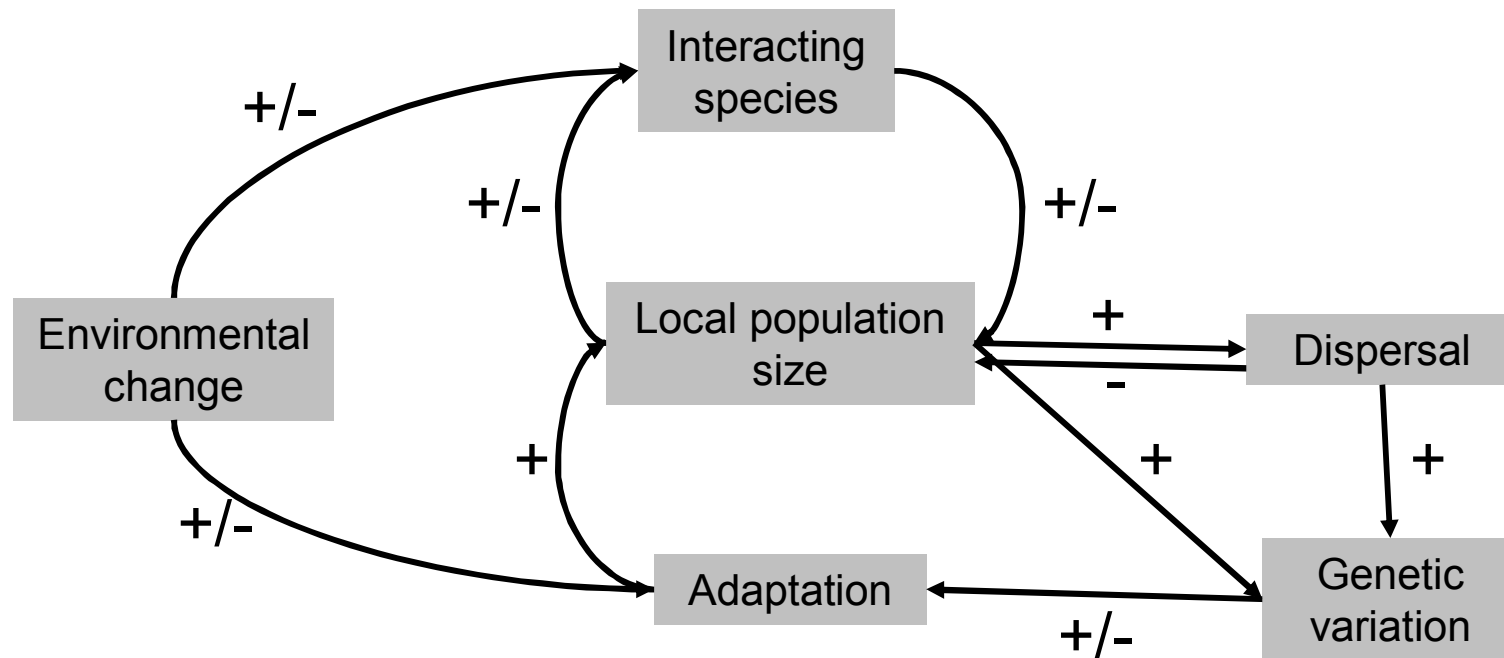
Patch models

Genetic models

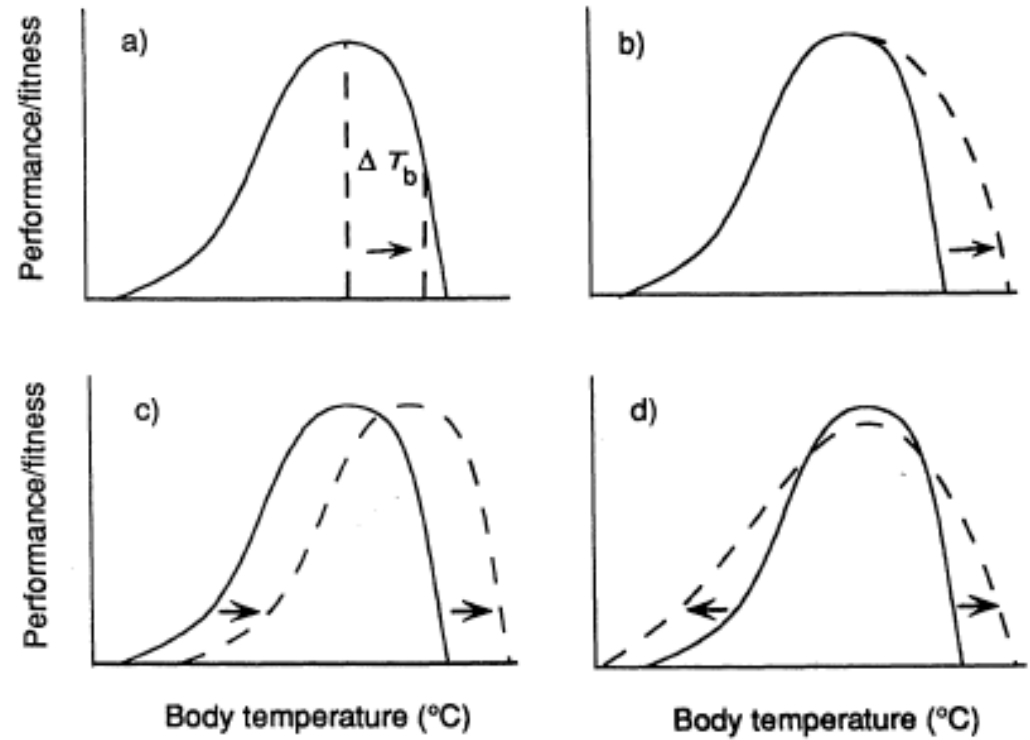
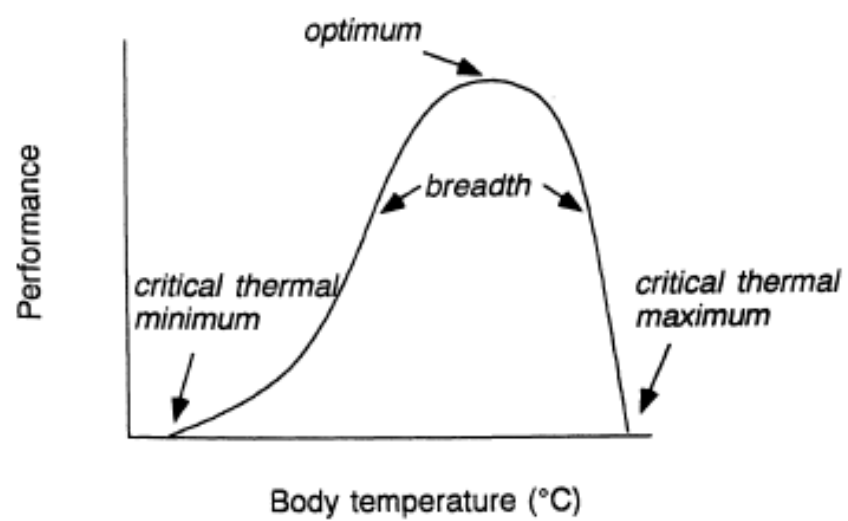
SPACE

TIME (no genotypic variation)

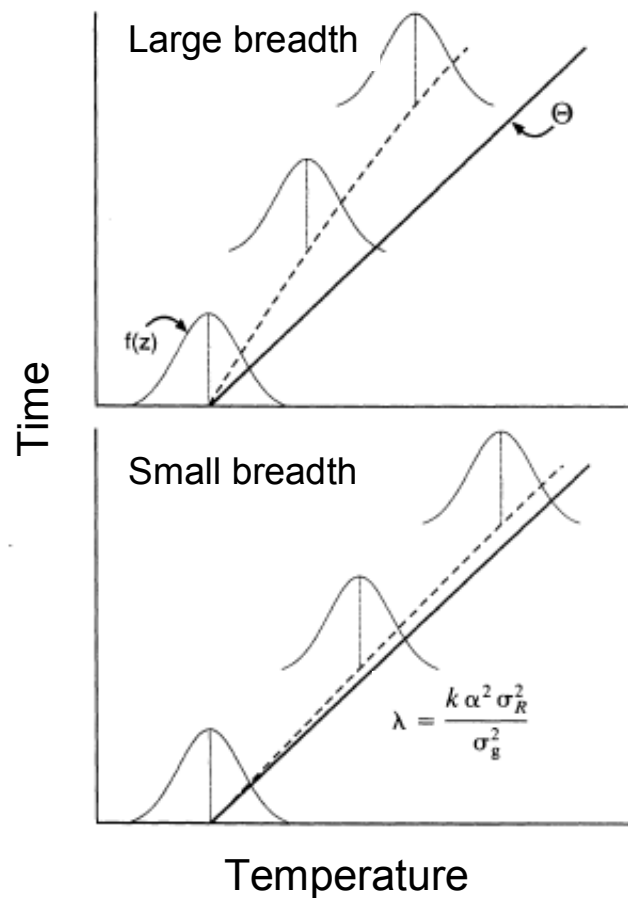
Trait evolution in response to climate change



(Lynch and Lande 1993)

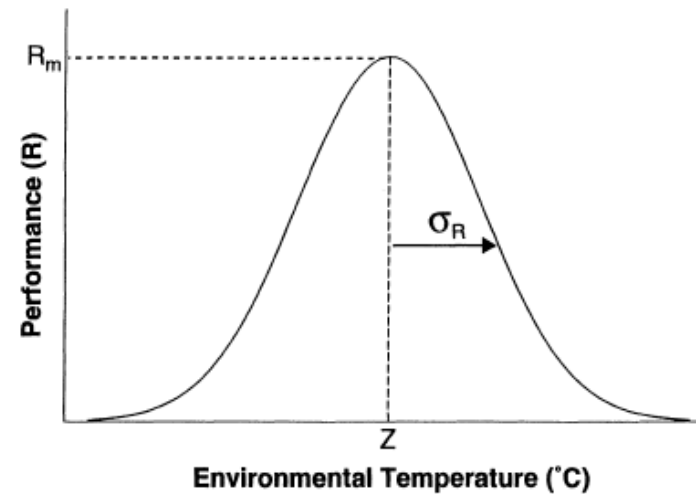


(Huey and Kingsolver 1993)



- Mean phenotype
- Environmental optimum

Large performance breadth
 → Increase evolutionary lag



$$\lambda = \frac{k \alpha^2 \sigma_R^2}{\sigma_g^2}$$

Fitness directly proportional to performance

$$r = \alpha R$$

k- rate of change of environmental temperature

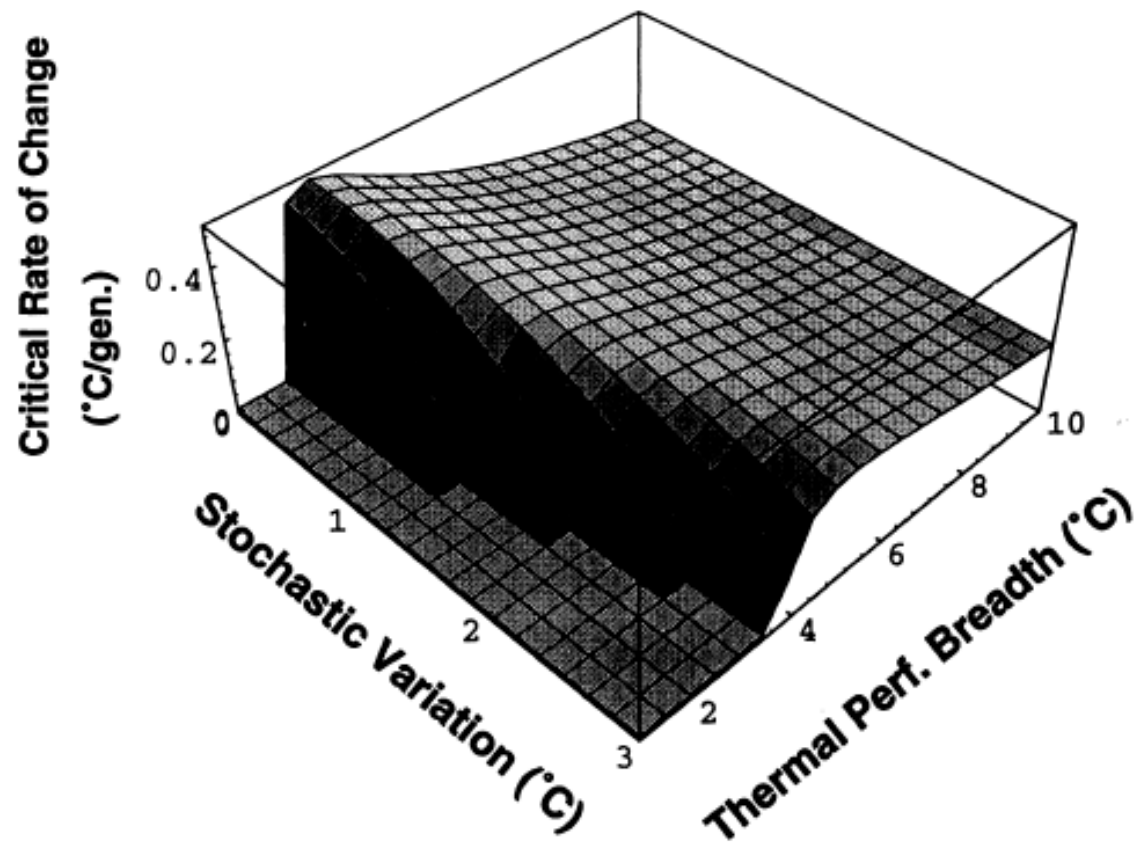
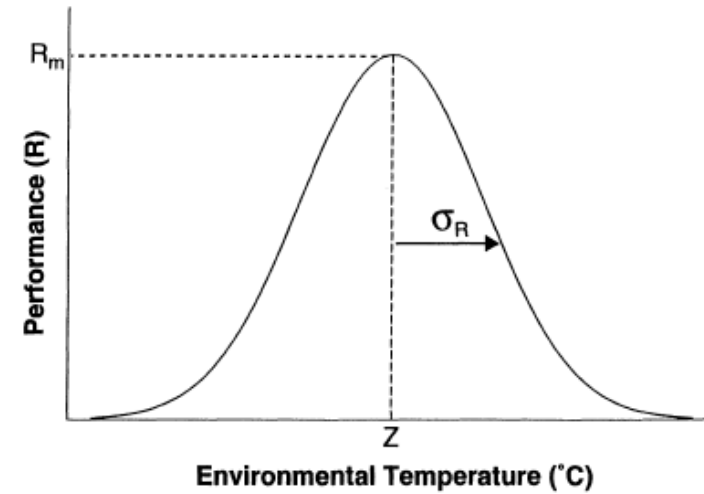
σ_g^2 - genetic variance

(Huey and Kingsolver 1993)

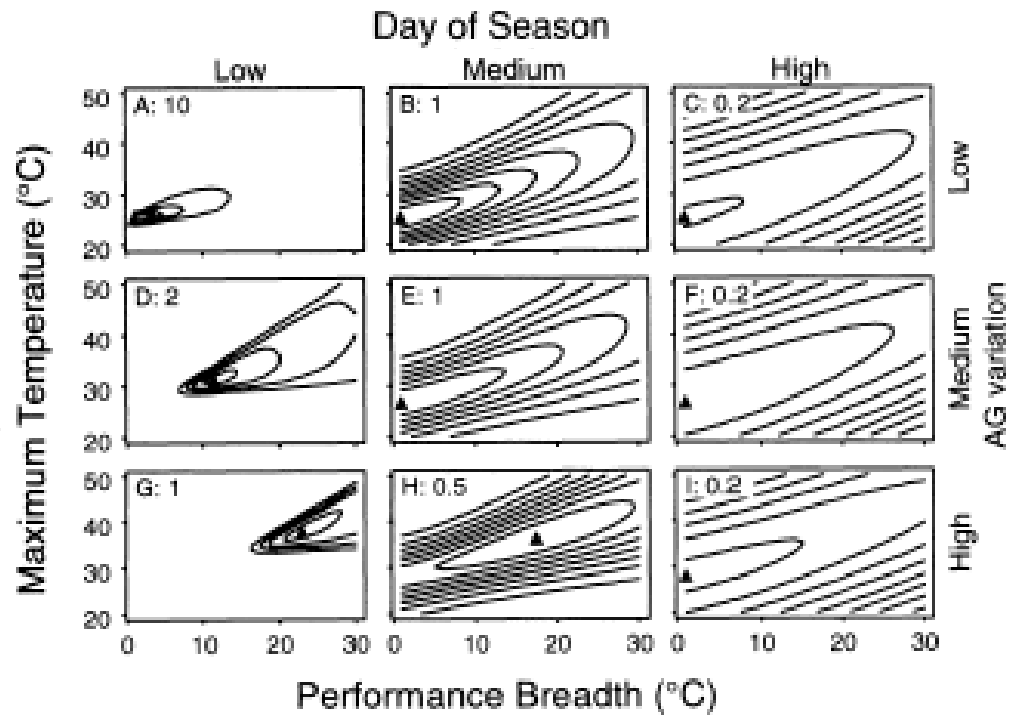
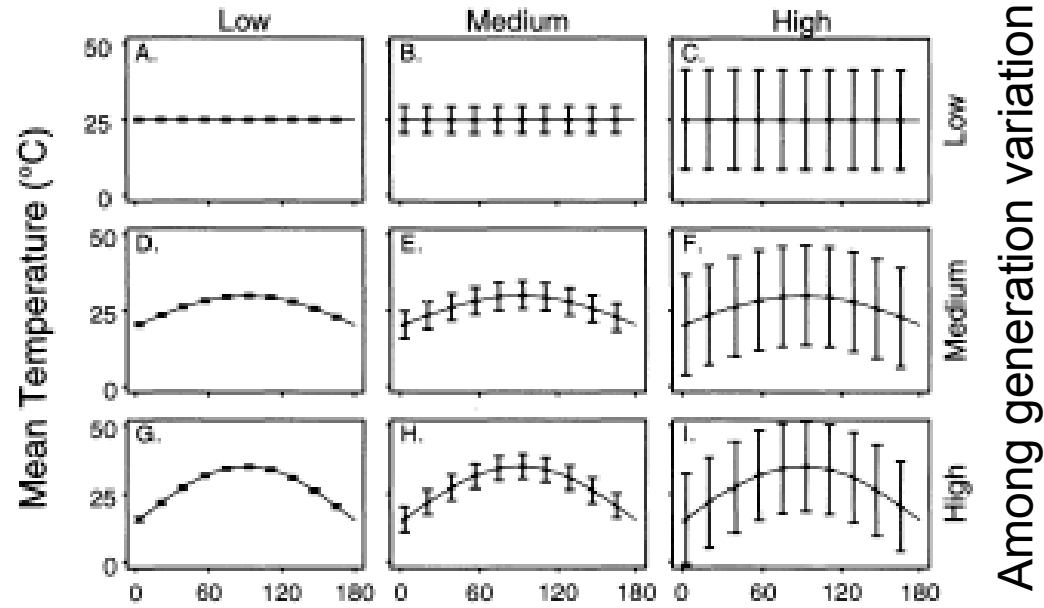
Critical rate of environmental change, k_c

$$k_c = \frac{h^2 \sigma_Z^2}{\alpha \sigma_R} \left(2\alpha R_m - \frac{\sigma_Z^2}{\alpha^2 \sigma_R^2} \right)^{0.5}$$

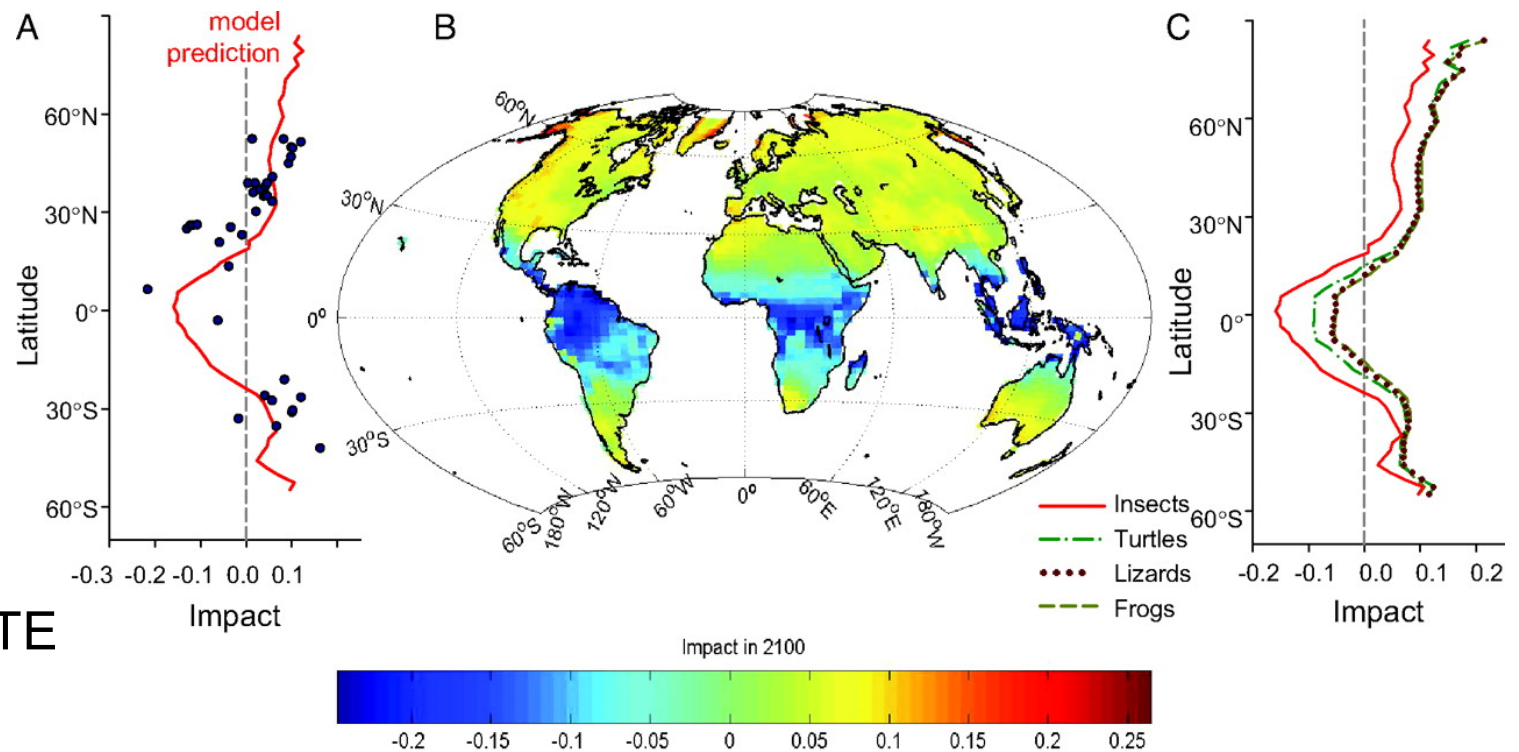
σ_g^2 - genetic variance
 Heritability $h^2 = \sigma_g^2 / \sigma_z^2$



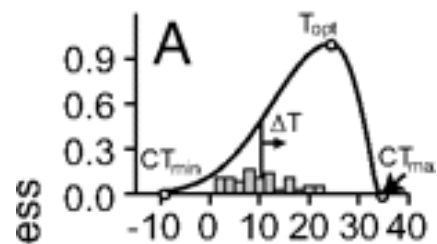
Within generation variation



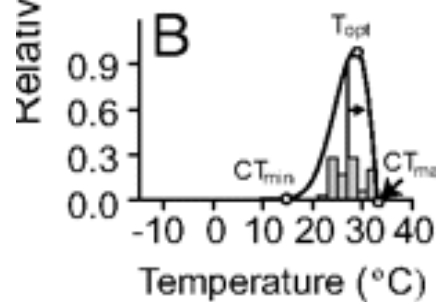
Numbers: contour interval
(Gilchrist 1995)



TEMPERATE



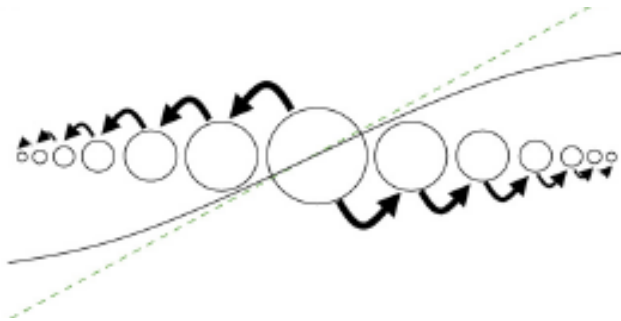
TROPICAL



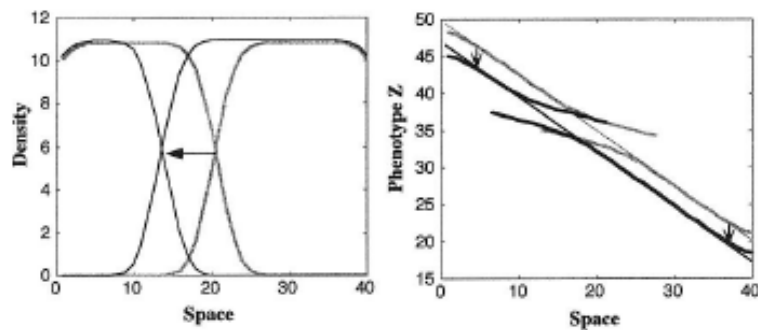
(Deutsch *et al.* 2008)

Theory

Gene flow along selection gradient
(Kirkpatrick & Barton 1997)



Incorporate species interactions
(Case & Taper 2000)



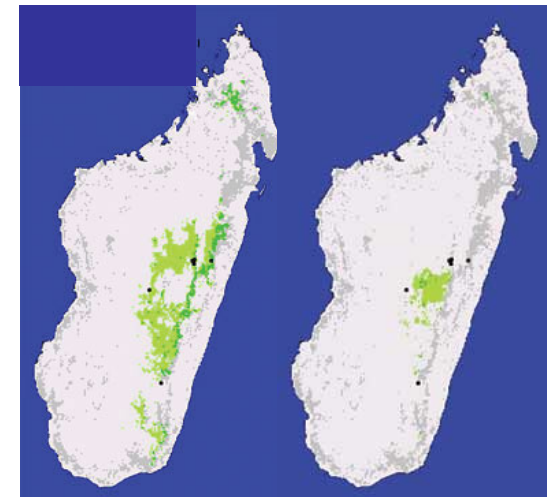
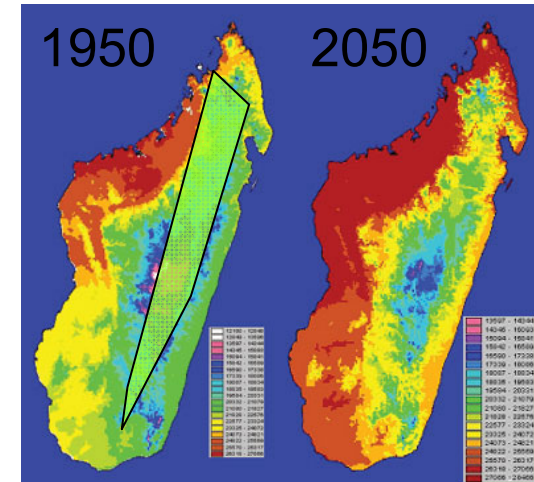
Practice: Correlative models

specimen localities



 Climatic envelope

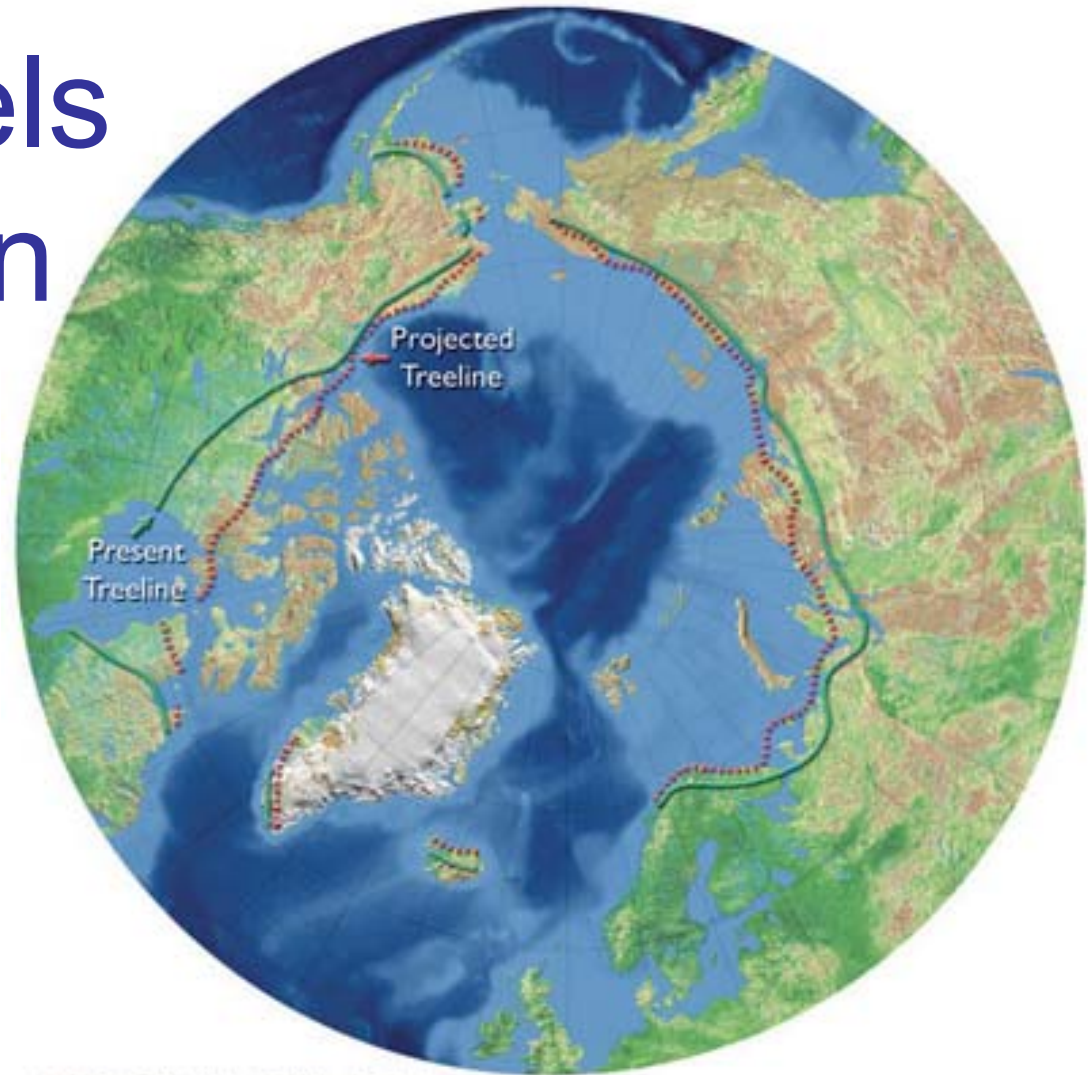
Temperature



(Lees 2002)

Range models

2. Application



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Lauren Buckley

buckley@bio.unc.edu

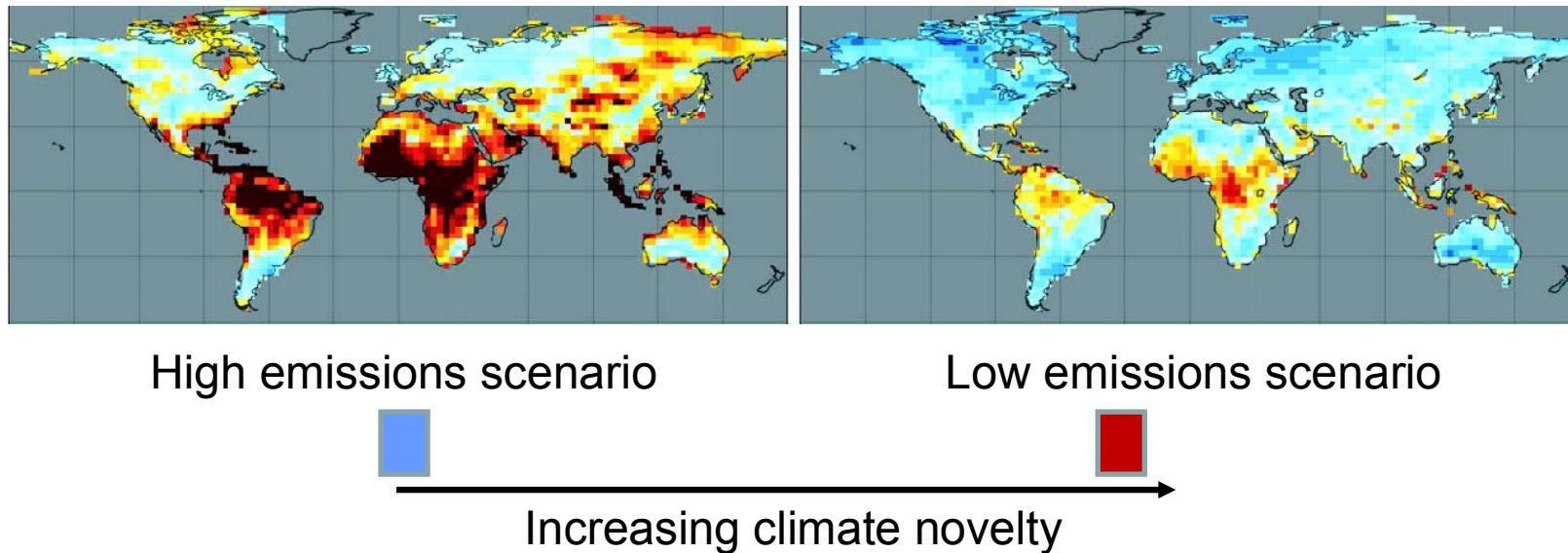
<http://www.bio.unc.edu/Faculty/Buckley/Lab/>



THE UNIVERSITY
of NORTH CAROLINA
at CHAPEL HILL

Climate-induced range shifts and extinctions?

Novel Climates by 2100



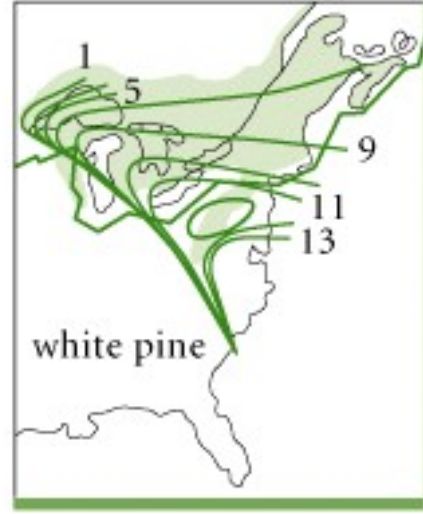
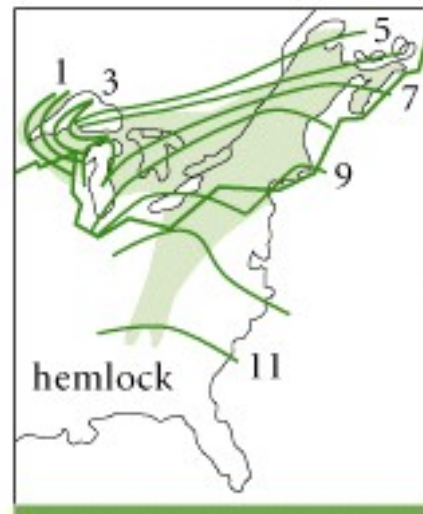
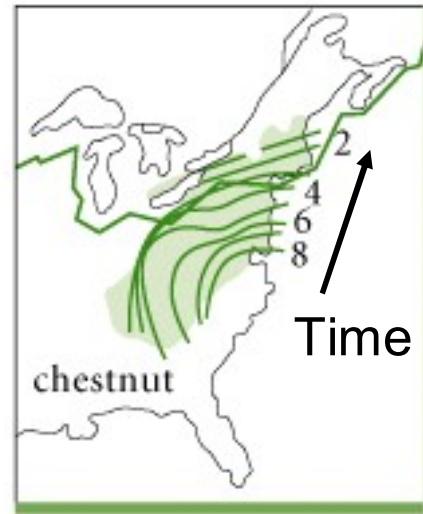
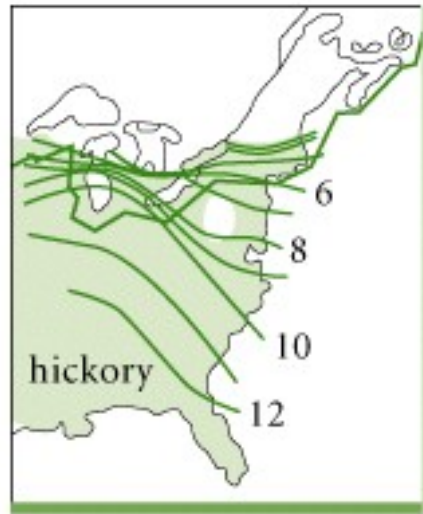
(Williams *et al.* 2007)



US mammals

Modern

Quaternary fossil
(Graham *et al.* 1996)



Trees
(Davis 1976)

Giant panda



Ferret



California condor



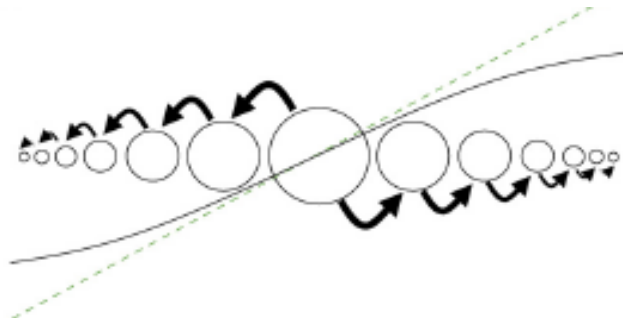
Whooping crane



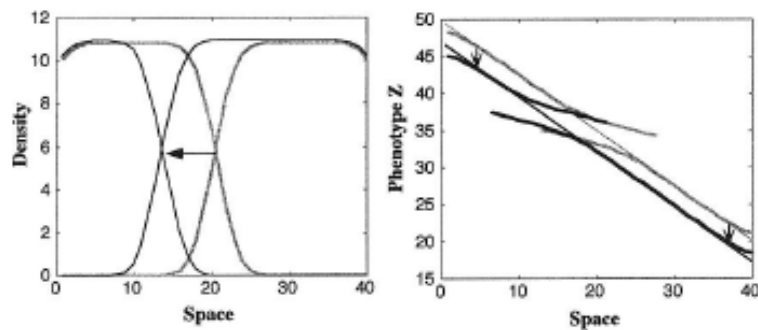
(Channell & Lomolino 2000)

Theory

Gene flow along selection gradient
(Kirkpatrick & Barton 1997)



Incorporate species interactions
(Case & Taper 2000)



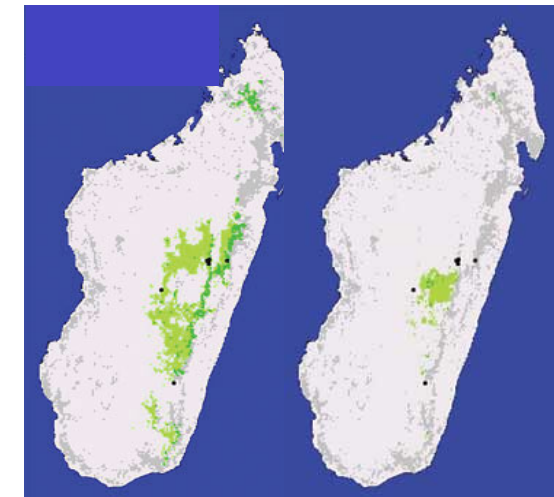
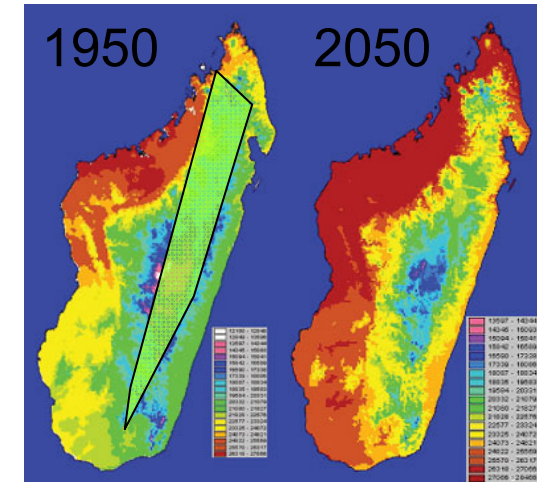
Practice: Correlative models

specimen localities



 Climatic envelope

Temperature



(Lees 2002)

Limitations of correlative models

- Other determinants of current range?
 - Extinctions?
- Nonlinear response to climate change?

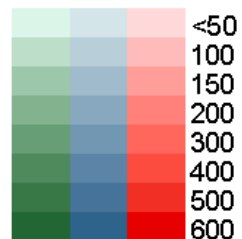
Trait variation and evolution

NORTH



Sceloporus undulatus
(Buckley 2008)

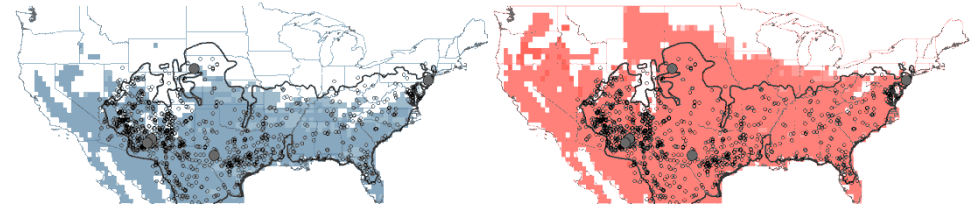
Lizards /1000m²



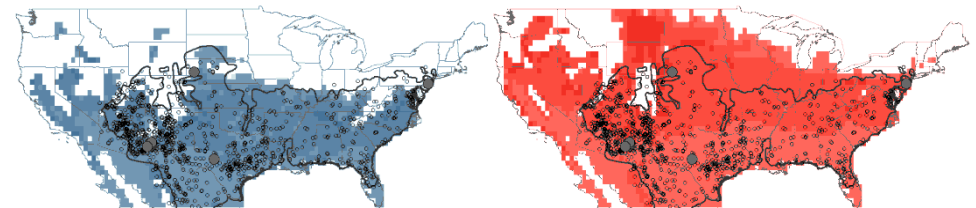
SOUTH

Nebraska traits

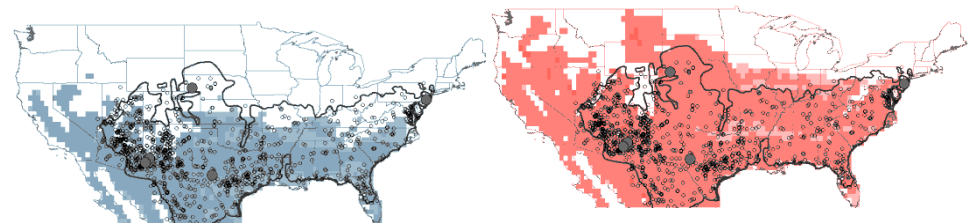
+ 3°C



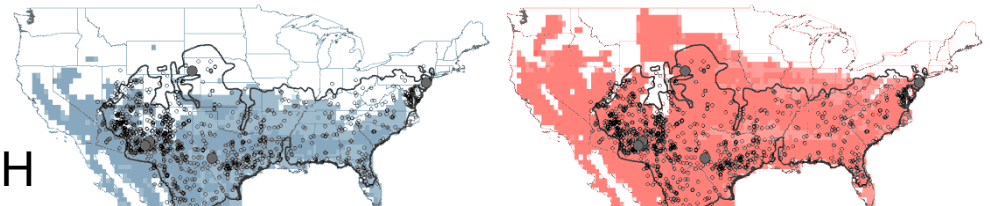
New Jersey traits



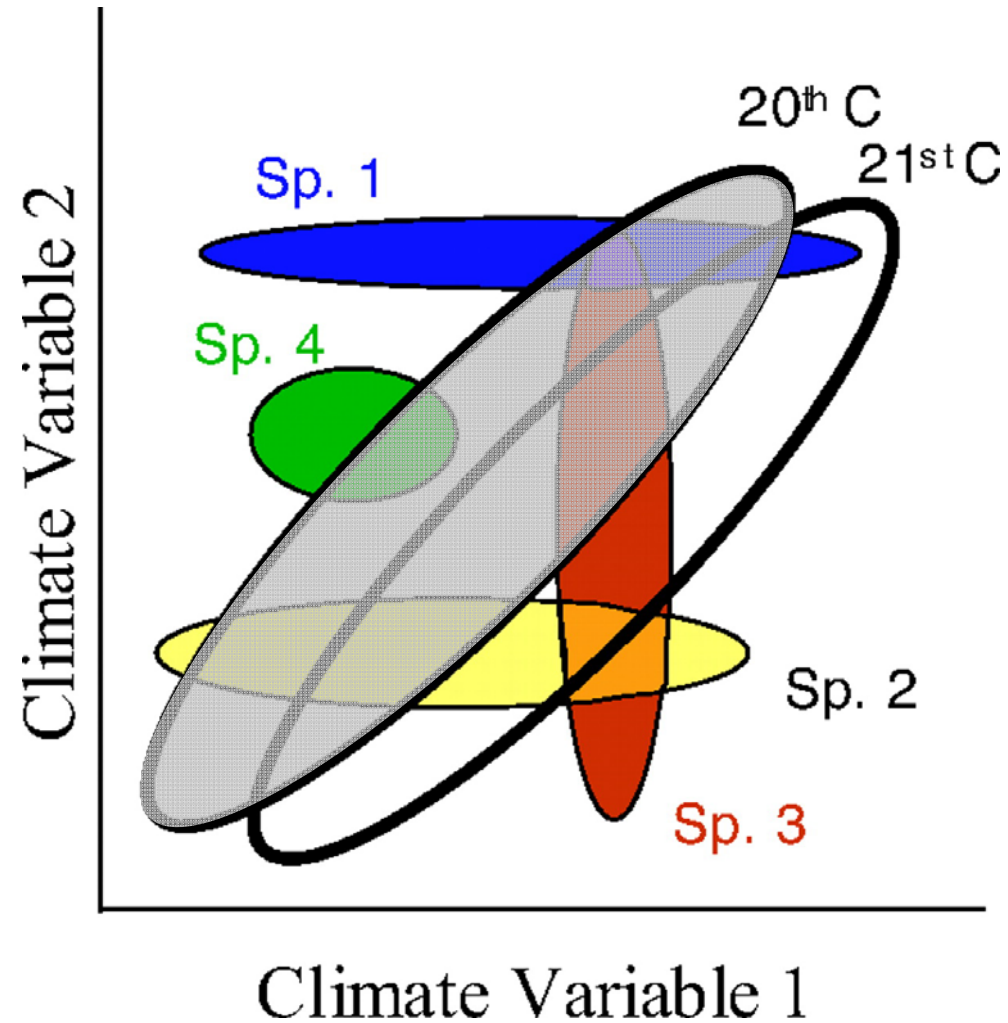
New Mexico traits



Texas traits

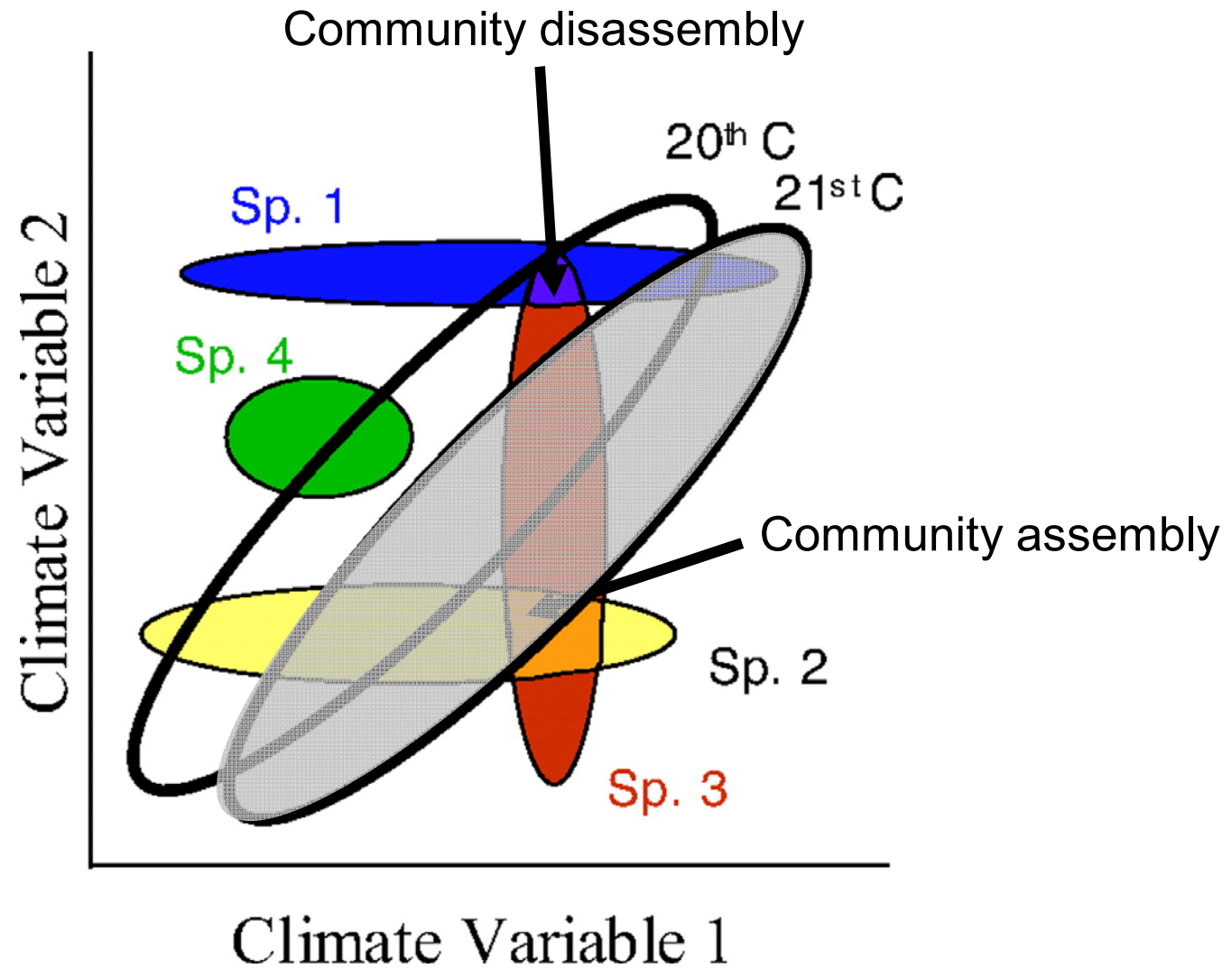


Species interactions



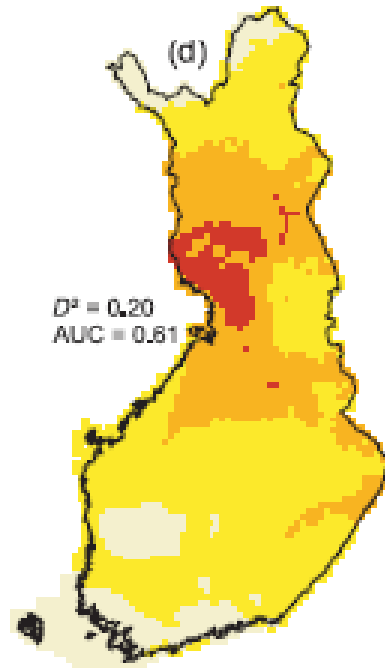
(Williams *et al.* 2007)

Species interactions

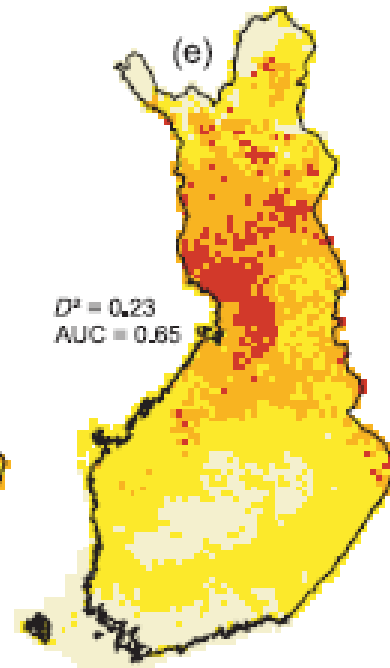


(Williams *et al.* 2007)

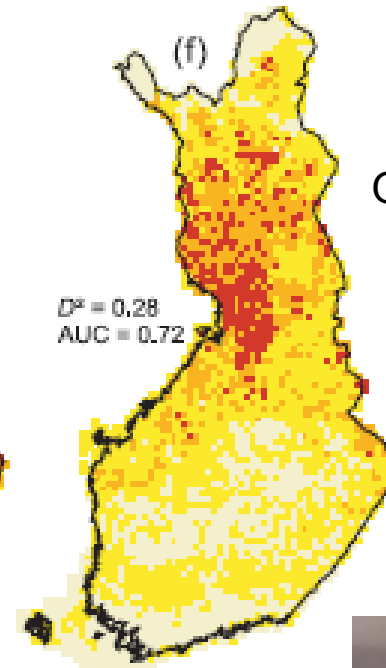
Climate



+habitat



+woodpeckers



Occurrence probability



Hawk owl
(Heikkinen *et al.* 2007)



Species Range Dynamics Working Group



MOVEMENT

- * Cost of movement
- * Dispersal barriers
- * Habitat quality
- * Spatial configuration of habitat



ENVIRONMENT



PHYSIOLOGY



DEMOGRAPHY

- * Downscaling climate models
- * Spatial and temporal variation in current and future environment
- * Translating environmental change into body temperature change

- * Acclimation and plasticity in physiology and behavior
- * Interactions of multiple changing environmental variables
- * Variation in environmental tolerance within and among populations
- * Hard climatic constraints verses cumulative effects

- * Growth, phenology, survival and recruitment
- * Metapopulation structure
- * Life stages
- * Mating
- * Time scales



BIOTIC INTERACTIONS

- * Loss and gain of competitors and predators
- * Interaction cascades
- * Phenological asynchrony
- * Disease
- * Hosts, mutualists, pollinators
- * Differential importance of biotic processes across range

(assumes no evolution)

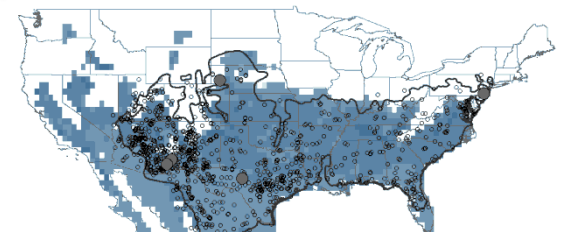
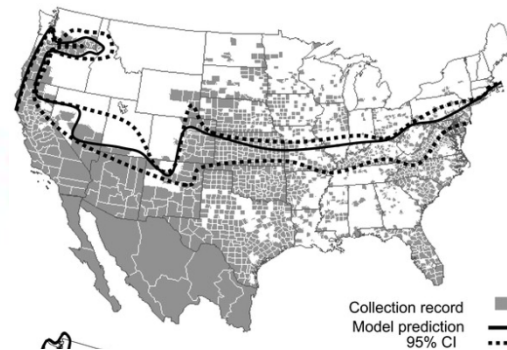
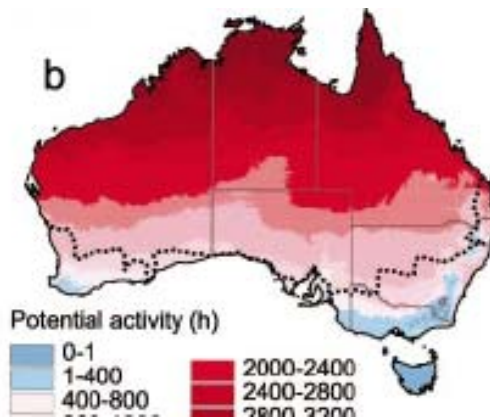
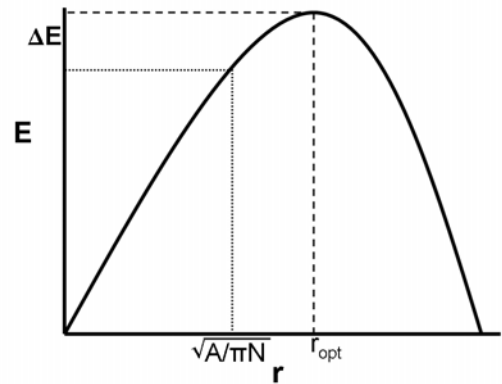
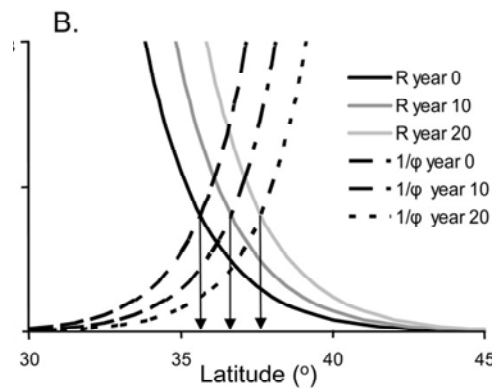
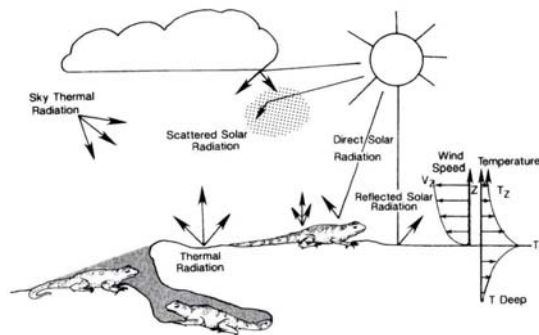
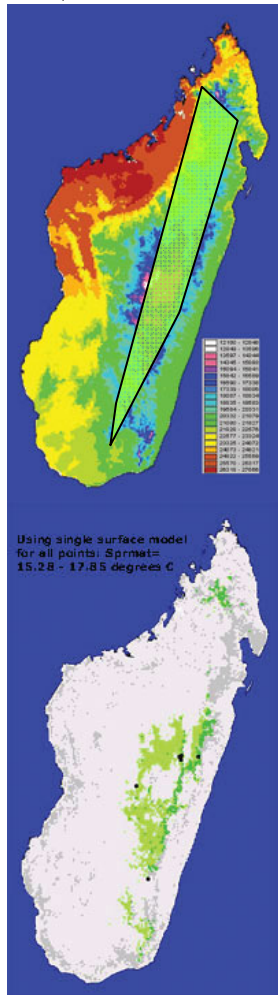
Range models

correlative
models

biophysical
models

performance
curves

foraging energetic
models



(Kearney & Porter 2004)

(Crozier & Dwyer 2006)

(Buckley 2008)



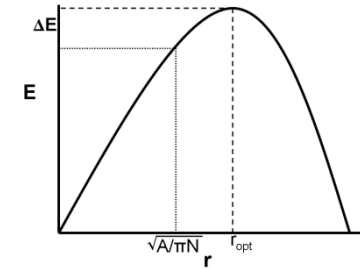
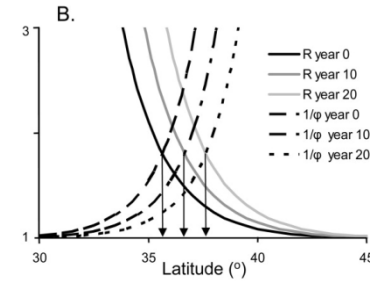
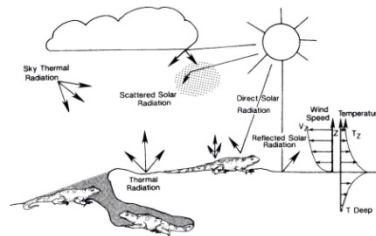
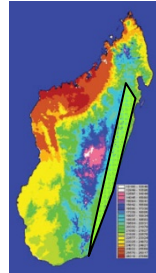
Range models

correlative
models

biophysical
models

performance
curves

foraging energetic
models



Data

localities

traits, thresholds

demography

traits

Demography

X

X

Geographic
Variation

X

X

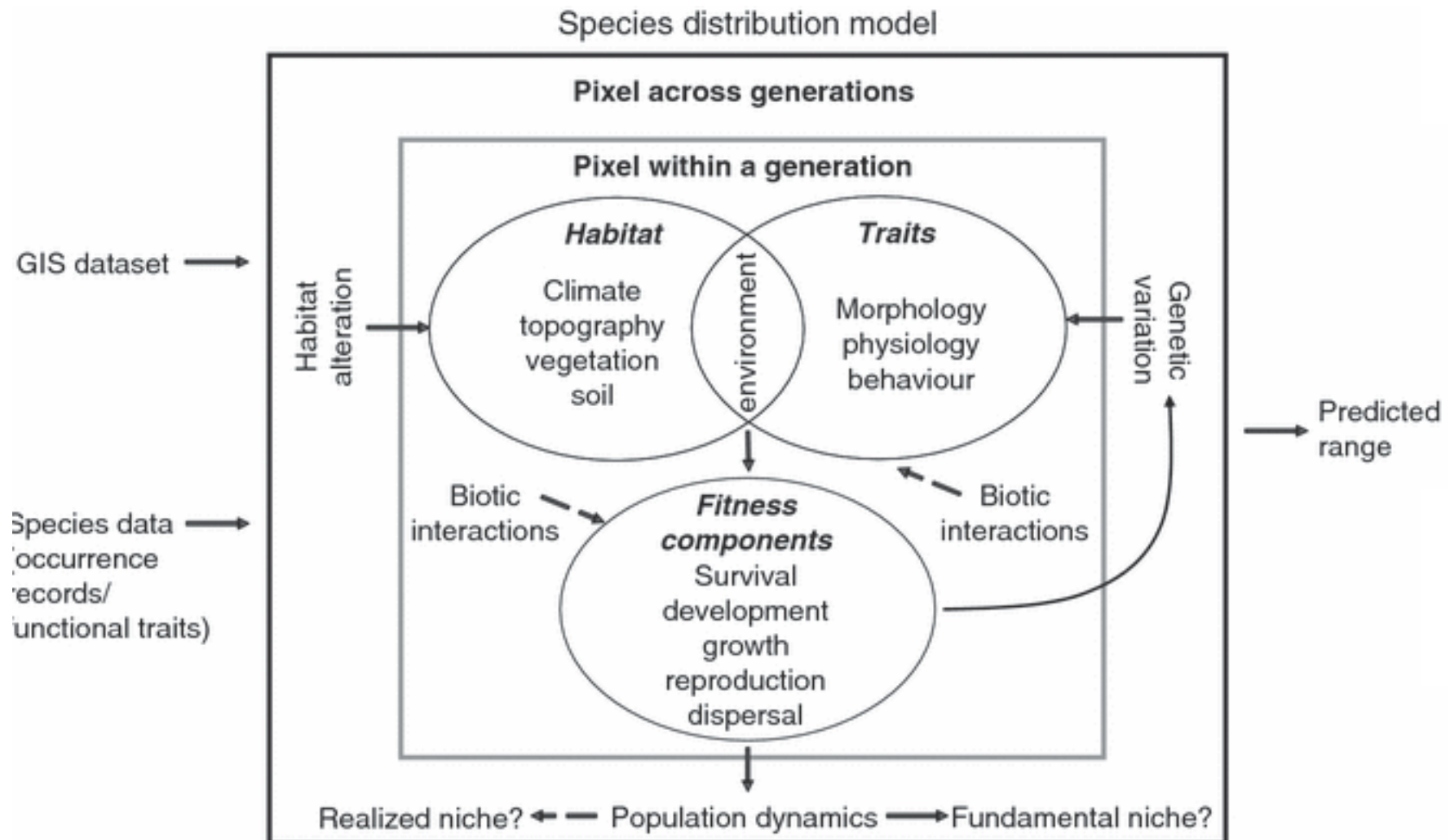
Biotic
Constraints

implicit

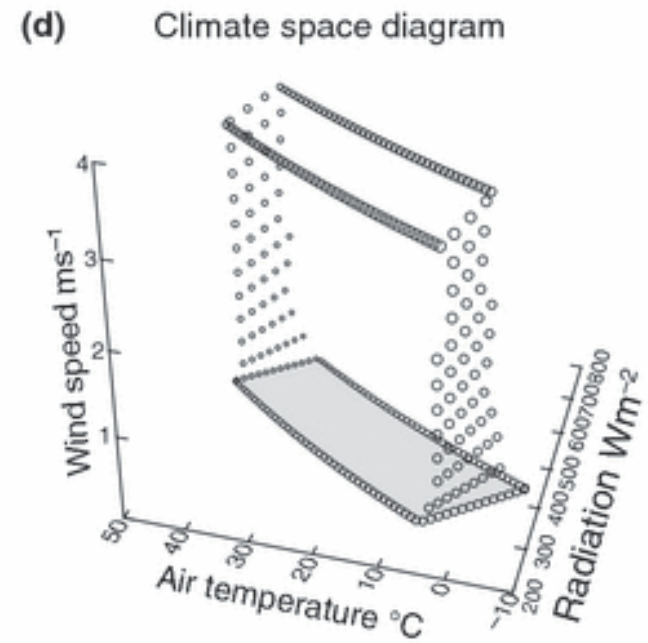
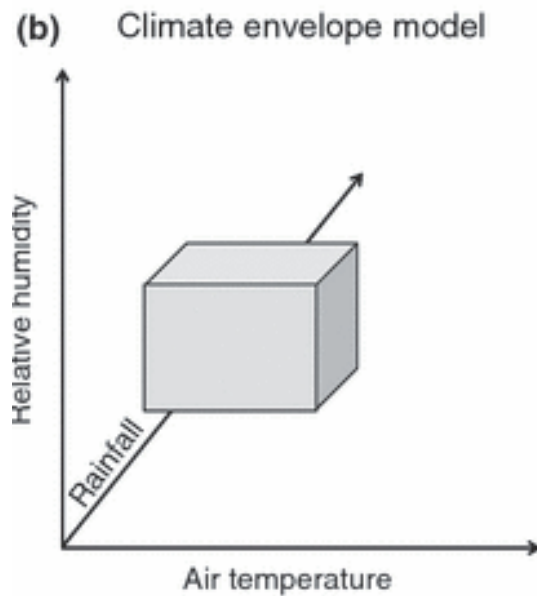
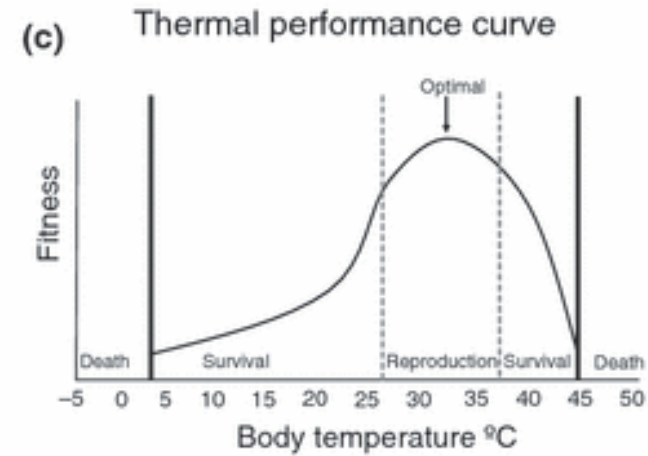
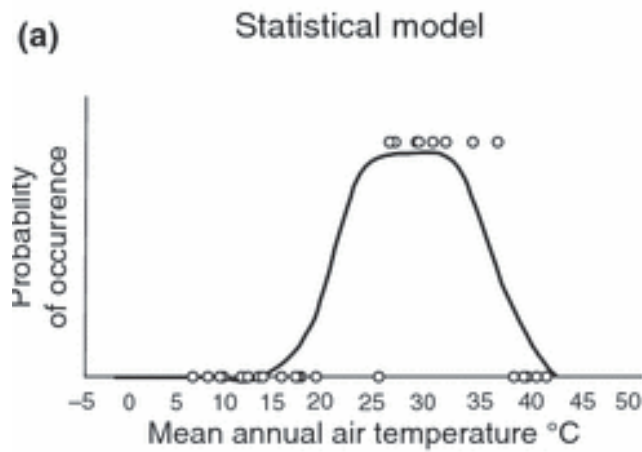
difficult

possible

possible



(Kearney and Porter 2009)



(Kearney and Porter 2009)

Model Comparison

correlative

biophysical
models

performance
curves

foraging energetic
models



Model species

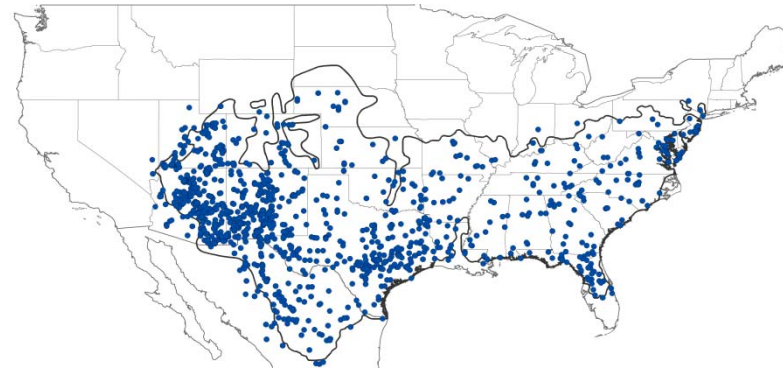
Fence lizard

Sceloporus undulatus



Distribution data

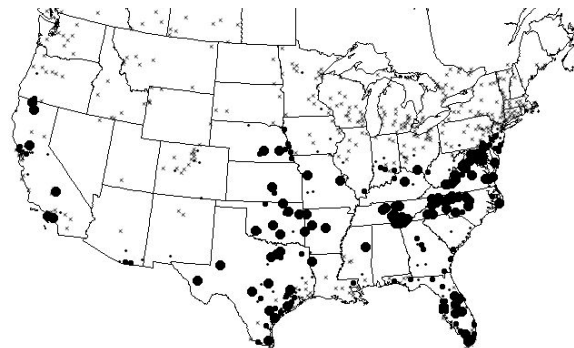
HerpNET



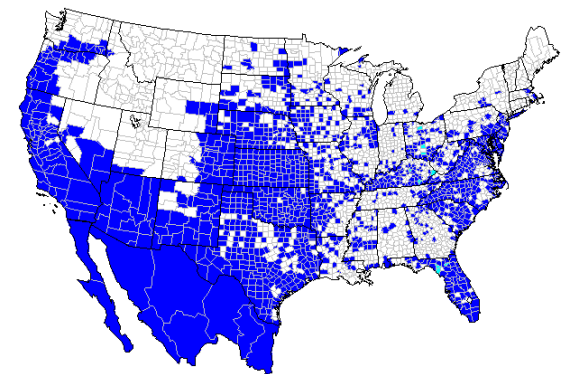
Sachem skipper butterfly
Atalopedes campestris



4th of July Butterfly counts



Butterflies and Moths of NA



Environmental data

10' resolution, New *et al.* 2002



Fence lizard range models

correlative
models

biophysical
models

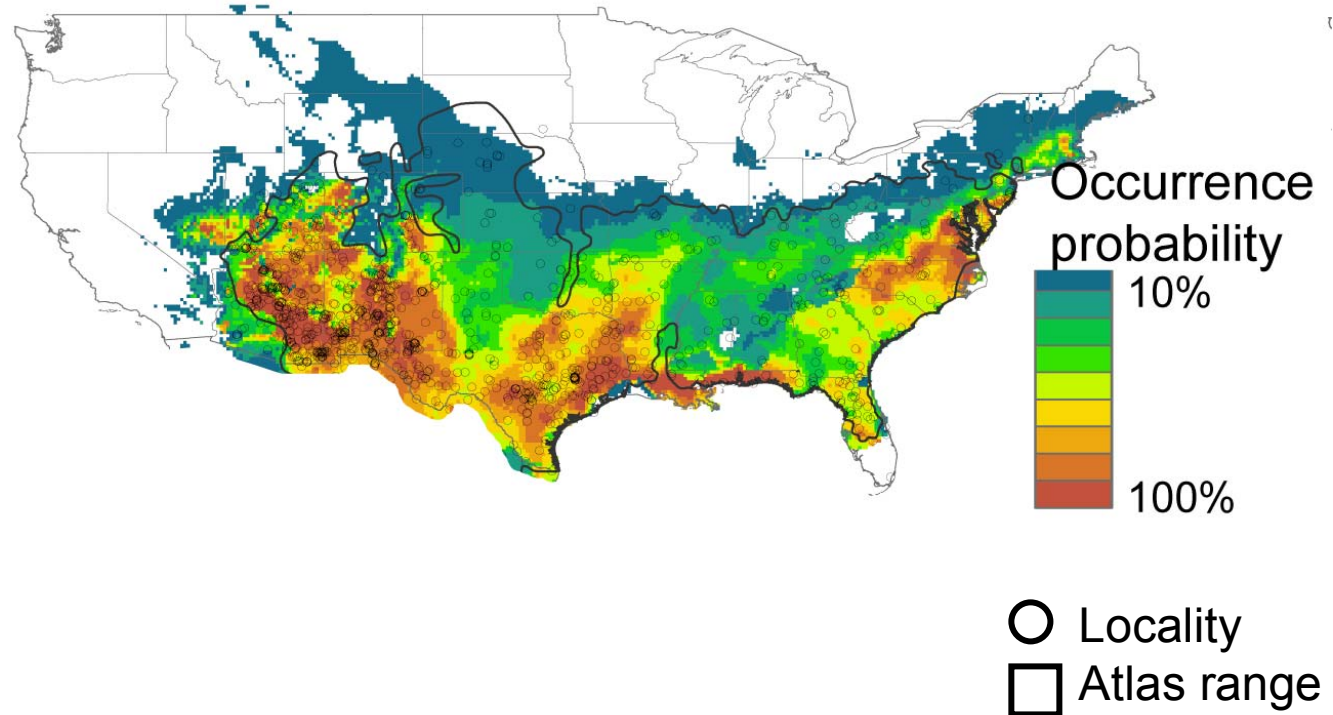
performance
curves

foraging energetic
models



Max ent

10% threshold



(NESCent/ NCEAS WG)



Sachem skipper range models

correlative
models

biophysical
models

performance
curves

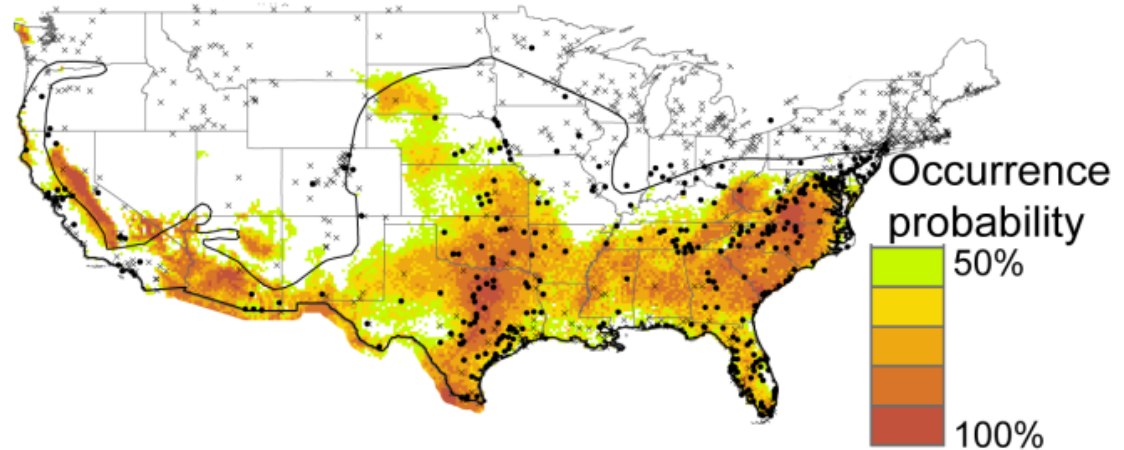
foraging energetic
models



GLM

50% Threshold

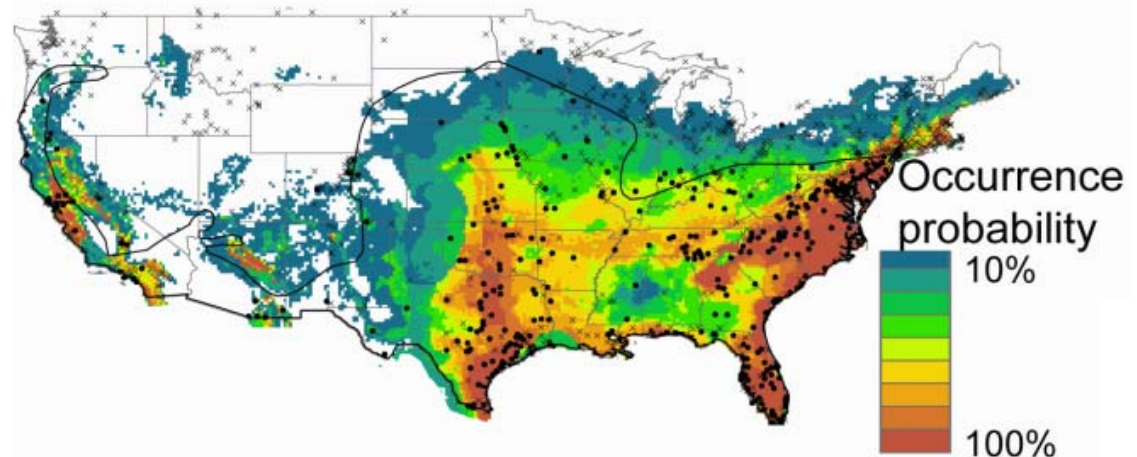
CORRELATIVE- GLM



MaxEnt

10% Threshold

CORRELATIVE- Maxent





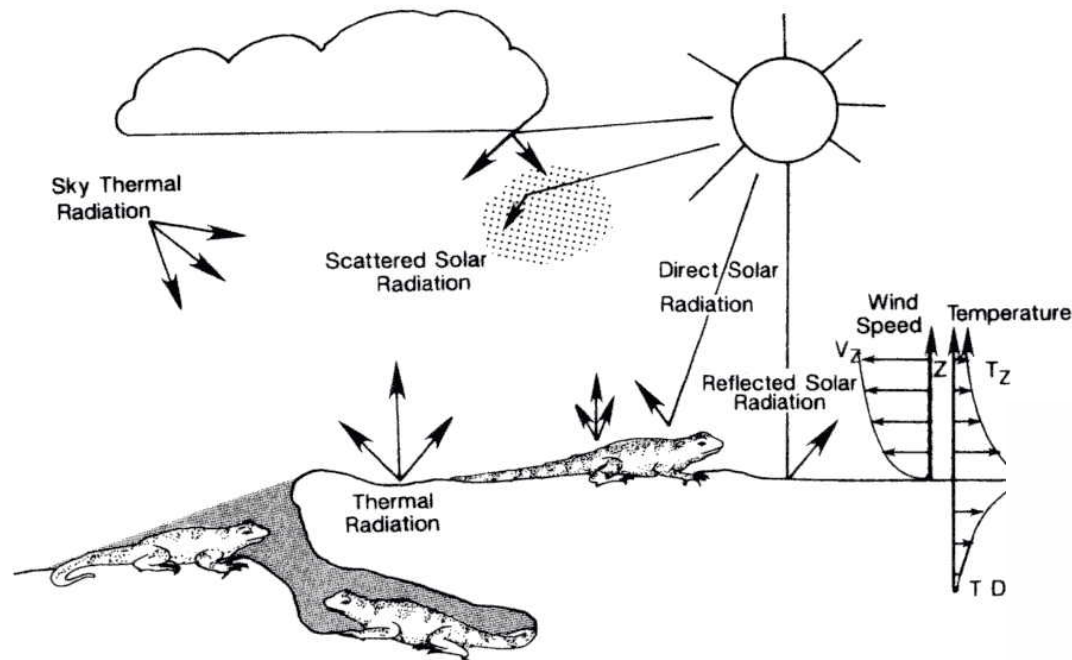
Fence lizard range models

correlative
models

biophysical
models

performance
curves

foraging energetic
models



Q = Heat
n = Mass
F = Food
W = Water
I = Ingested
D = Defecated
A = Absorbed
U = Urine
G = Growth
R = Reproduction
S = Stored

$$\begin{aligned}
 Q_{\text{SOLAR}} + Q_{\text{IR,in}} + m_{\text{O}_2} + \text{Work} \\
 m_{\text{F,I}} - m_{\text{F,D}} = m_{\text{F,A}} = \frac{Q}{m} + m_{\text{F,G}} + m_{\text{F,R}} + m_{\text{F,S}} \\
 \text{METAB} \\
 m_{\text{F,CO}_2} = Q_{\text{IR,out}} + Q_{\text{CONV}} \\
 + m_{\text{F,NH}_3} + m_{\text{F,W}} \\
 m_{\text{W,I}} - m_{\text{W,D}} = m_{\text{W,A}} + m_{\text{F,W}} = \frac{Q}{m} + m_{\text{W,U}} + m_{\text{W,S}} \\
 + Q_{\text{COND}} + Q_{\text{S}}
 \end{aligned}$$

(Grant & Porter 1992)



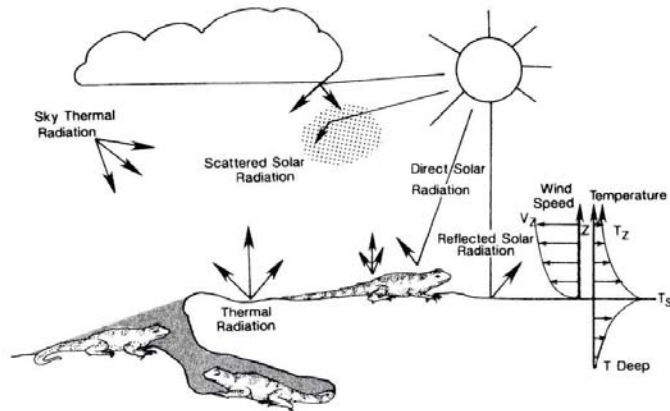
Fence lizard range models

correlative
models

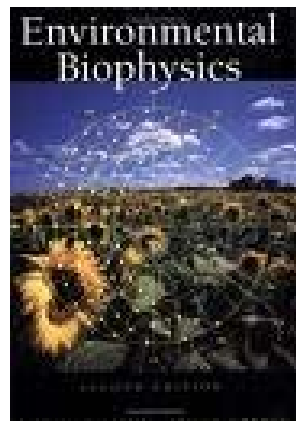
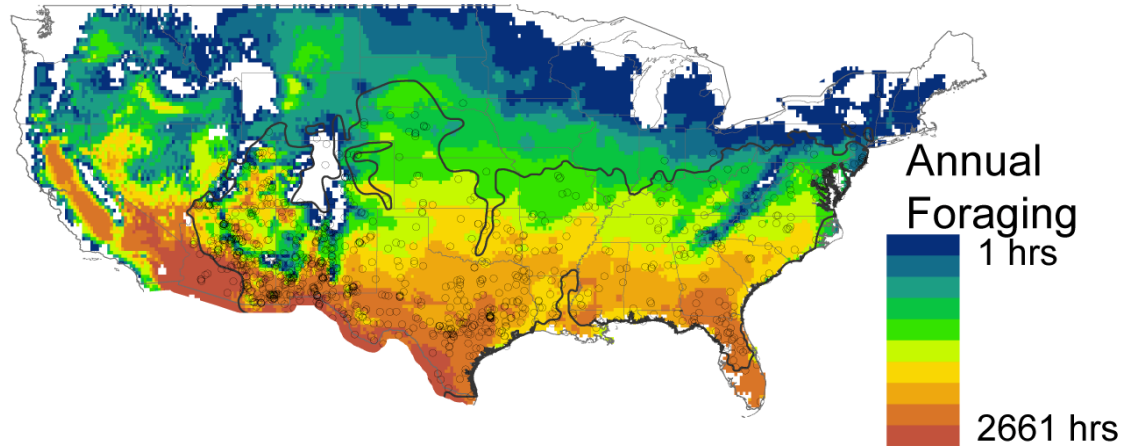
biophysical
models

performance
curves

foraging energetic
models



(Grant & Porter 1992)



(Campbell and Norman 1988
Implemented by Buckley 2008)



1315 hr/yr threshold



Sachem skipper range models

correlative
models

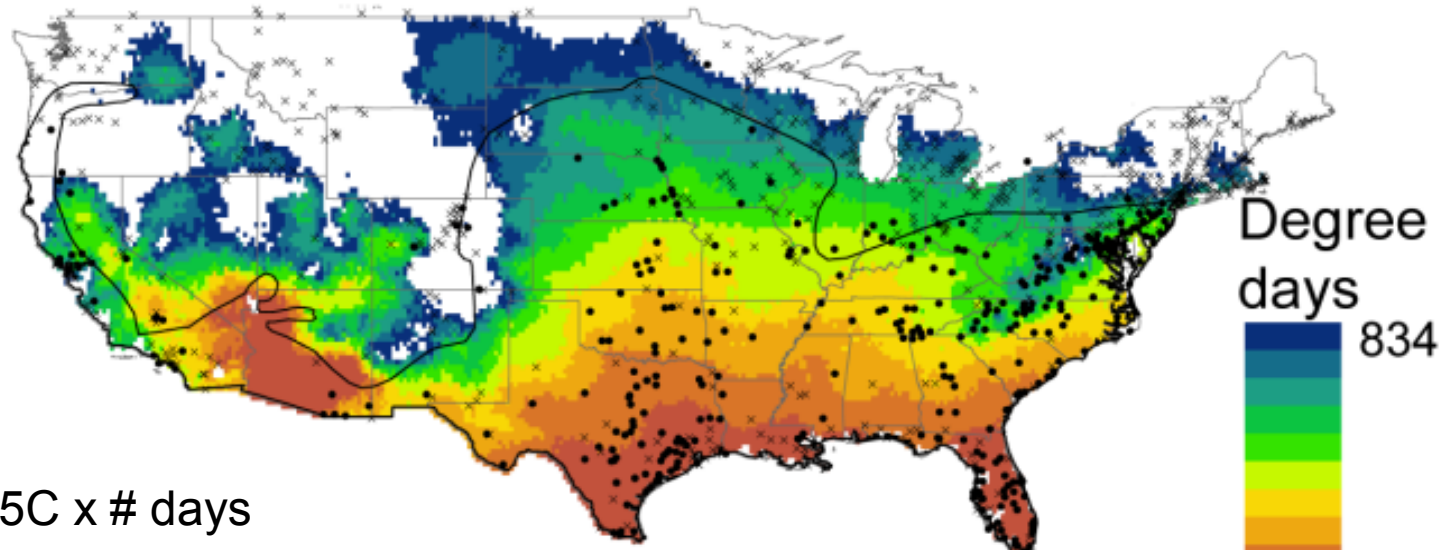
biophysical
models

performance
curves

foraging energetic
models



BIOPHYSICAL THRESHOLD



degrees over 15.5C x # days



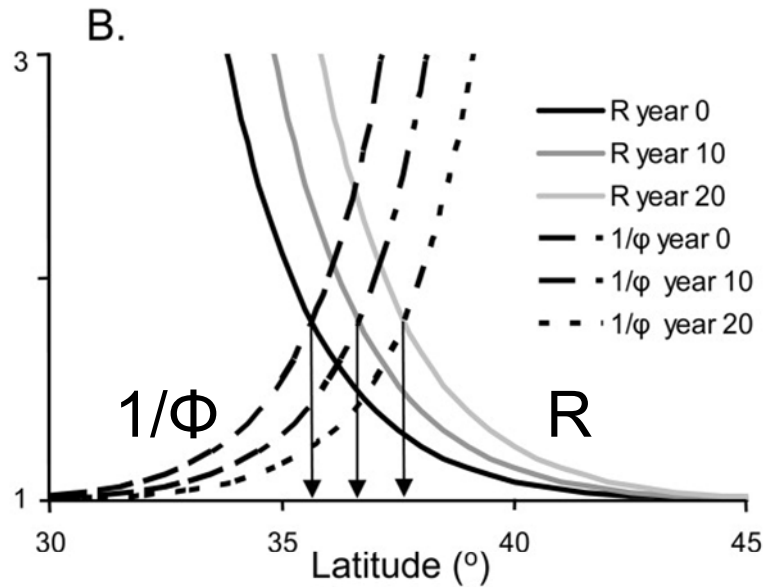
Sachem skipper range models

correlative
models

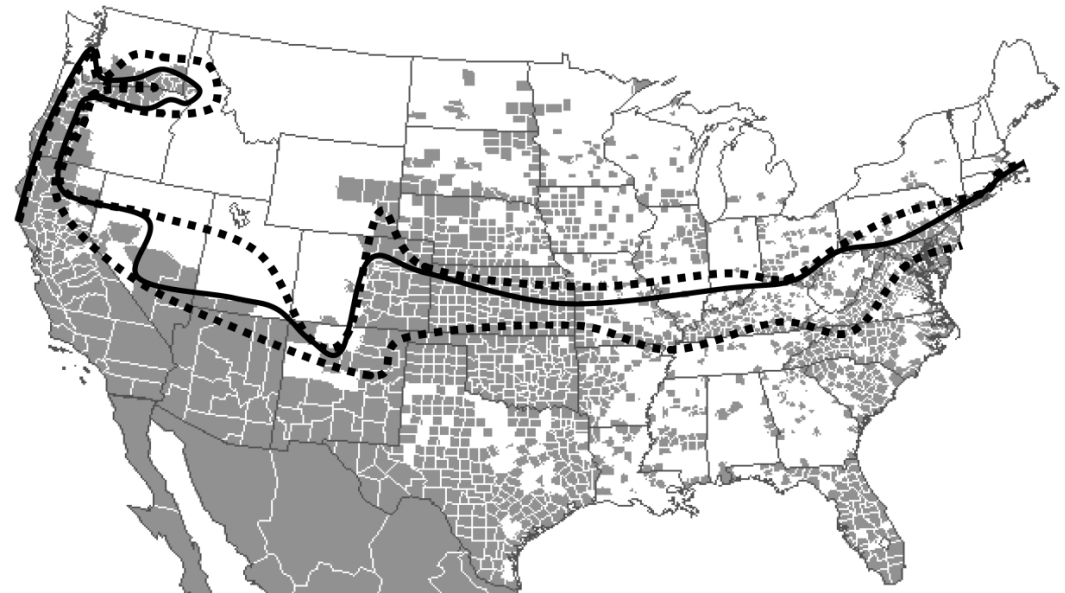
biophysical
models

performance
curves

foraging energetic
models



R- net summer recruitment
 Φ - winter survival



■ Collection record
 — Model prediction
 - - 95% CI

(Crozier & Dwyer 2006)



Sachem skipper range models

correlative
models

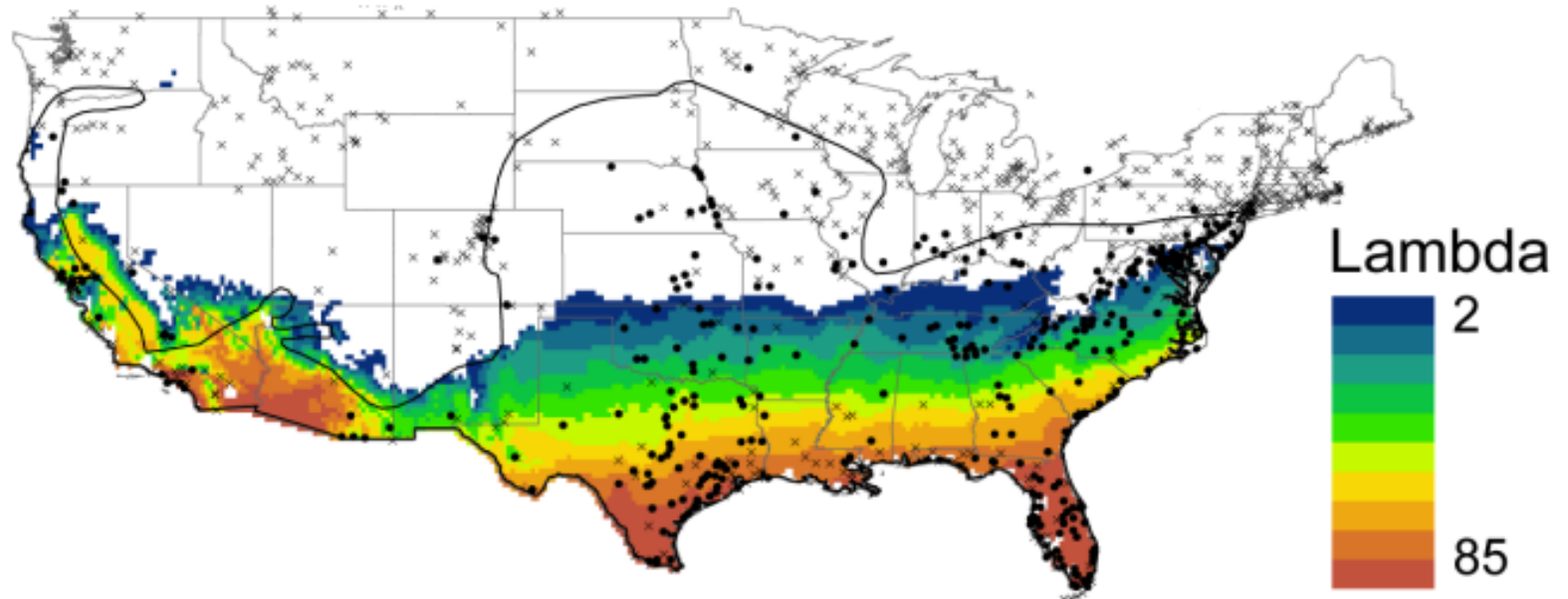
biophysical
models

performance
curves

foraging energetic
models



LIFE HISTORY





Fence lizard range models

correlative
models

biophysical
models

performance
curves

foraging energetic
models



net reproductive
rate

sum annually
maturity to death

$$R_0 = \sum_{x=\alpha}^L S_j S_a(T)^{(x-\alpha)} m(T)$$

juvenile and adult (T)
survival

annual
fecundity (T)



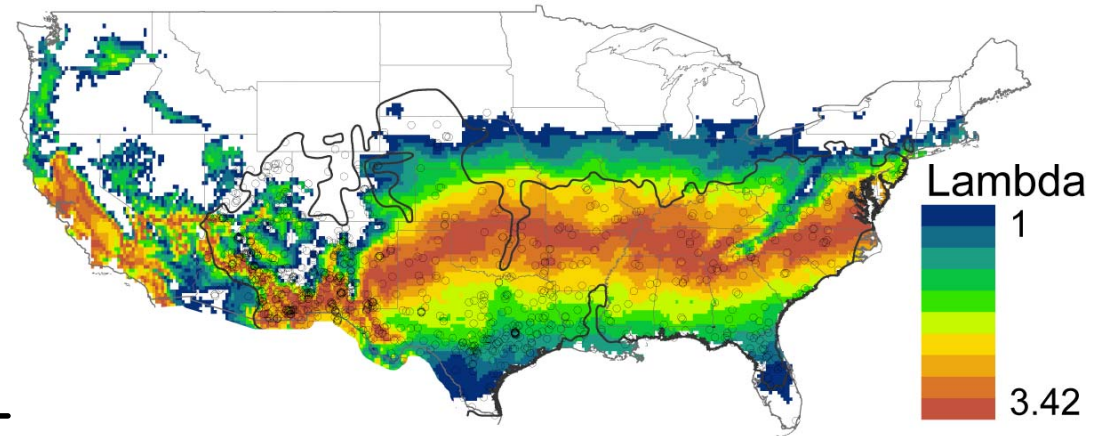
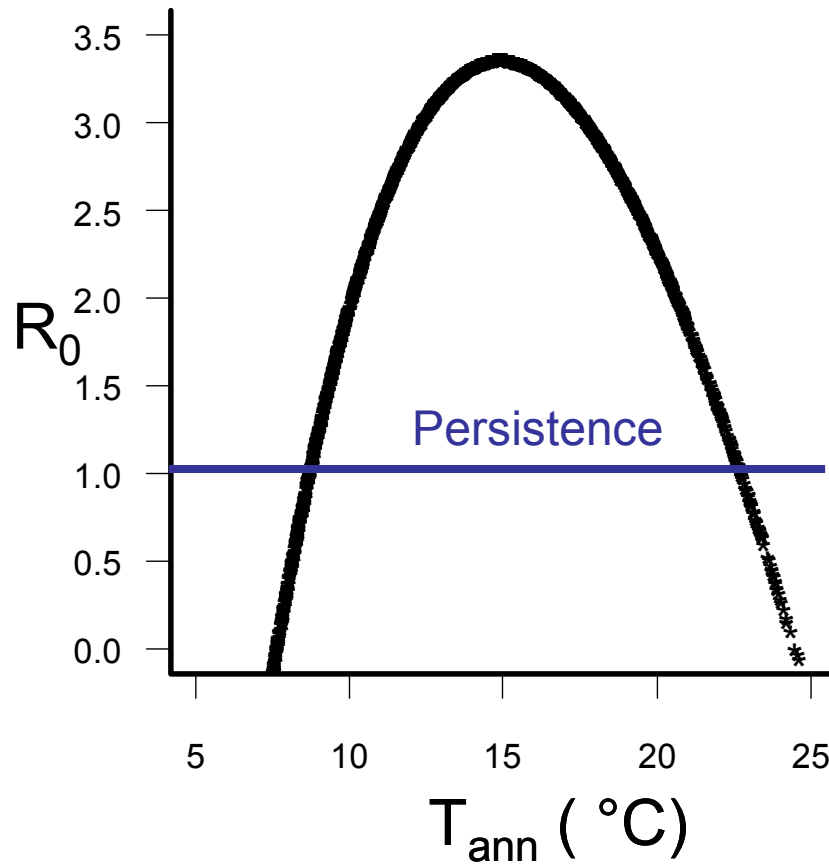
Fence lizard range models

correlative
models

biophysical
models

performance
curves

foraging energetic
models





Fence lizard range models

correlative
models

biophysical
models

performance
curves

foraging energetic
models



environmental
conditions



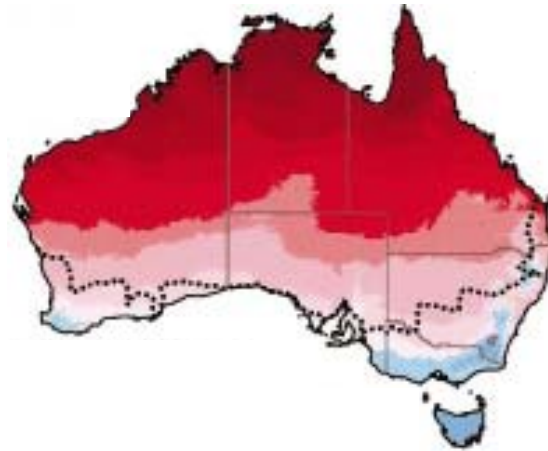
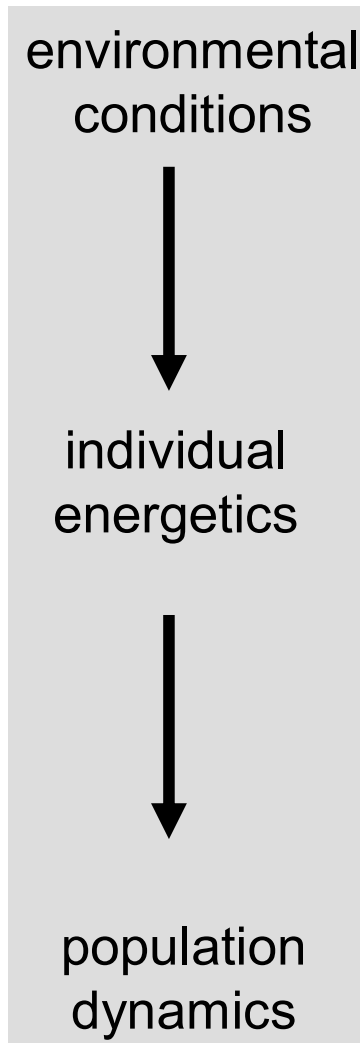
individual
energetics



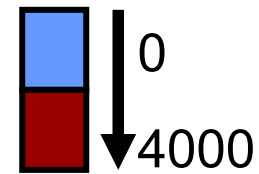
population
dynamics

can extend to include biotic factors
(adaptation, movement, species' interactions, ...)

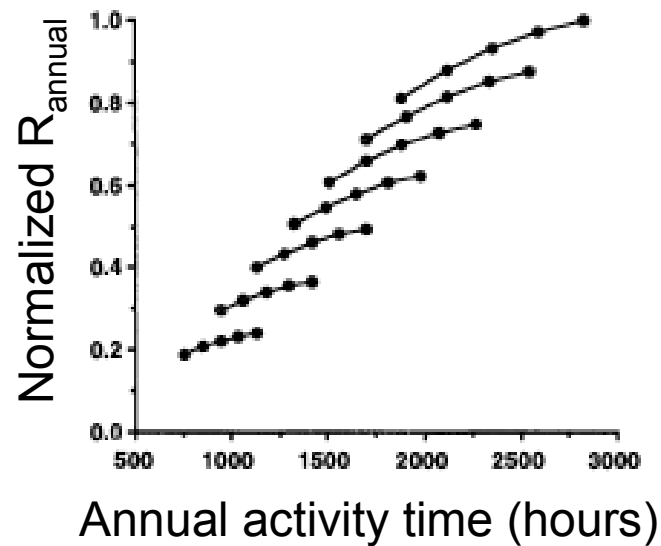
Mechanistic range models



Potential activity (h)



Gecko
(Kearney & Porter 2004)



Fence lizard
(Adolph & Porter 1993)

Dynamic foraging energetic model

Foraging duration & performance

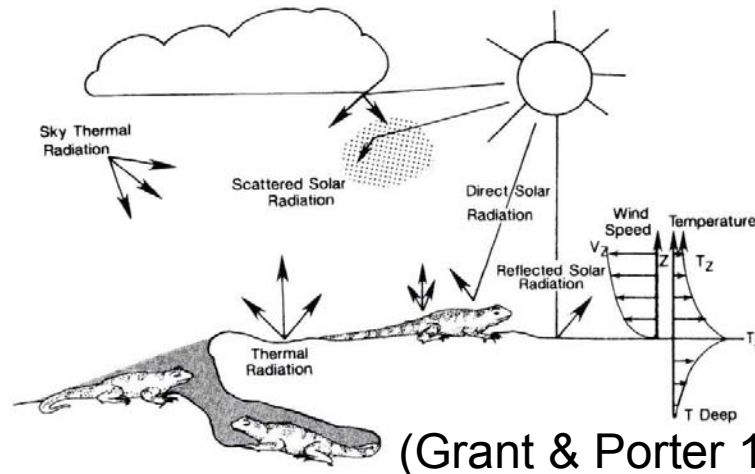
environmental
conditions



individual
energetics



population
dynamics



PARAMETERS

Body temperature range, T_b

Body size → Metabolic rate

Sprint speed

Prey size

Gut capacity

Environmental temperature, T_e (Campbell & Norman 2000)

Assume forage when T_e within 20-80% quantiles of T_b

Dynamic foraging energetic model

Foraging duration & performance

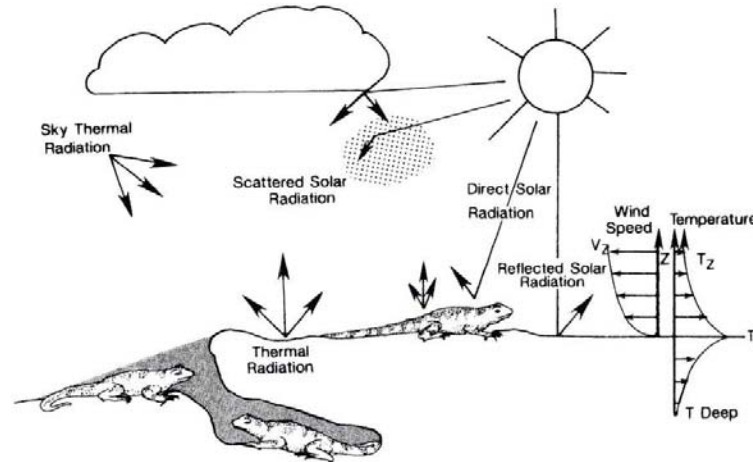
environmental
conditions



individual
energetics



population
dynamics



PARAMETERS

Body temperature range, T_b
Body size $\rightarrow$ Metabolic rate
Sprint speed
Prey size
Gut capacity

Optimal foraging model

Optimize energetic yield / time

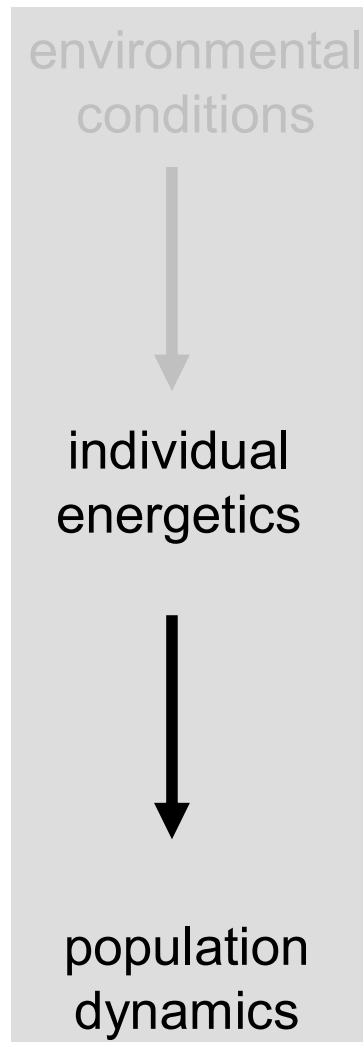
$$E(r) = \frac{\text{energy intake} - \text{pursuit costs}(r) - \text{waiting costs}(r)}{\text{pursuit time}(r) + \text{waiting time}(r)}$$

$r \rightarrow$ foraging radius

PARAMETERS

Insect abundance

Dynamic foraging energetic model



Population dynamic model

$$\Delta N = \underbrace{[bE(r) t_f]}_{\text{Birth}} - \underbrace{[c(t-t_f) - m]}_{\text{Death}} N$$

$t_f \rightarrow$ foraging duration

$t \rightarrow$ total time

$c \rightarrow$ costs (offspring) for maintenance while not foraging

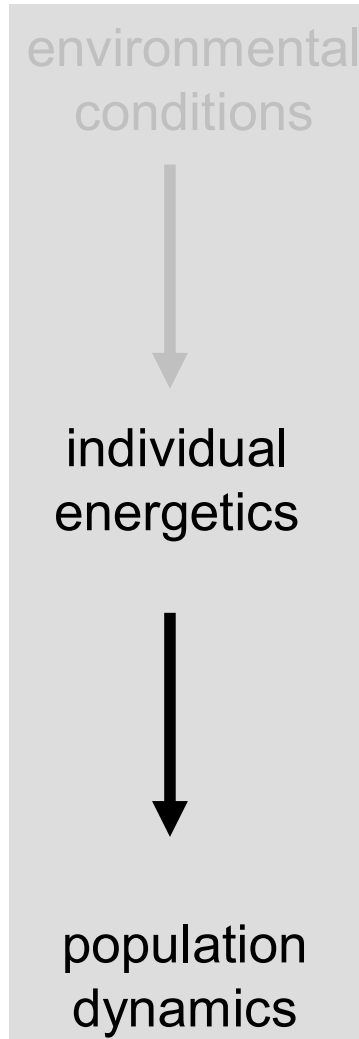
PARAMETERS

$b \rightarrow$ converts energy into offspring
(eggs/J x S_{maturity})

$m \rightarrow$ mortality

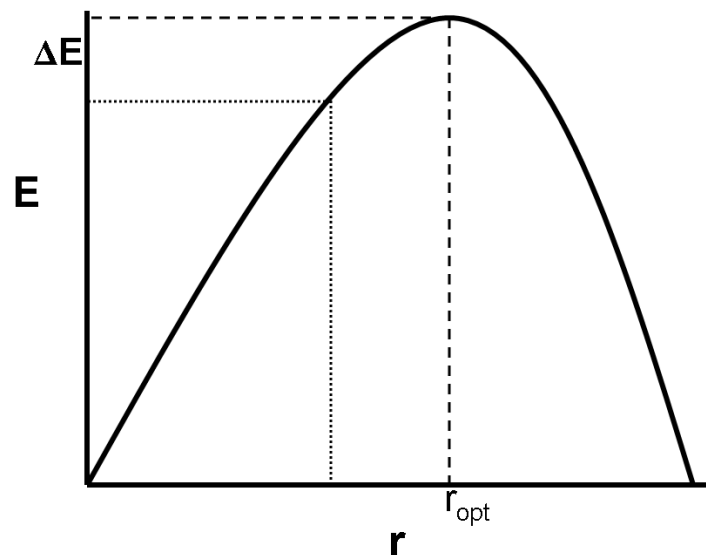
(Buckley & Roughgarden JAE 2005)

Dynamic foraging energetic model



Population dynamic model

$$\Delta N = [bE(r) t_f - c(t - t_f) - m]N$$



$t_f \rightarrow$ foraging duration

$t \rightarrow$ total time

$c \rightarrow$ costs (offspring) for maintenance while not foraging

PARAMETERS

$b \rightarrow$ converts energy into offspring
(eggs/J $\times S_{\text{maturity}}$)

$m \rightarrow$ mortality

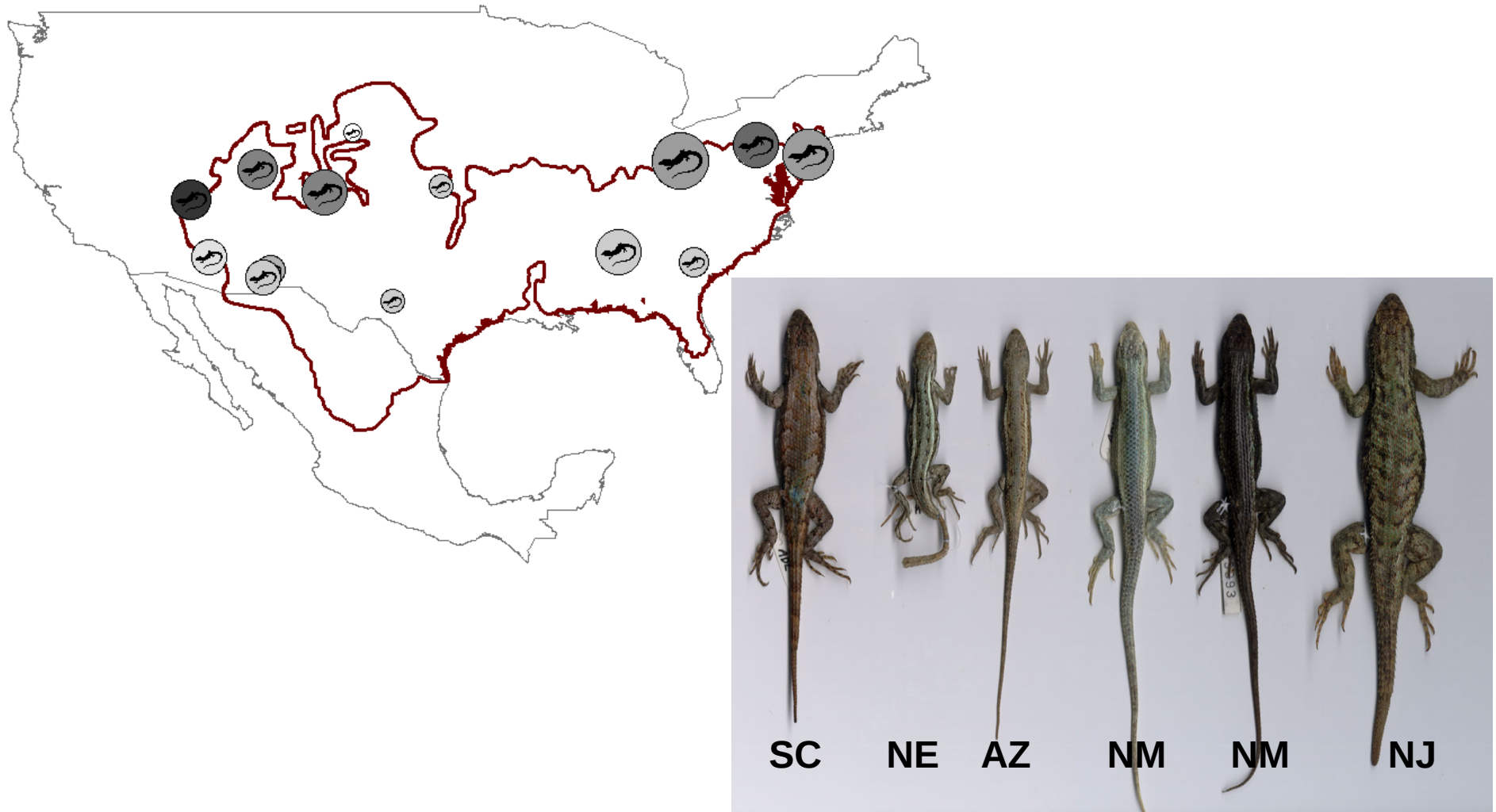
Density-dependence in $E(r)$

Can solve for equilibrium population size

Abundance patterns of solitary and sympatric anoles along elevation gradients?

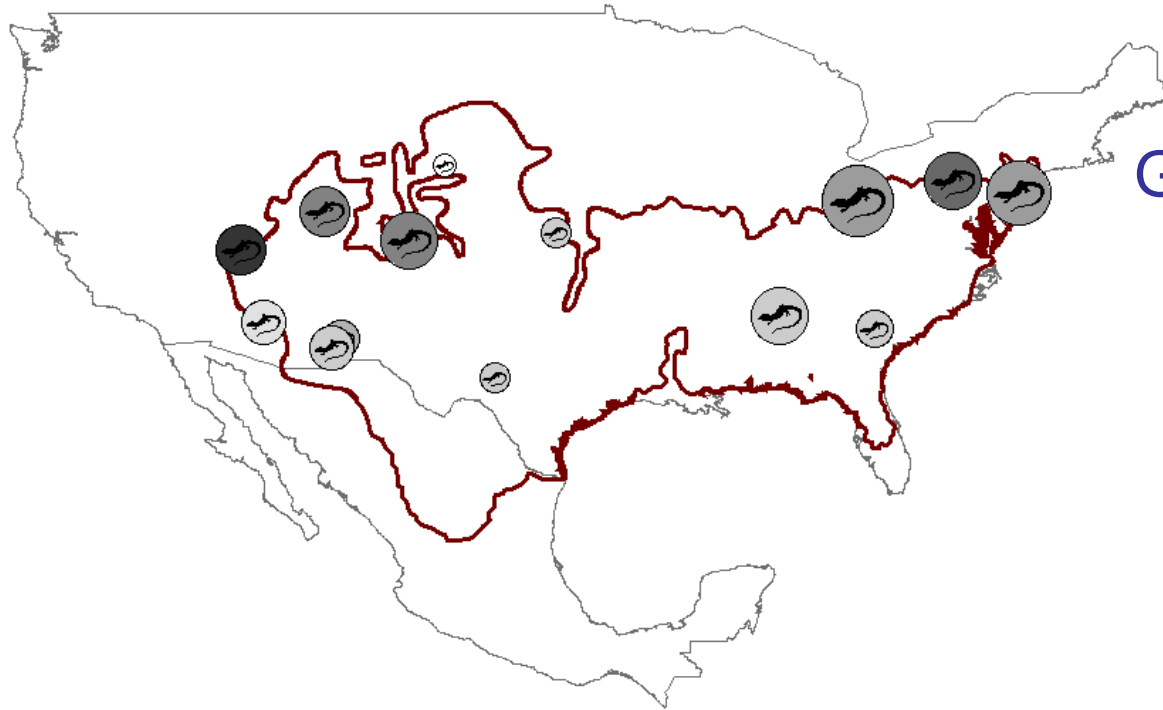


Does biology matter when predicting climate change responses?



Sceloporus undulatus
(Leaché and Reeder 2002)

Does biology matter when predicting climate change responses?

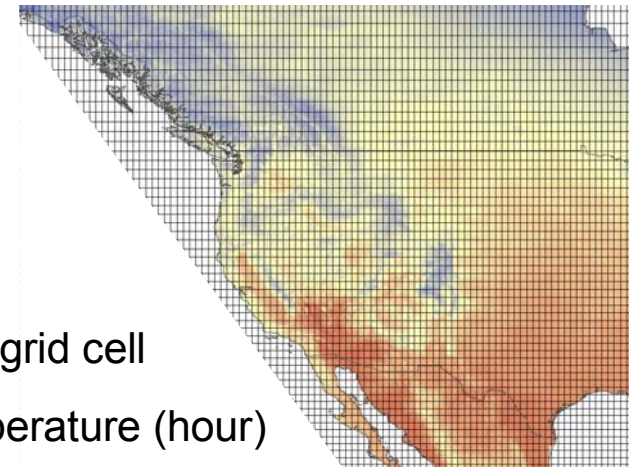


Geographic variation in
Size (SVL)
Survival
Development rate

Insect abundance

Assumes no trait plasticity

0.5° grid cell
temperature (hour)

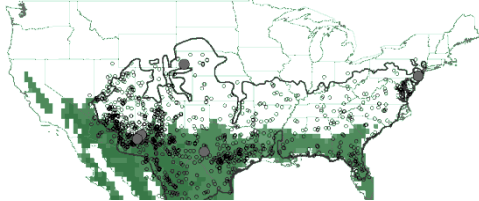


(Buckley AmNat 2008)

Dynamic foraging energetic model

Population SVL

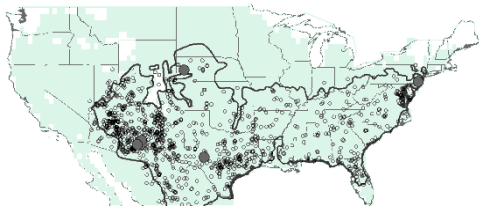
Nebraska traits



$\frac{SVL}{(mm)}$ $\frac{Age_{mat}}{(mo)}$ $\frac{S_{annual}}{(\%)}$

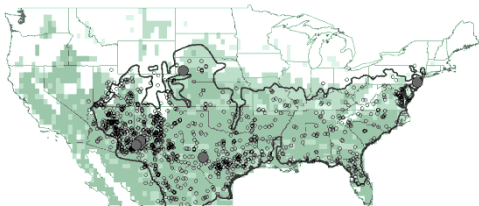
55

New Jersey traits



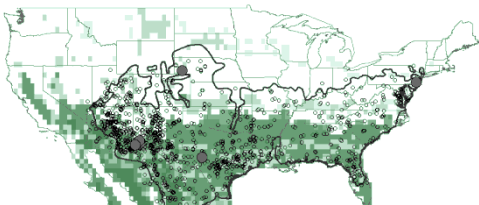
73

New Mexico traits



63

Texas traits



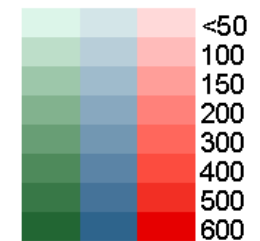
57

NORTH



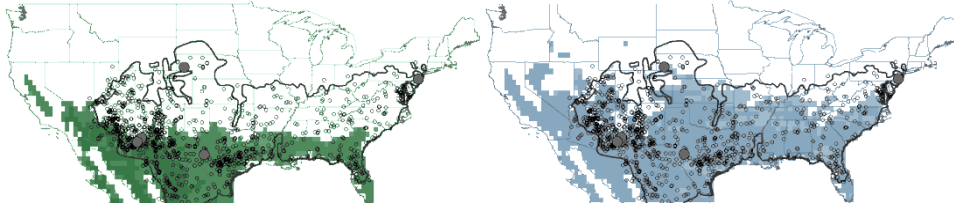
SOUTH

Lizards /1000m²



Population SVL + Population Life History

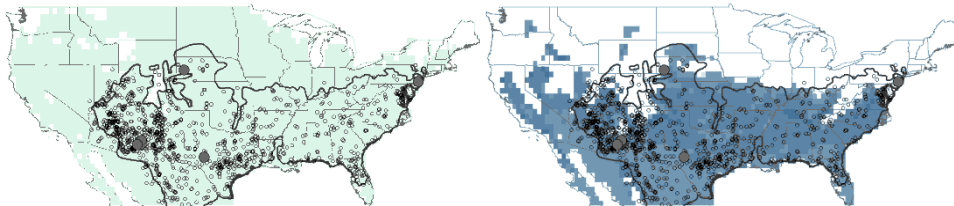
Nebraska traits



SVL
(mm) Age_{mat}
(mo) S_{annual}
(%)

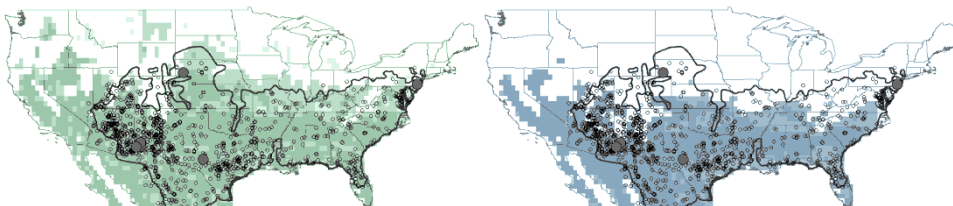
55 10 0.33

New Jersey traits



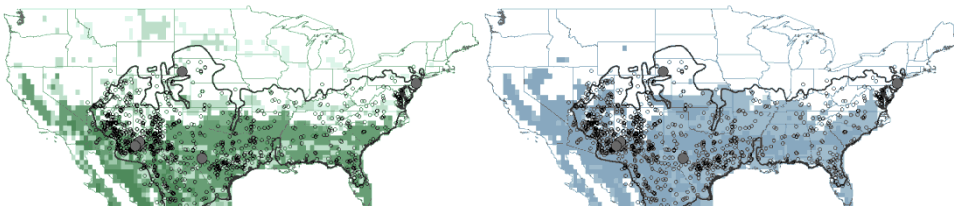
73 20 0.44

New Mexico traits



63 18 0.32

Texas traits



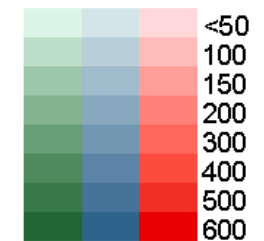
57 12 0.11

NORTH



SOUTH

Lizards /1000m²

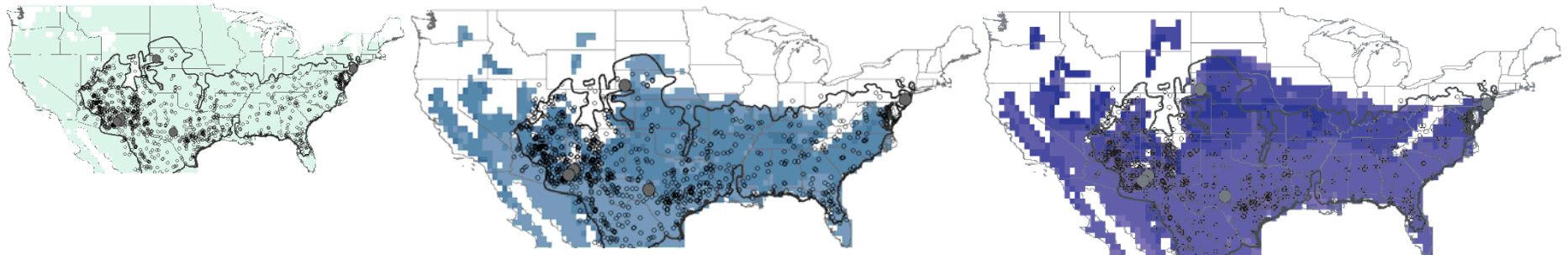


Population SVL

+ Population Life History

+ Population Insect Abundance

New Jersey traits

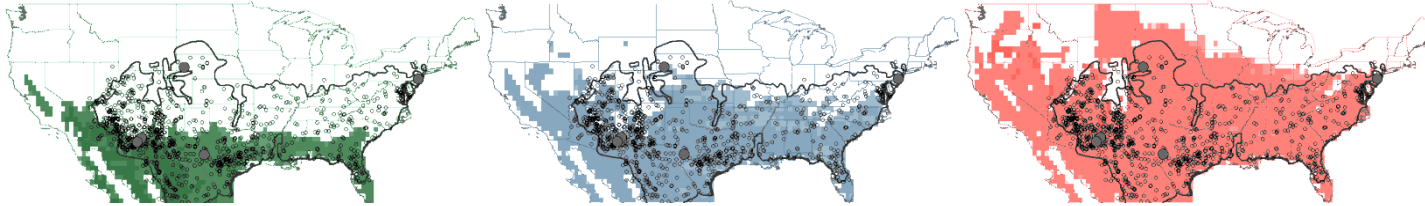


Population SVL

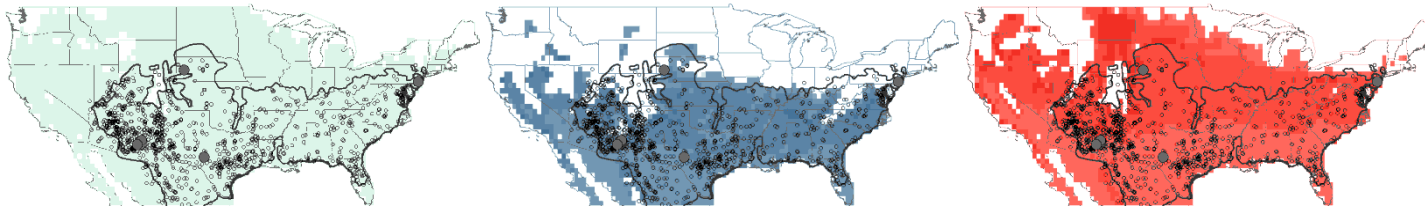
+ Population Life History

+ 3°C

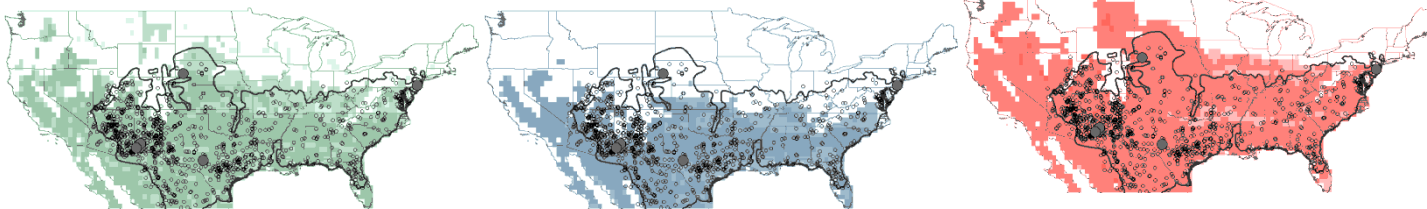
Nebraska traits



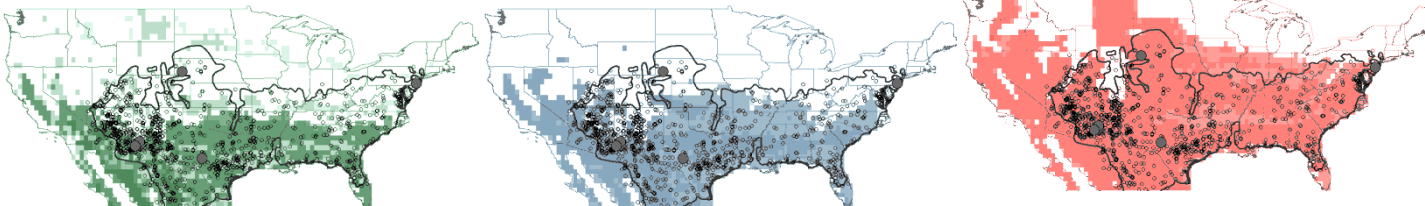
New Jersey traits



New Mexico traits



Texas traits

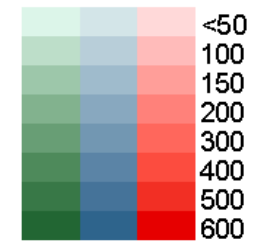


NORTH

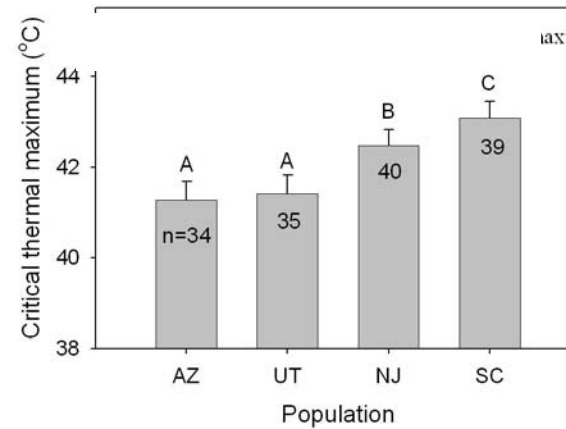
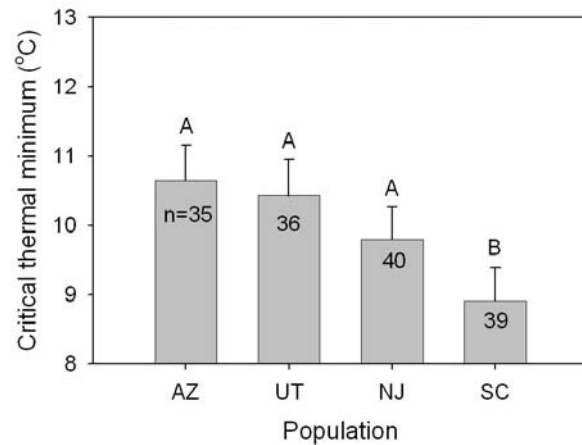


SOUTH

Lizards /1000m²



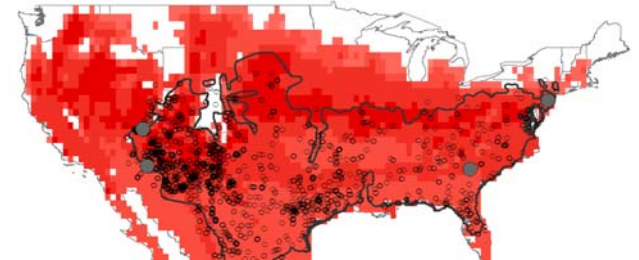
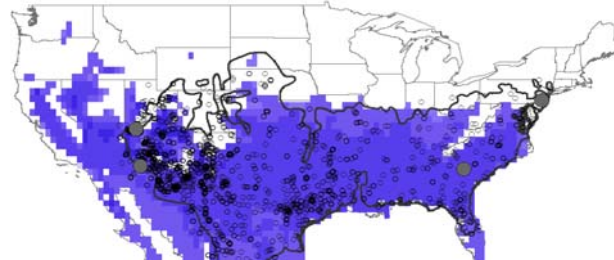
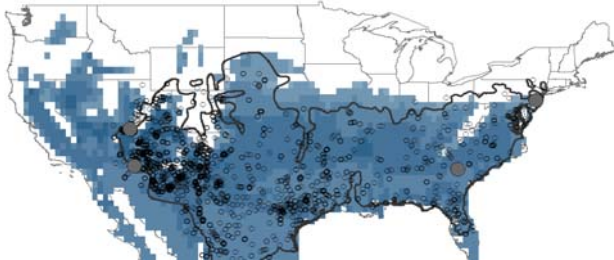
Range implications of physiological variation?



New Jersey SVL + Thermal Limits

+ NJ Metabolic Rates

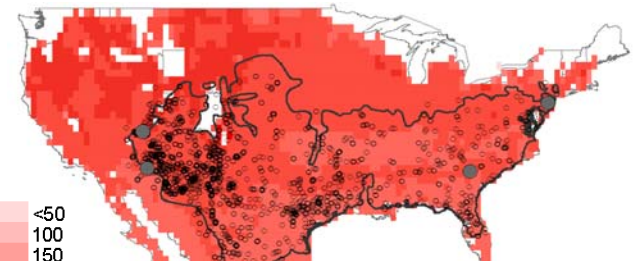
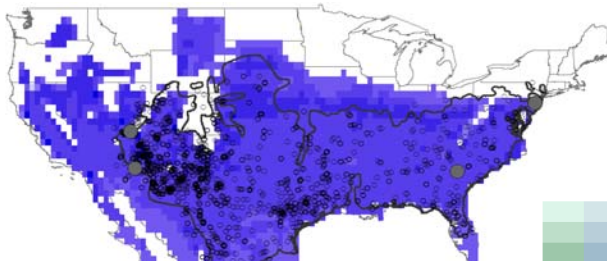
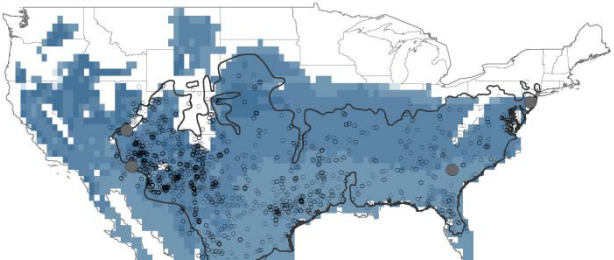
+ 3°C



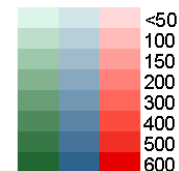
South Carolina SVL + Thermal Limits

+ SC Metabolic Rates

+ 3°C

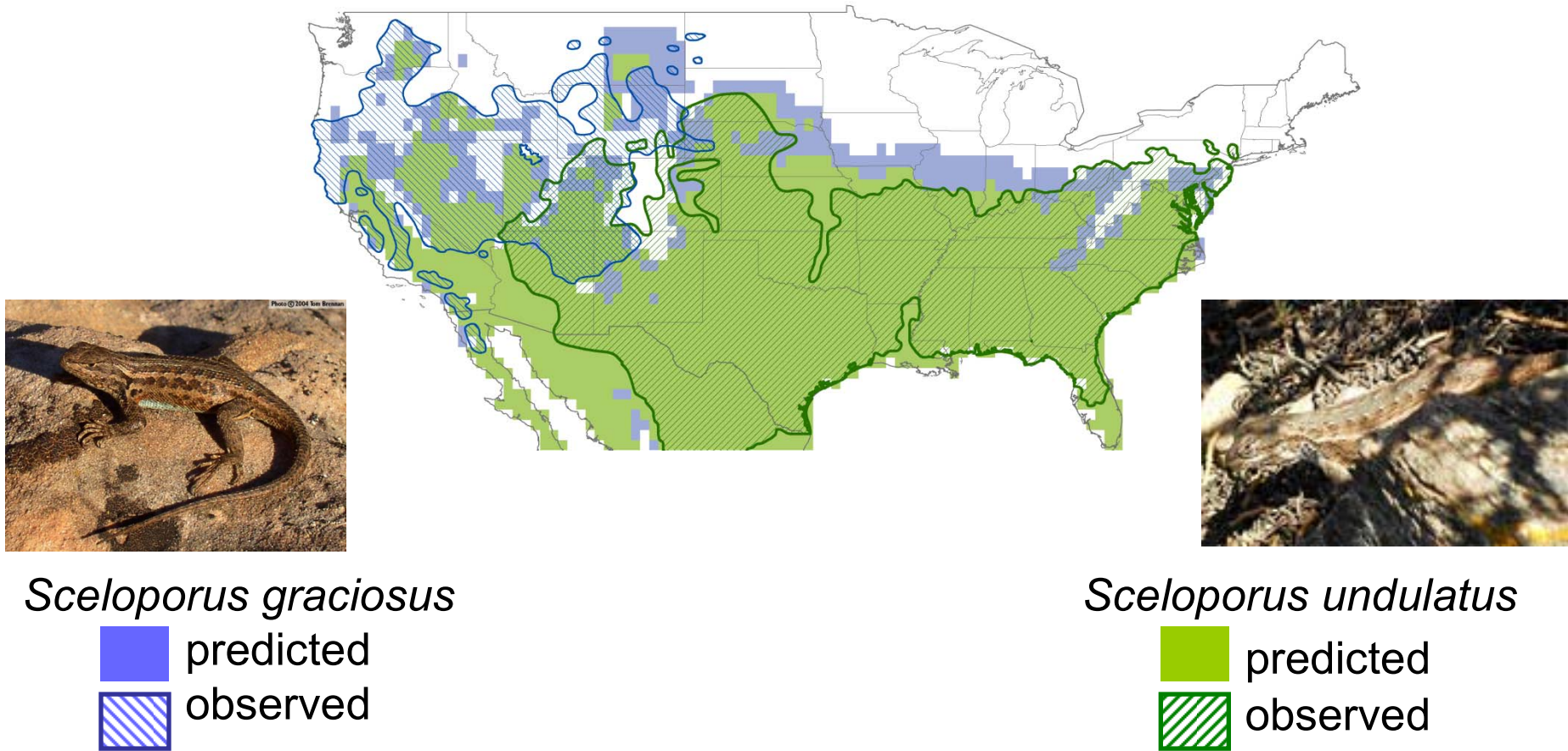


(Ehrenberger, Buckley, and Angilletta)



Lizards /1000m²

Species interactions?



(Buckley 2008)



Sachem skipper range models

correlative
models

biophysical
models

performance
curves

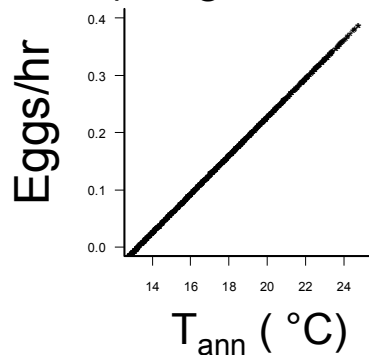
foraging energetic
models



$$R_0 = bE - Y$$

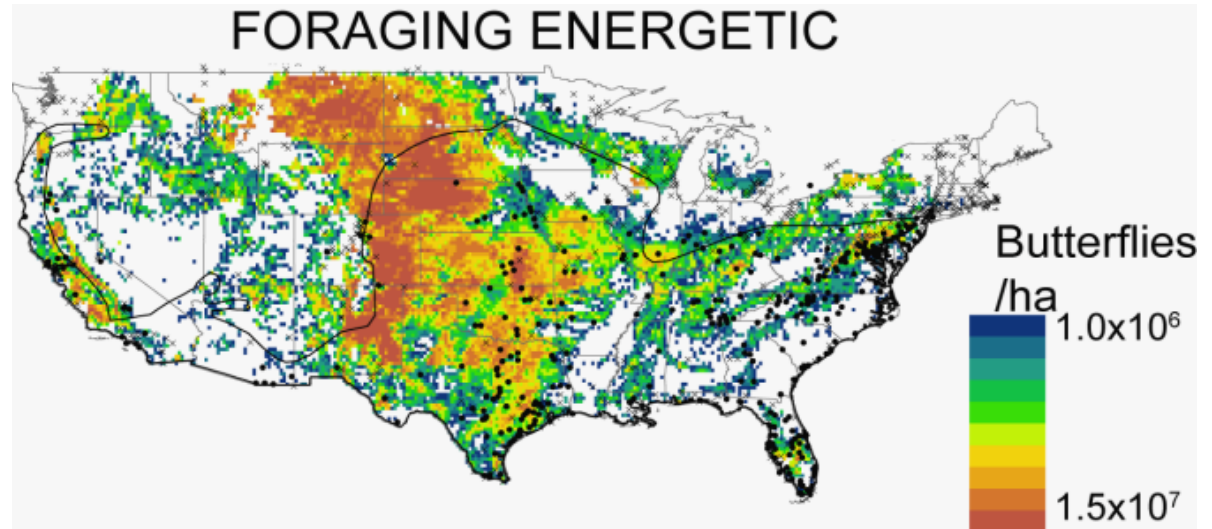
eggs/J J - J

Thermal constraints
Egg maturation
(Berger 2008)

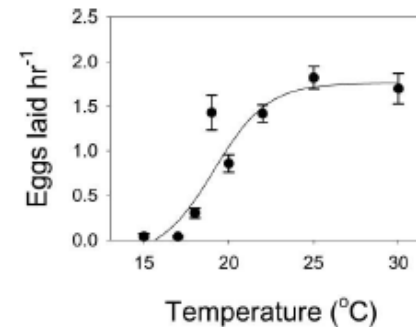


Density dependence in nectar resources

$$\left(1 - \frac{N_{forrate} r_{enrate}}{P}\right)$$



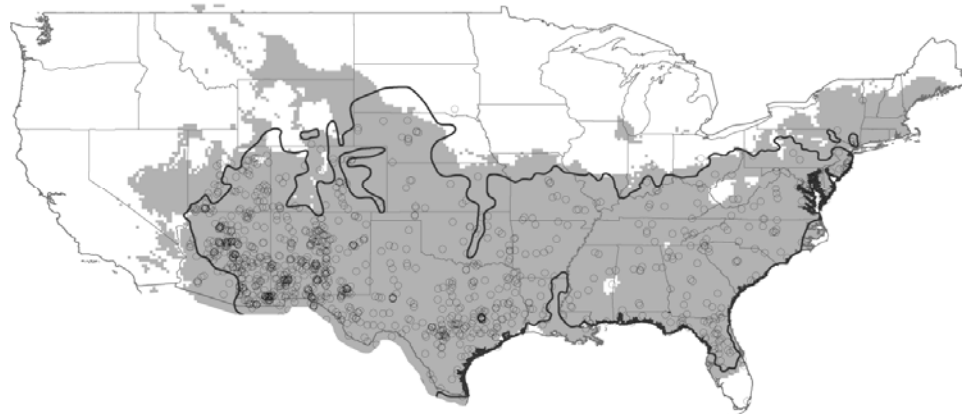
Ovipositing
(Gotthard 2007)



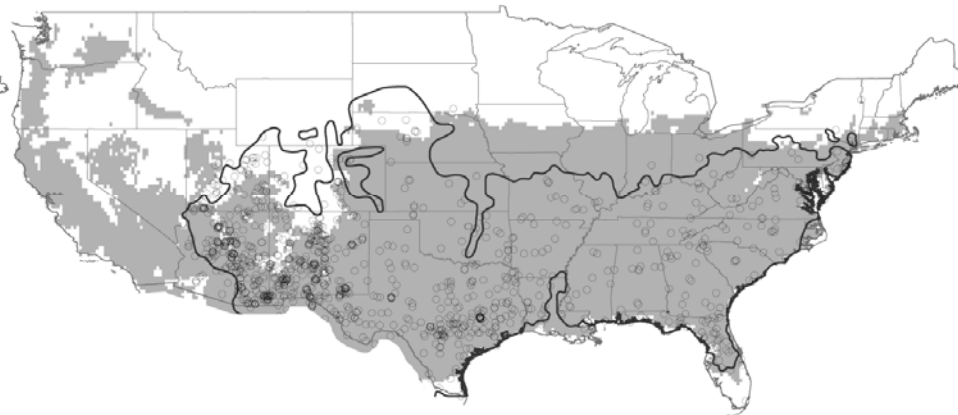


Fence lizard range models

CORRELATIVE- Maxent



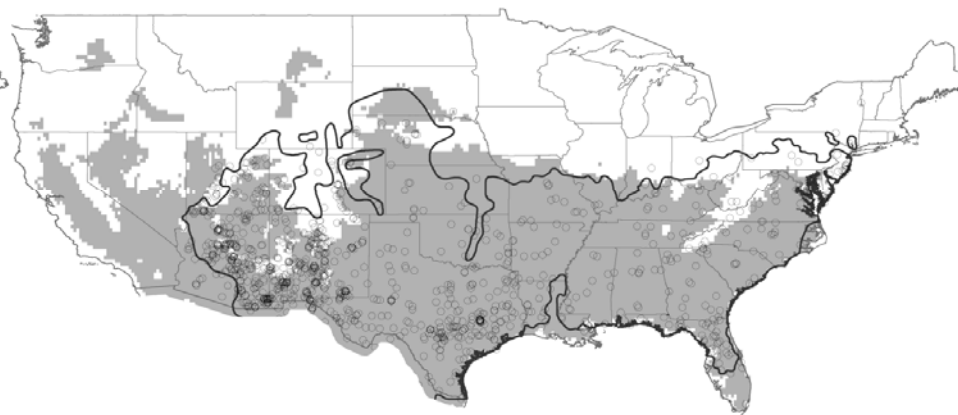
LIFE HISTORY



BIOPHYSICAL THRESHOLD



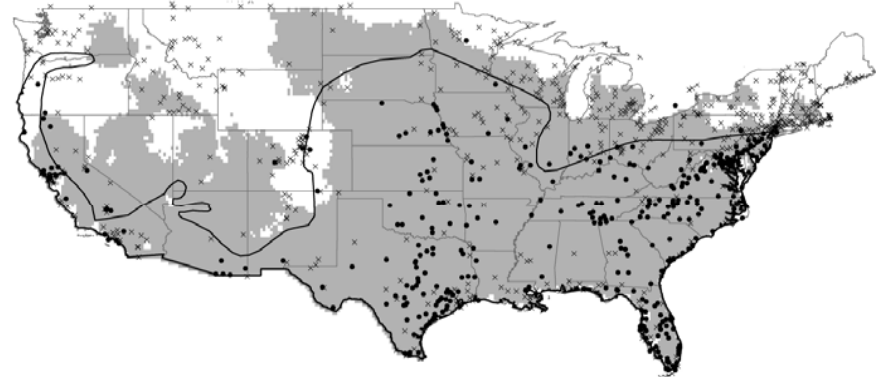
FORAGING ENERGETIC



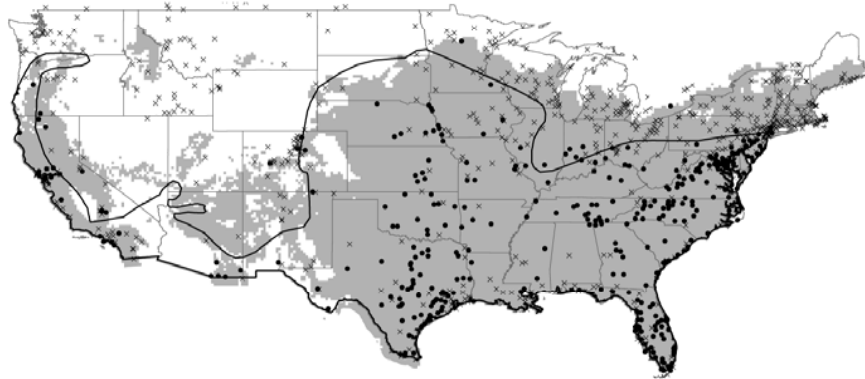


Sachem skipper range models

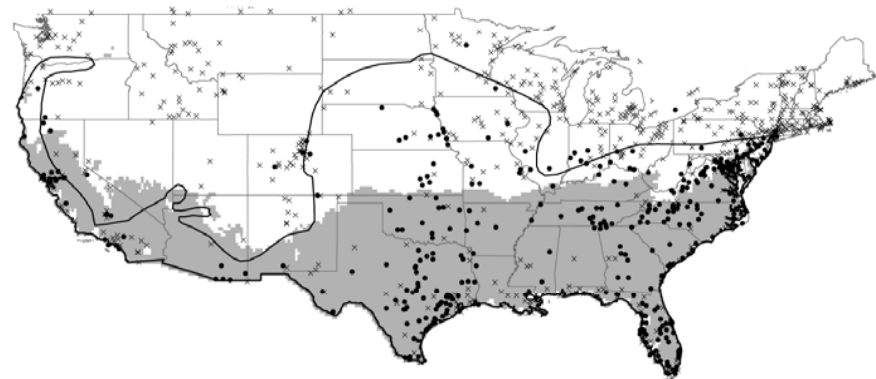
BIOPHYSICAL THRESHOLD



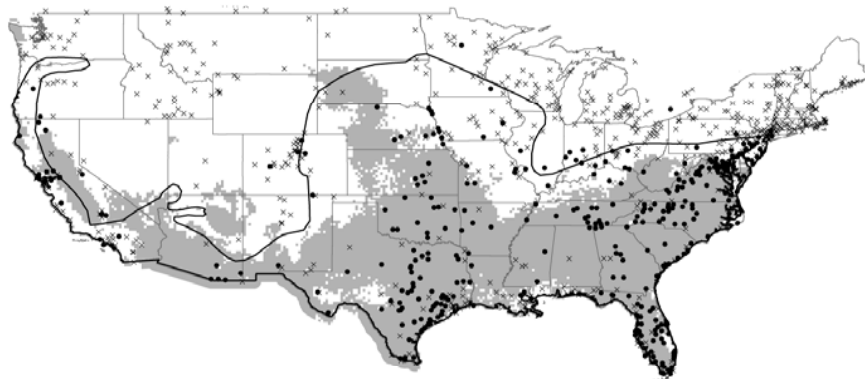
CORRELATIVE- Maxent



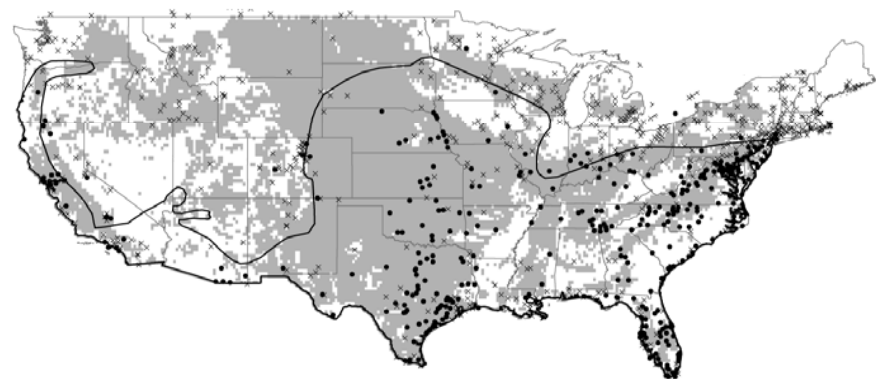
LIFE HISTORY

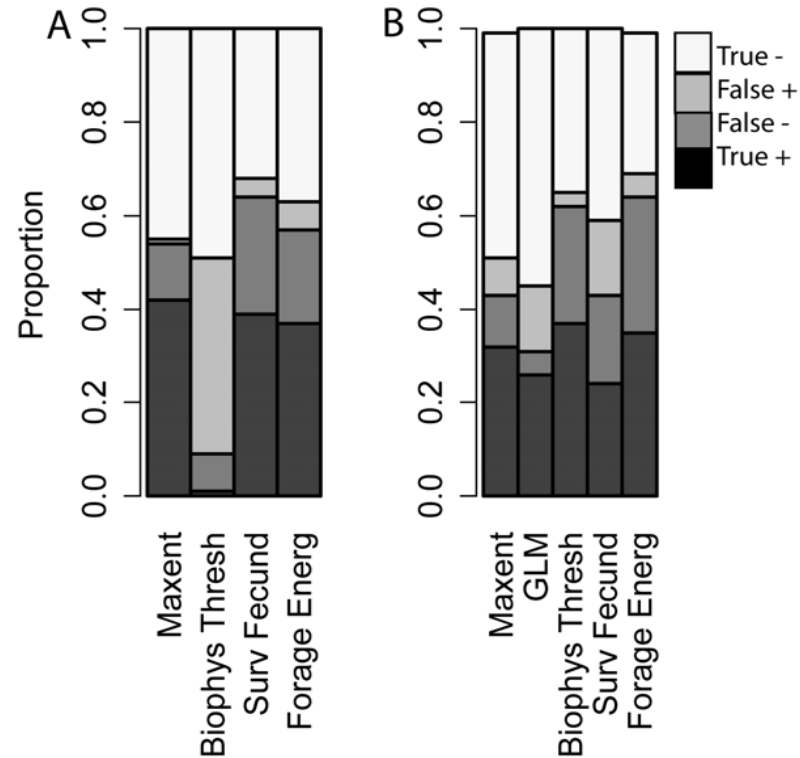


CORRELATIVE- GLM



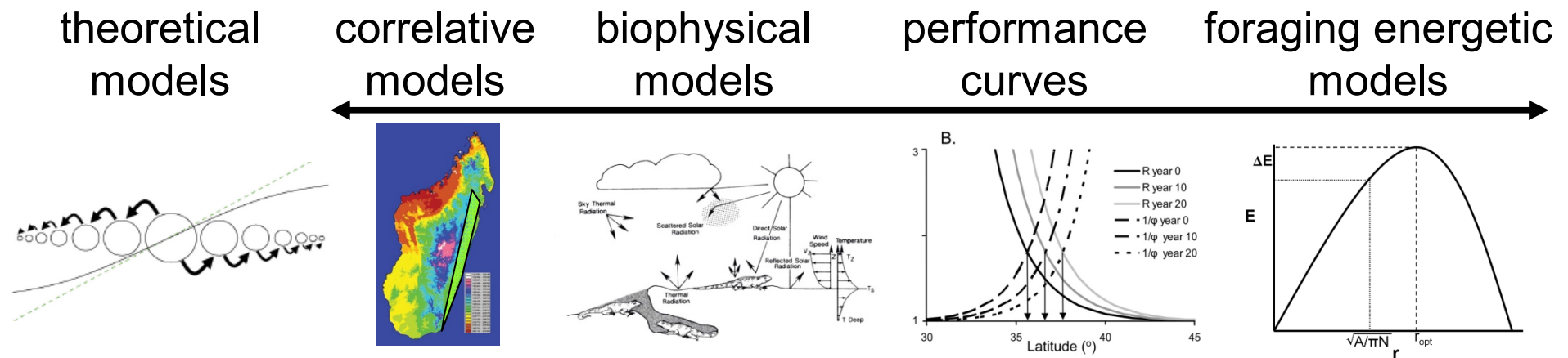
FORAGING ENERGETIC





Model	<i>S. undulatus</i>				<i>A. campestris</i>				
	% +	% -	% true	% localities	% +	% -	% true	% survey +	% survey -
Correlative- Maxent	0.98	0.78	0.87	0.99	0.80	0.81	0.81	0.97	0.44
Correlative- GLM					0.64	0.92	0.81	0.73	0.81
Biophysical threshold	0.02	0.86	0.50	0.04	0.92	0.58	0.72	0.94	0.43
Life history	0.90	0.56	0.71	0.88	0.61	0.68	0.65	0.77	0.69
Foraging energetic	0.85	0.65	0.74	0.84	0.88	0.51	0.66	0.90	0.31

Range models



- Large but narrowing divide between theory and application
- Correlative models readily provide abundant range information
Mechanistic models emerging as viable alternative
- Mechanistic models enable incorporating physiology, & demography, biotic constraints & evolution, individualistic responses

Assisted migration ?



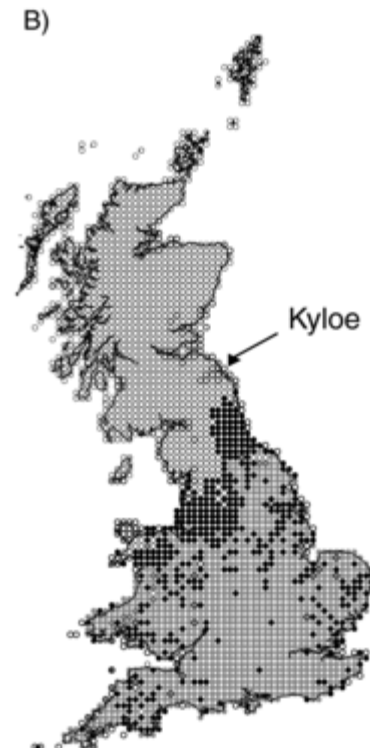
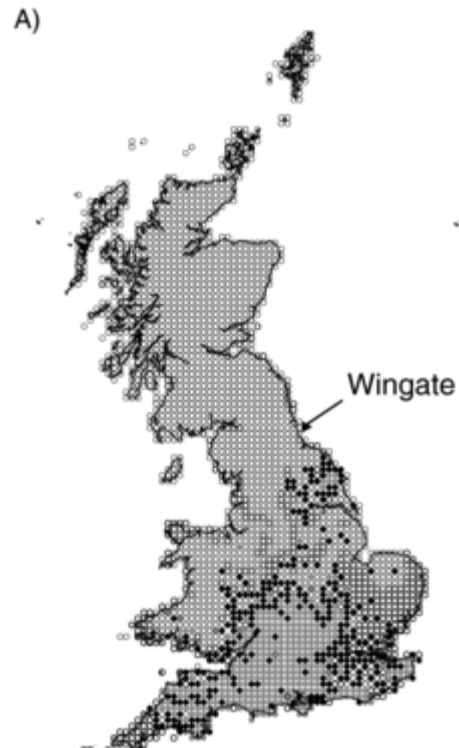
(Nature 2008)



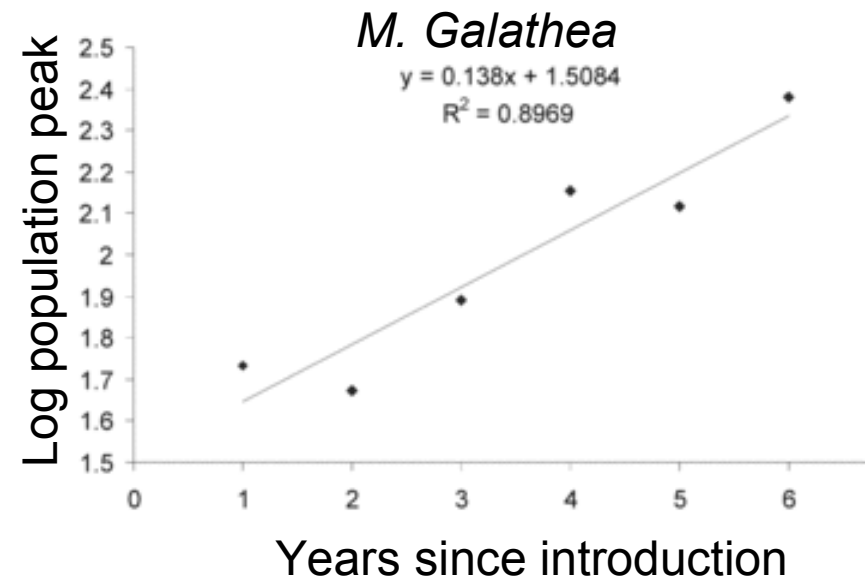
Melanargia galathea
(marbled white)



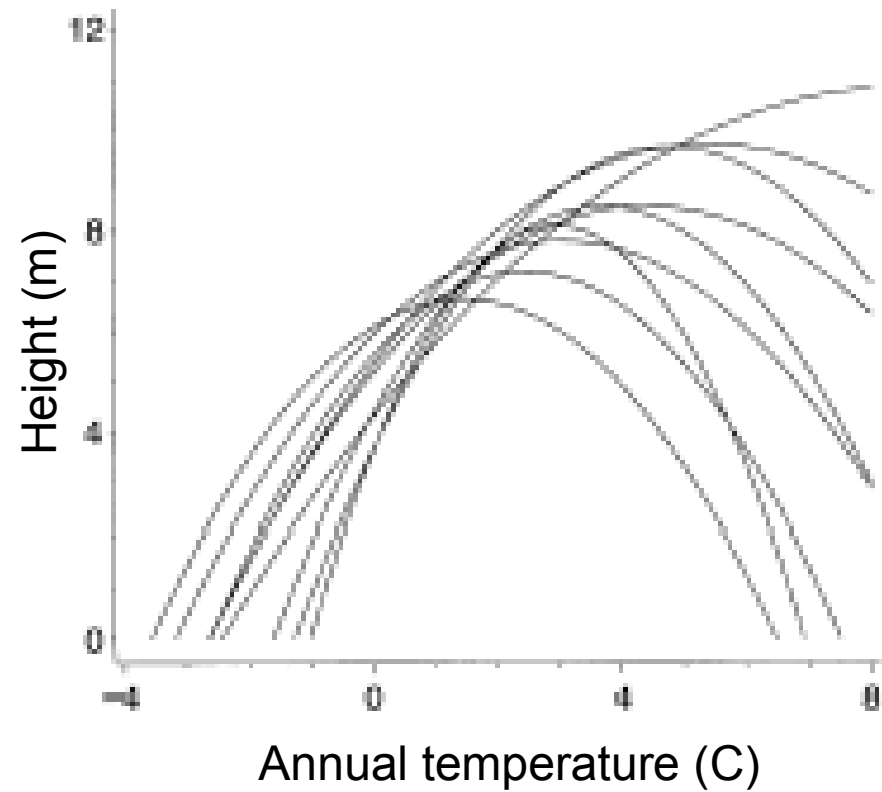
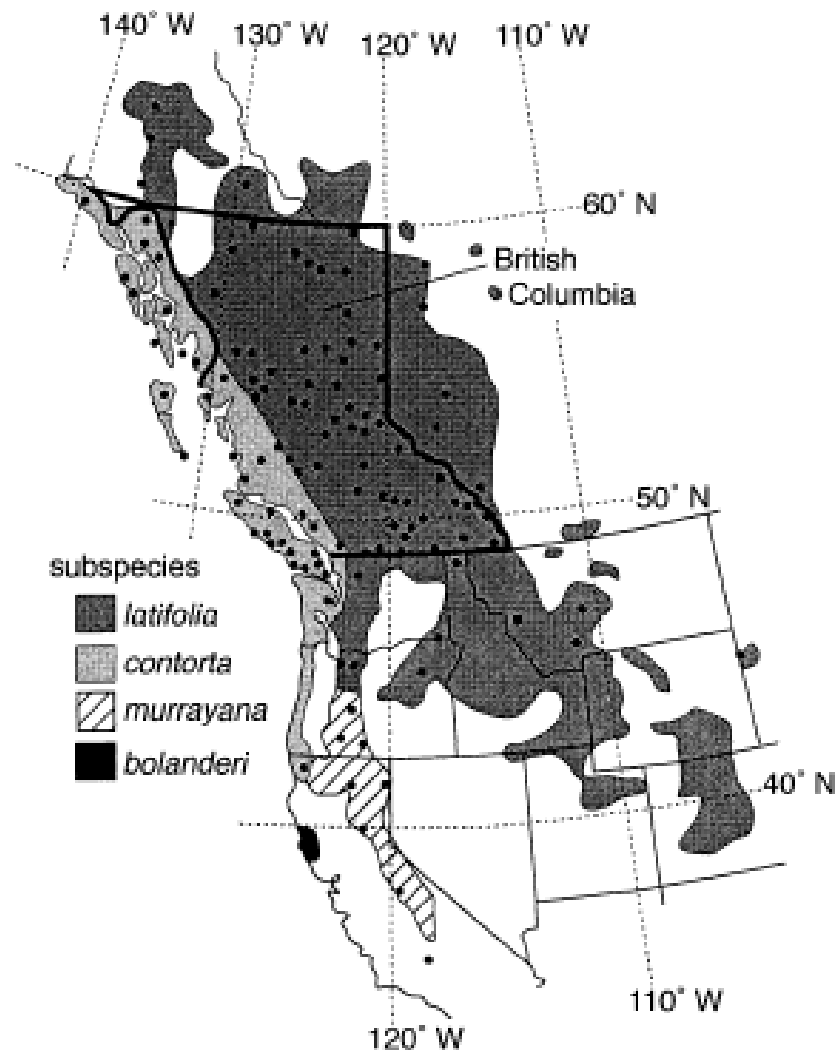
Thymelicus sylvestris
(small skipper)



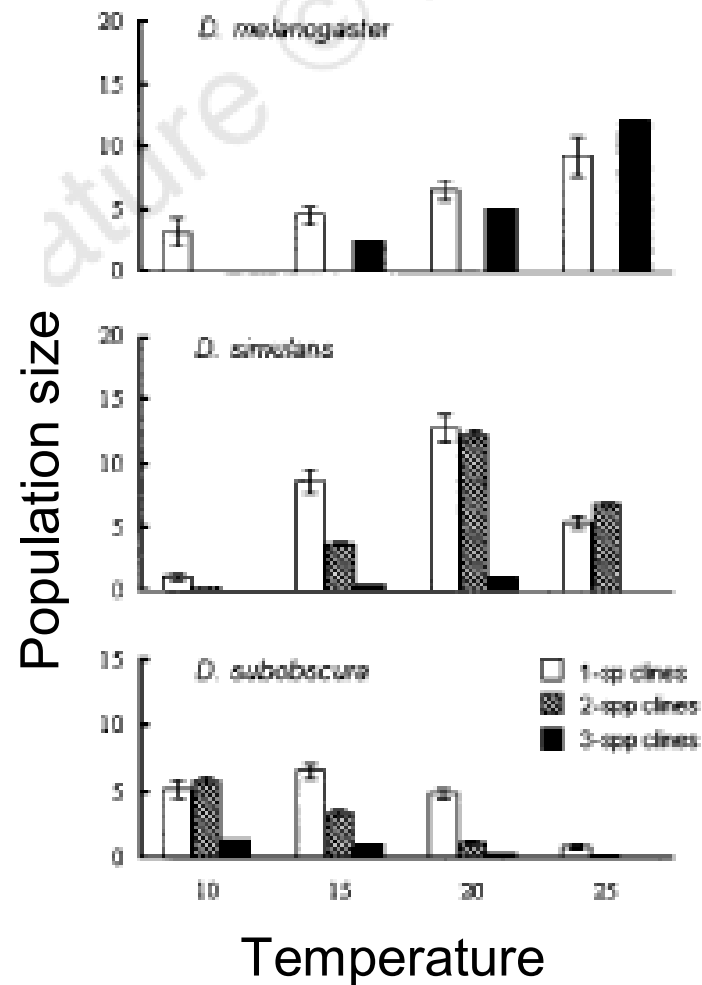
■ Newly occupied 1995-1999



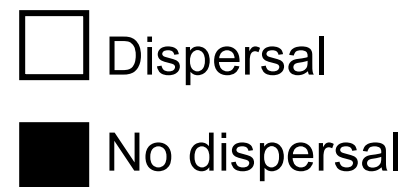
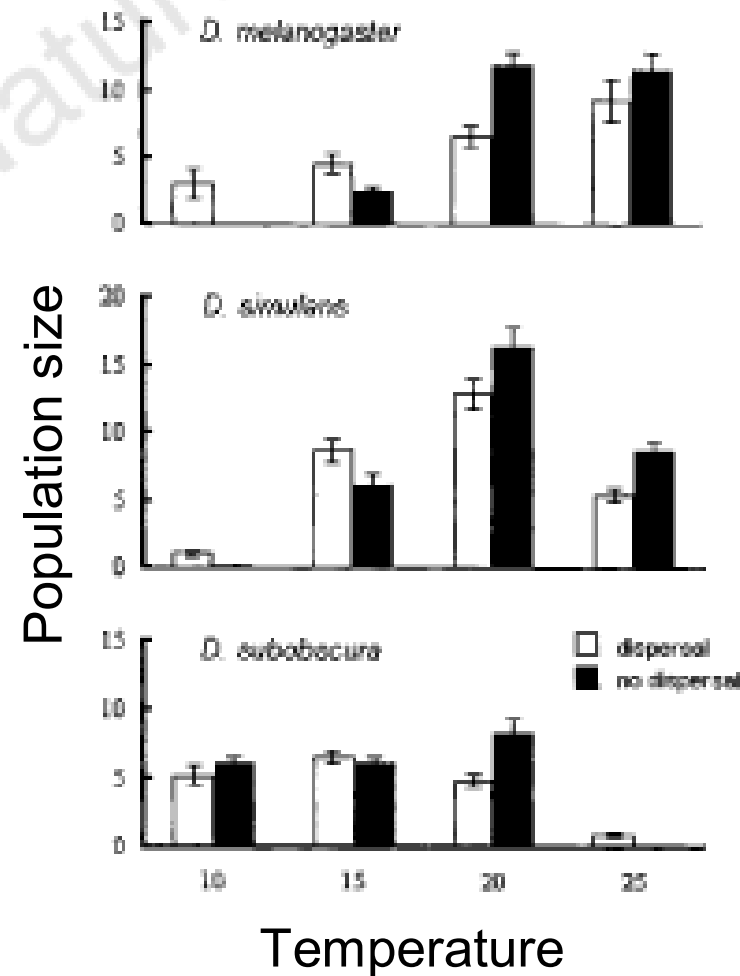
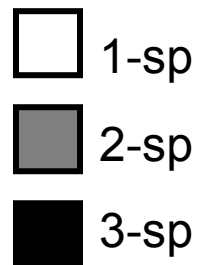
(Willis et al. 2009)



Pinus contorta
(Rehfeldt et al. 1999)



Species interactions



Drosophila
(Davis 1998)

Model sensitivity

Insect abundance

Energy / insect

T_b quantiles

0.5 a

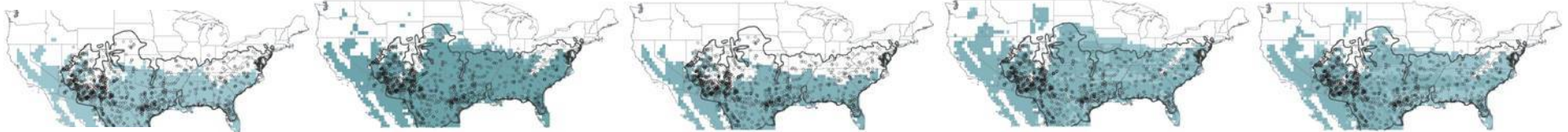
2 a

0.5 ei

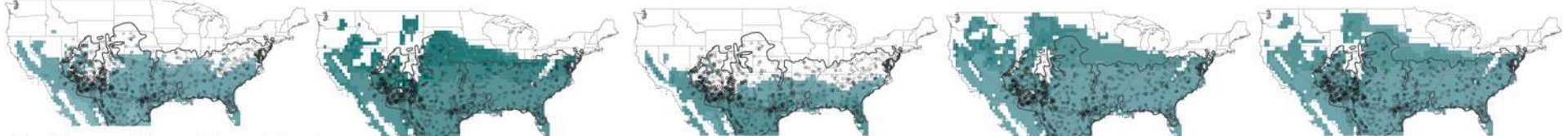
2 ei

10% -90% T_b

Nebraska traits



New Jersey traits



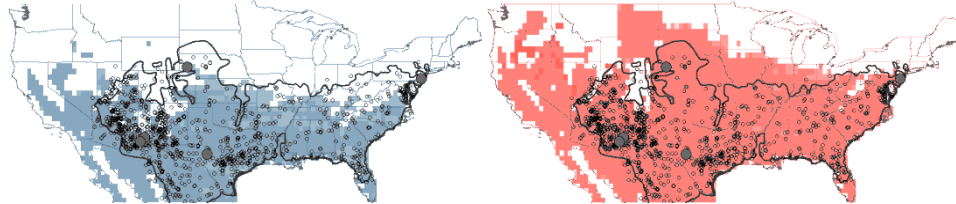
Lizards /1000m2

<50 100 150 200 300 400 500 600 700

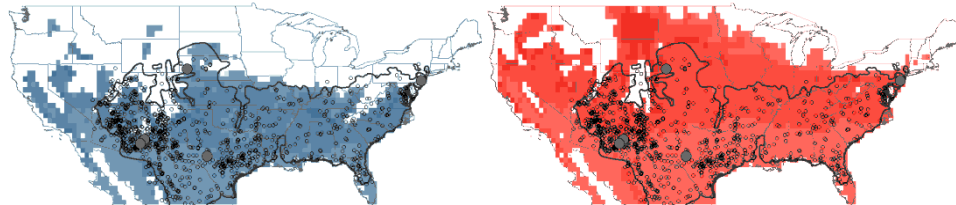


+ Population Life History + 3°C

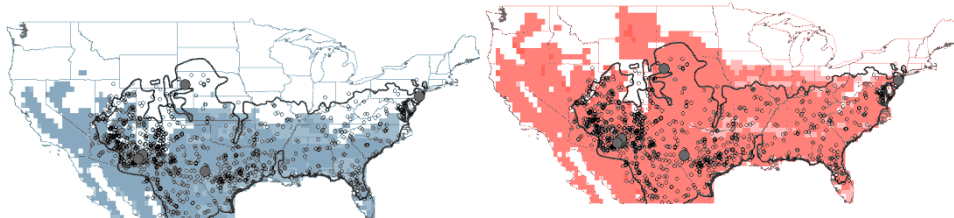
Nebraska traits



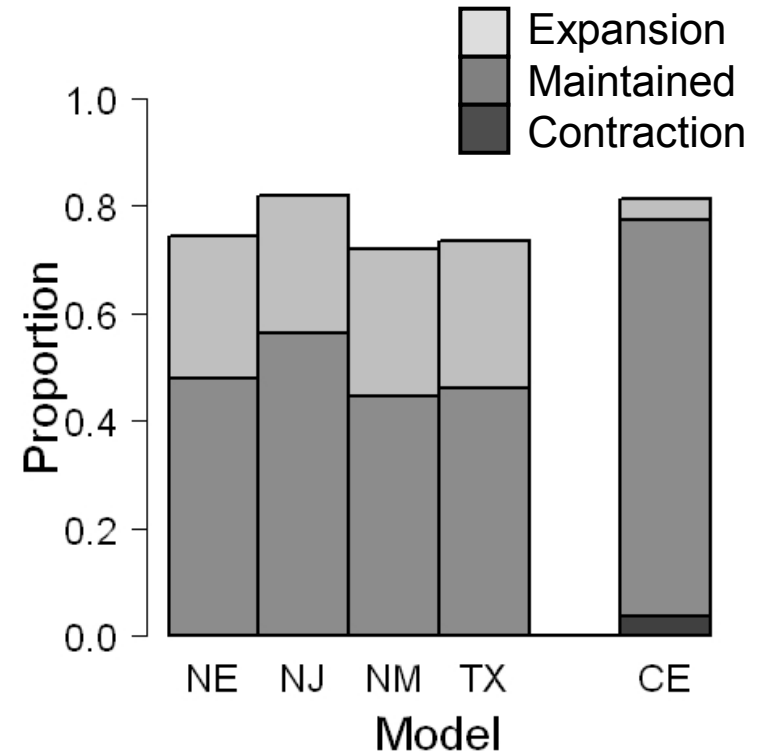
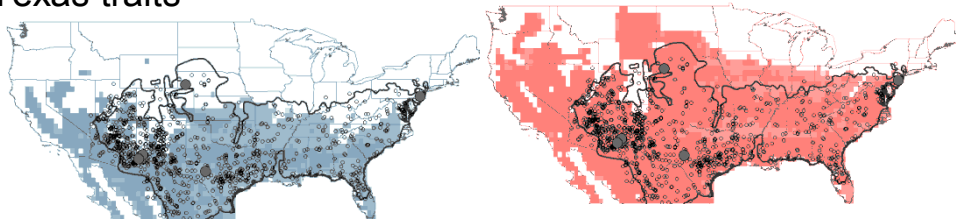
New Jersey traits



New Mexico traits



Texas traits



Dynamic model better omission / commission performance

Species interactions?
Adaptive potential?