

Integrating neontological and paleontological data to study species' responses to global change



Erin E. Saupe, Yale University

Outline:

(1) Study background

(2) Niche evolution

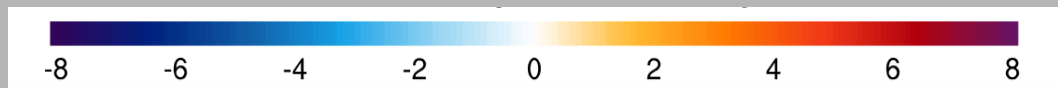
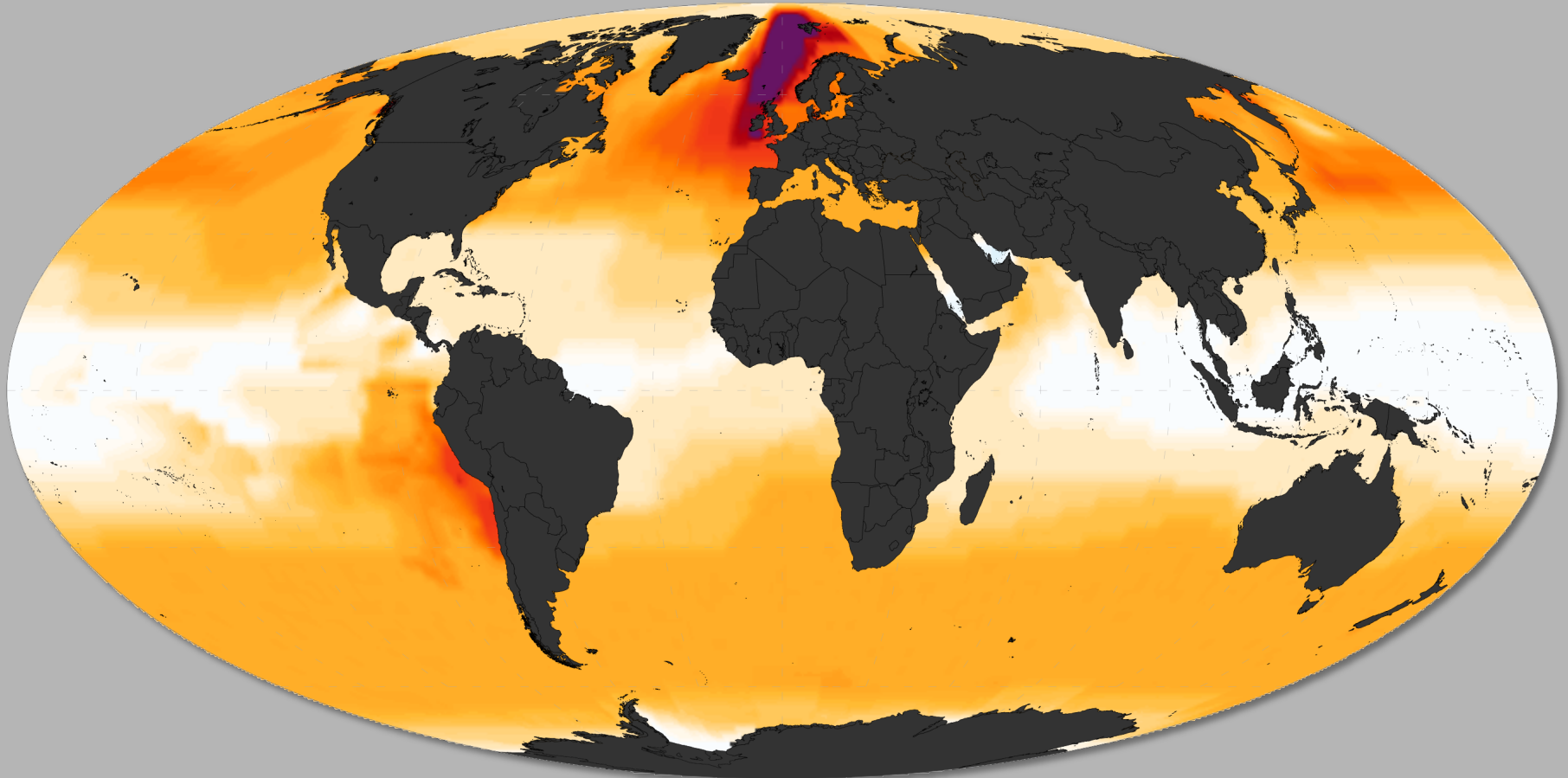
(3) Extinction selectivity



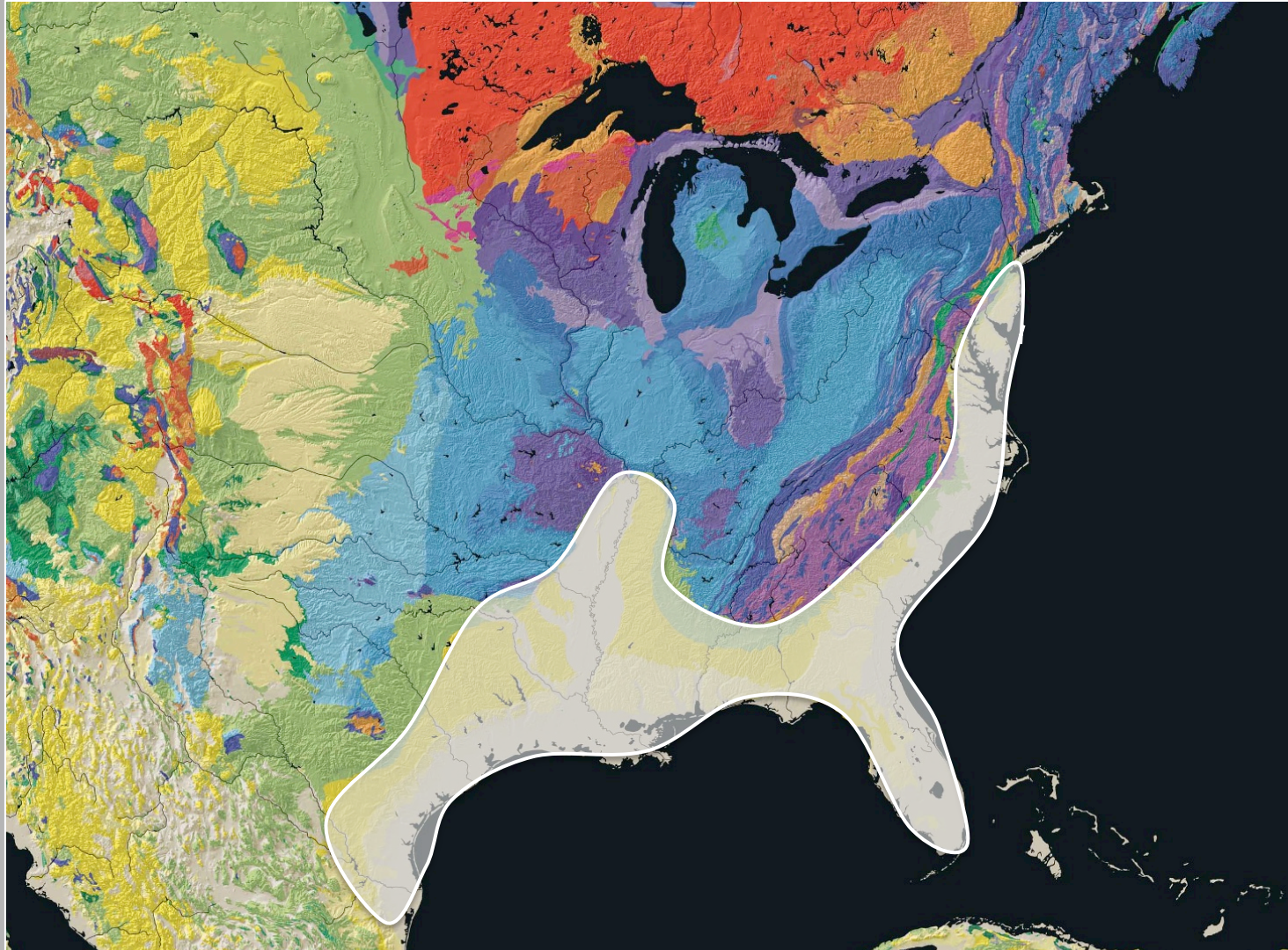
Learning
from the
fossil
record



Pliocene SST ($^{\circ}\text{C}$) Anomaly ~ 3.2 Ma



Atlantic & Gulf Coastal Plain



Study System



Crassostrea virginica



Terebra dislocata



Crepidula fornicata



Dinocardium robustum



Anomia simplex



Bulla occidentalis



Mercenaria campechiensis



Neverita duplicata

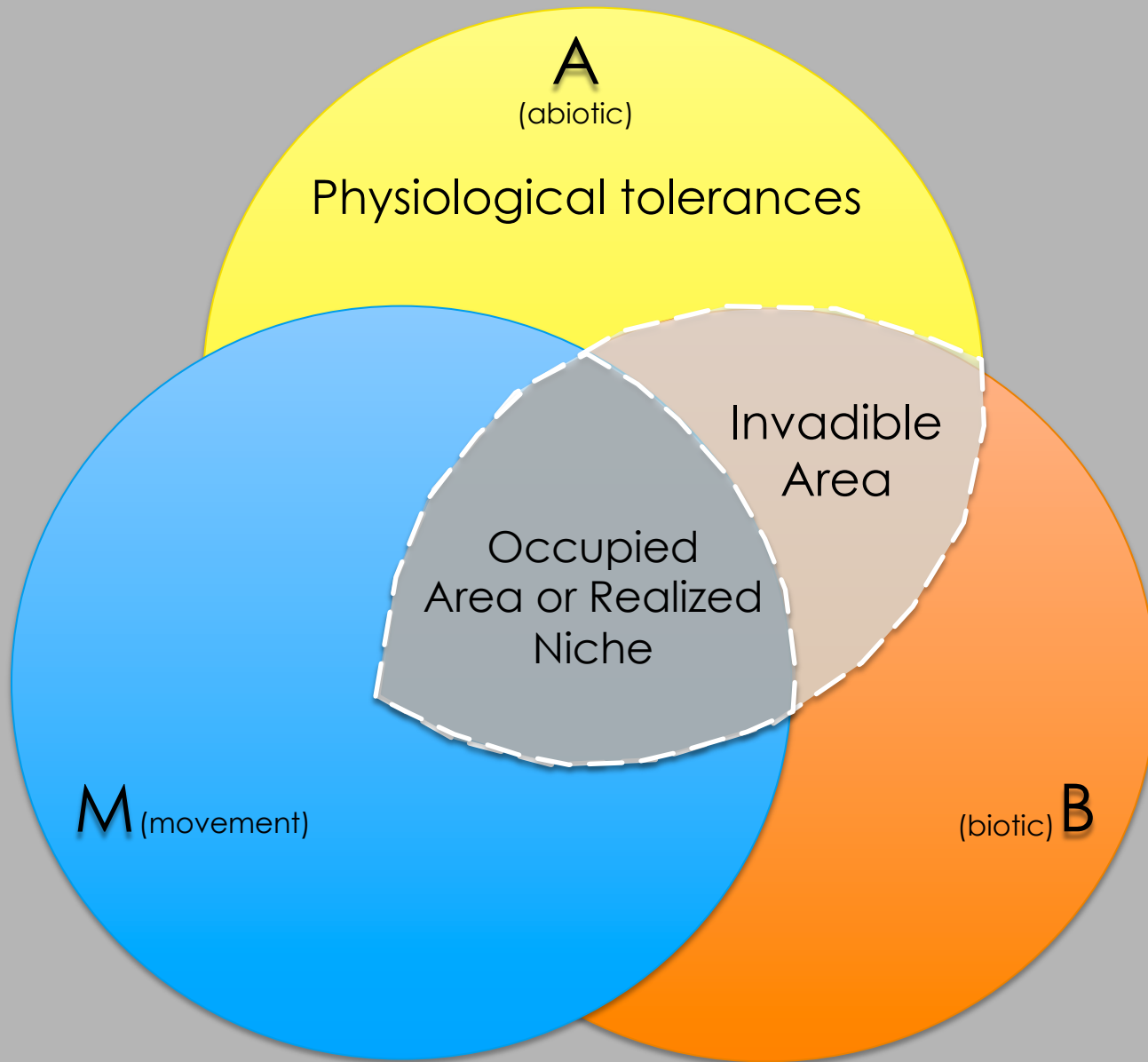


Lucina pensylvanica

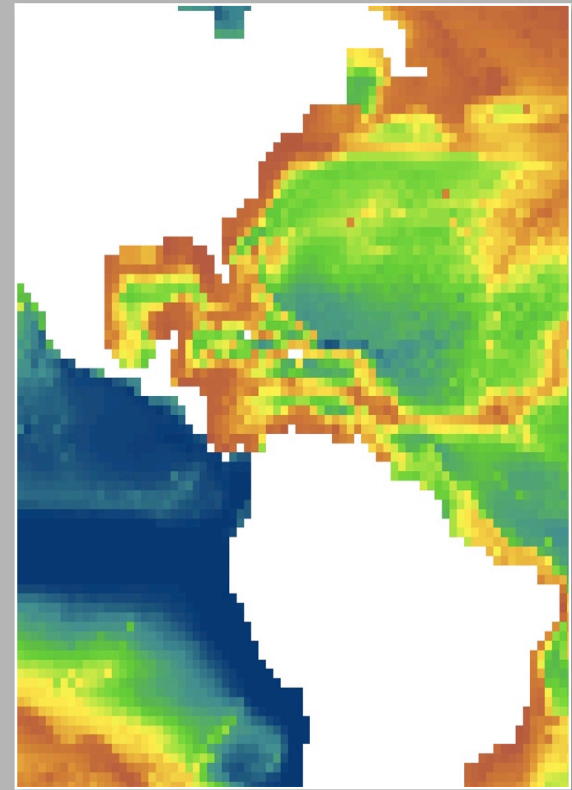
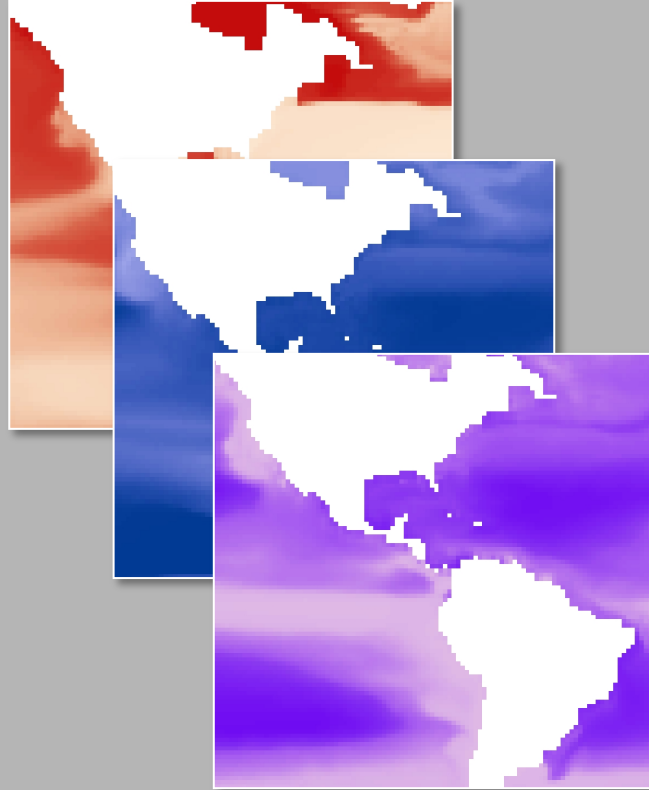


Oliva sayana





Ecological Niche Modeling



Outline:

(1) Study background

(2) Niche evolution

(3) Extinction selectivity

Frequency of niche shifts?



Species



Crassostrea virginica



Terebra dislocata



Crepidula fornicata



Dinocardium robustum



Anomia simplex



Bulla occidentalis



Mercenaria campechiensis



Neverita duplicata



Lucina pensylvanica



Oliva sayana



Study Interval

EON	ERA	PERIOD	EPOCH		
Phanerozoic	Cenozoic	Quaternary	Holocene		
			Pleistocene	Late	
		Early			
		Tertiary	Neogene	Pliocene	Late
				Early	
				Miocene	Late
					Middle
					Early
			Paleogene	Oligocene	Late
					Early
				Eocene	Late
					Middle
					Early
		Paleocene	Late		
	Early				
	Mesozoic	Cretaceous	Late		
			Early		
		Jurassic	Late		
			Middle		
			Early		
		Triassic	Late		
Middle					
Early					

* Present

* ~130 Ka

* ~3.2–3.0 Ma

Hypotheses



Pliocene

Pleistocene

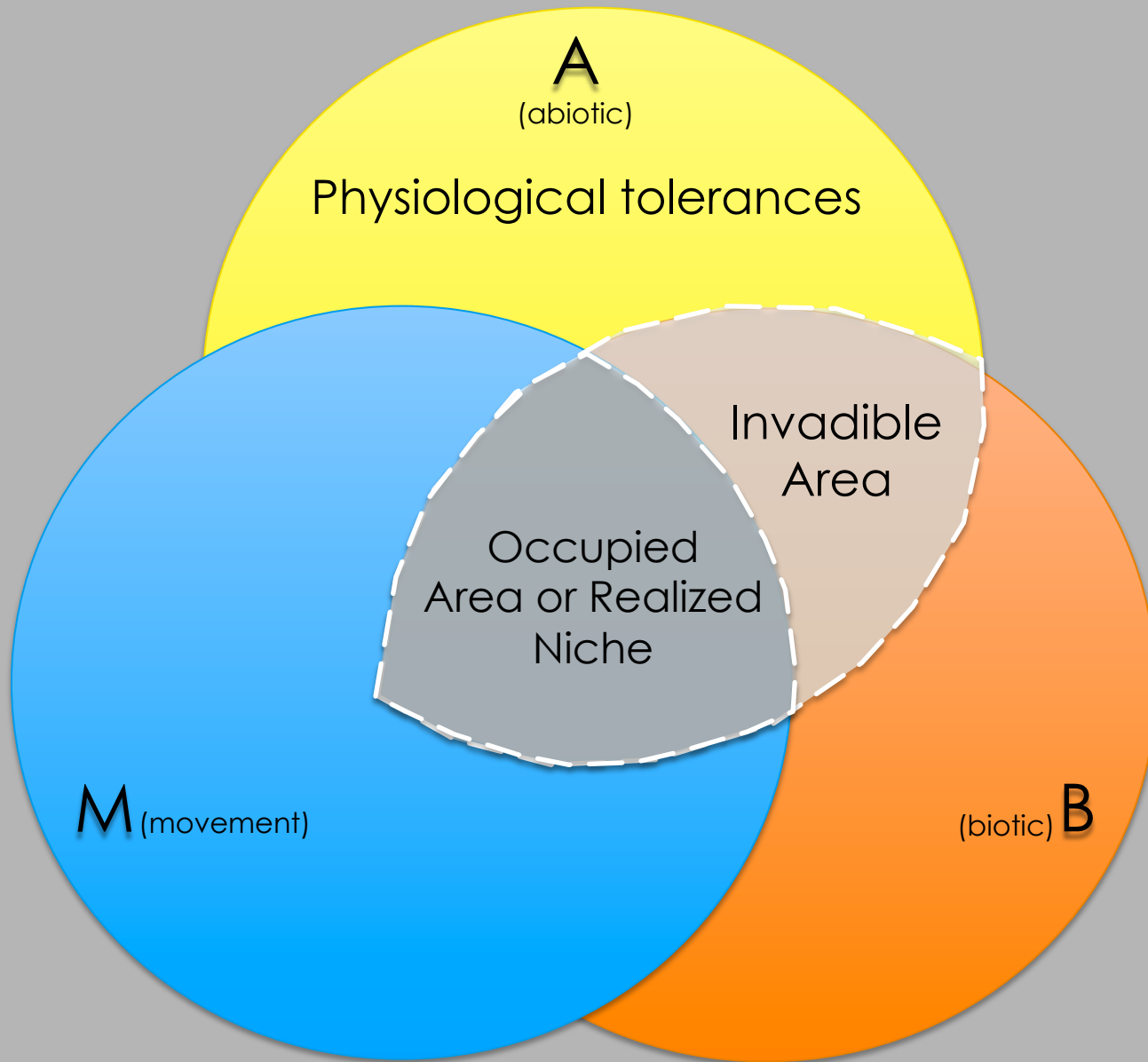
Present

x 10 species
x comparison type






Species environmental preferences remained stable over this interval

The upper and lower thermal tolerance limits for these species remained constant across the interval



Occurrence data

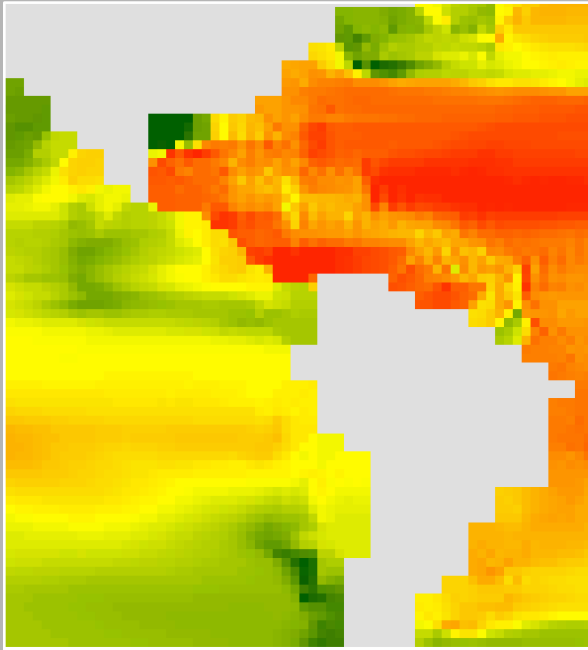


Species	Pliocene		Pleistocene		Present	
	All	Unique	All	Unique	All	Unique
	219	13	39	15	157	31
	82	6	60	9	331	58
	42	7	24	10	73	37
	156	16	31	10	157	42
	59	7	46	11	119	22
	127	7	29	8	72	37
	187	14	94	12	120	24
	134	16	54	14	169	28
	76	9	28	8	177	25
	58	9	37	13	147	20

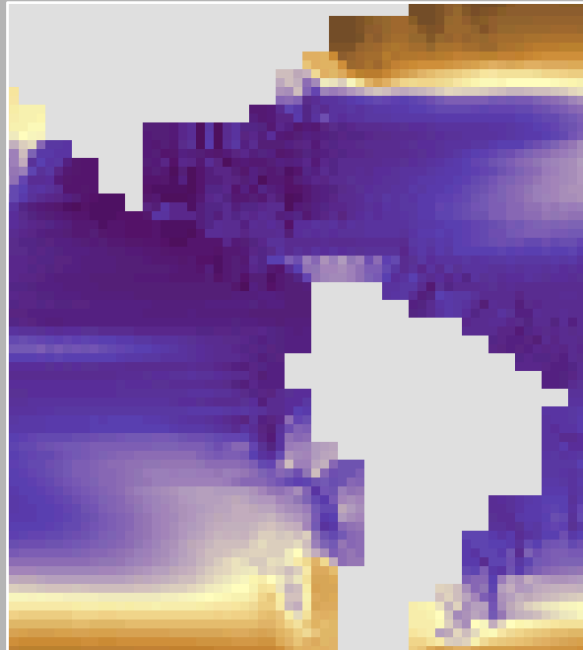
Georeferenced 3,104 records

Environmental Data

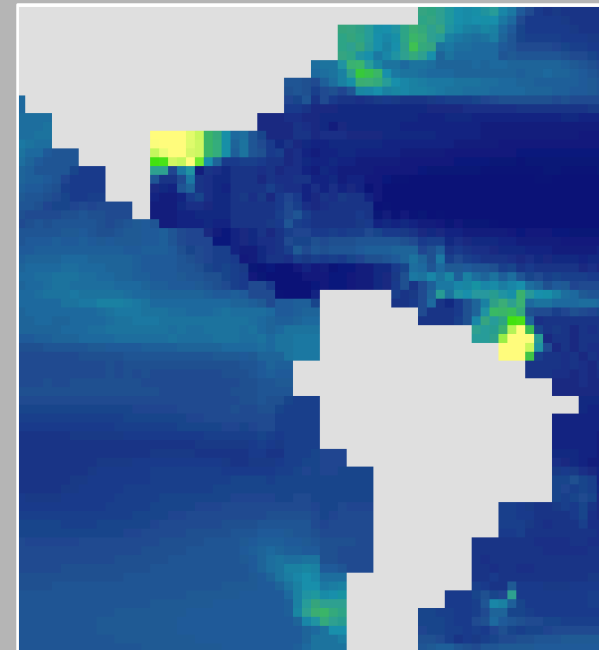
Max salinity



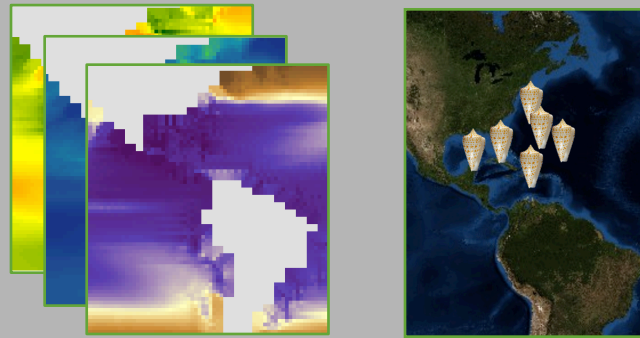
Max temperature



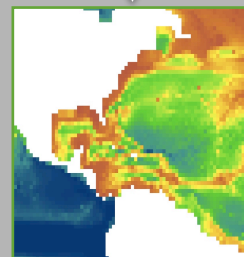
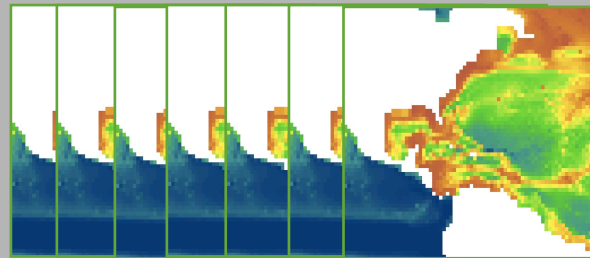
Min salinity



Model algorithm

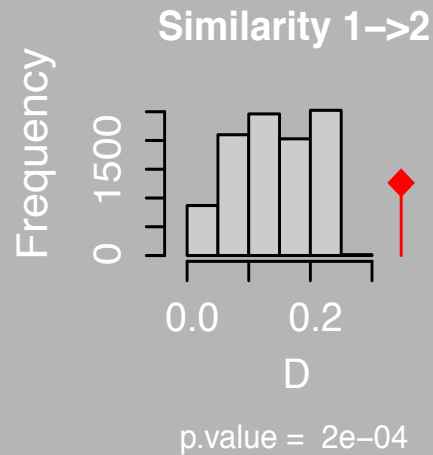
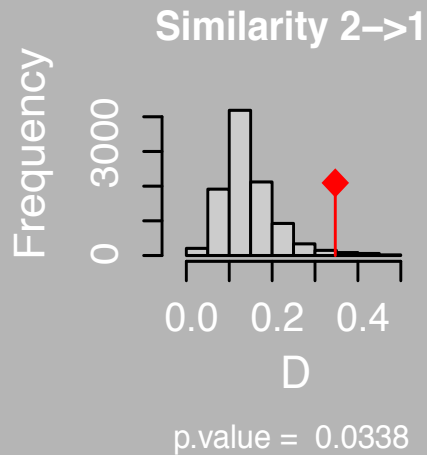
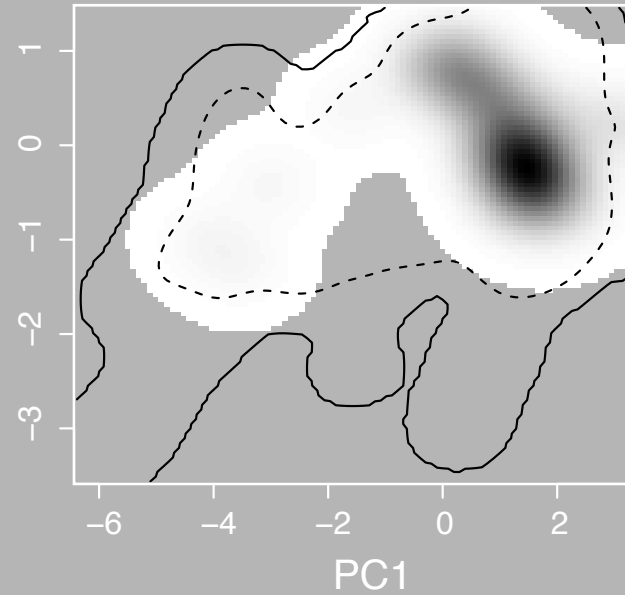
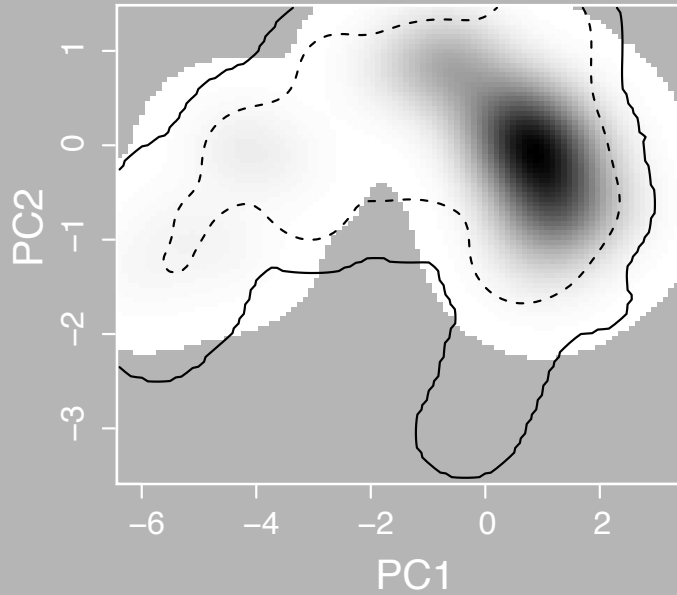


Maxent



Median model

Pleistocene to Present



Anomia simplex

Niche evolution

Geographic – ENM Tools

	Plio-Pleis		Pres-Pleis		Plio-Pres	
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	NS	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	NS
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05

Environmental – PCA

	Plio-Pleis		Pleis-Pres		Plio-Pres	
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	0.26	0.34
	<.05	<.05	<.05	<.05	<.05	<.05
	0.27	<.05	<.05	<.05	0.14	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	0.12	<.05
	<.05	<.05	<.05	<.05	<.05	<.05
	<.05	<.05	<.05	<.05	<.05	<.05

The background of the slide features a collection of various seashells, including bivalves, gastropods, and nautilus-like shells, rendered in a light, semi-transparent style. The shells are arranged in a somewhat horizontal line across the top and middle of the slide, with some overlapping. The overall aesthetic is scientific and naturalistic.

Species niche characteristics conserved within lineages

Buffeted by environment, not adapting to environment

Pattern congruent with habitat tracking in fossil record & modern distributional shifts

Conservation implications:



Extinction will result if environmental changes occur too rapidly

Validates ENM methodology to predict future responses of species to climate change

Outline:

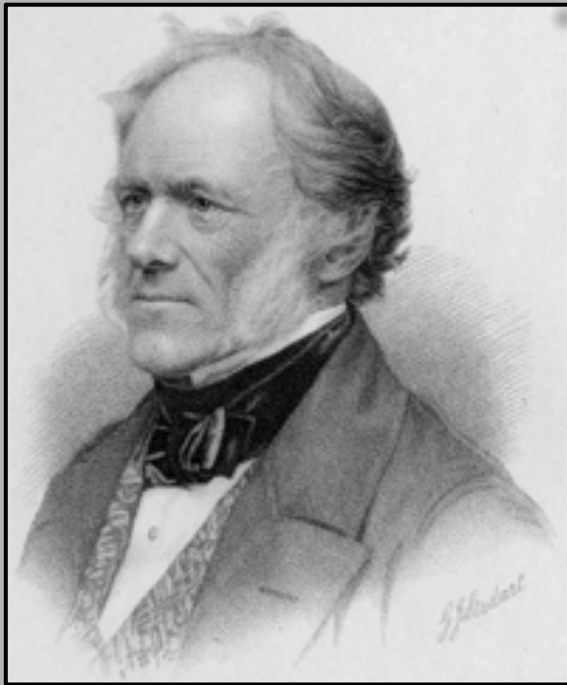
(1) Study background

(2) Niche evolution

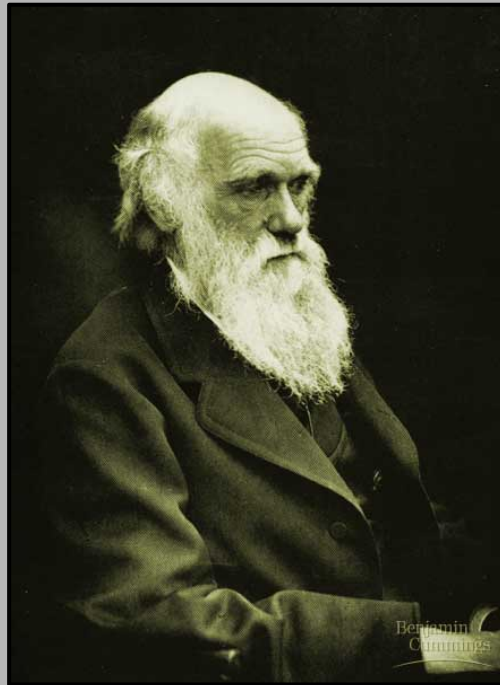
(3) Extinction selectivity

Extinction selectivity: historical perspective

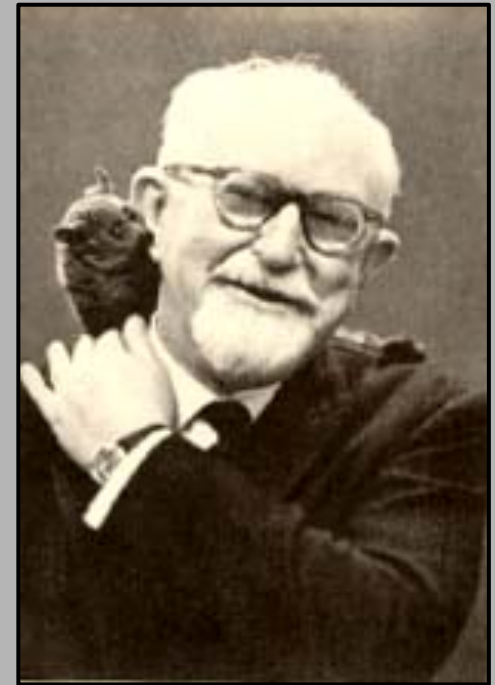
Charles Lyell



Charles Darwin



G.G. Simpson

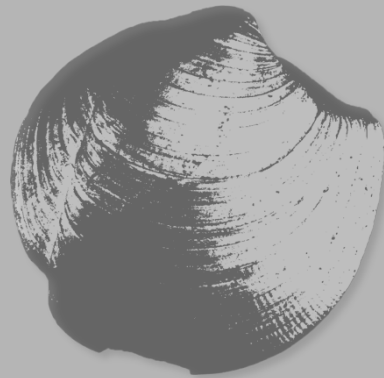


50 bivalve genera (8 families)

Anomiidae
Arcidae
Cardiidae
Carditidae
Lucinidae
Ostreidae
Tellinidae
Veneridae

16 gastropod genera (8 families)

Bullidae
Calyptraeidae
Conidae
Fasciolariidae
Muricidae
Naticidae
Olividae
Terebridae



Collections & field work



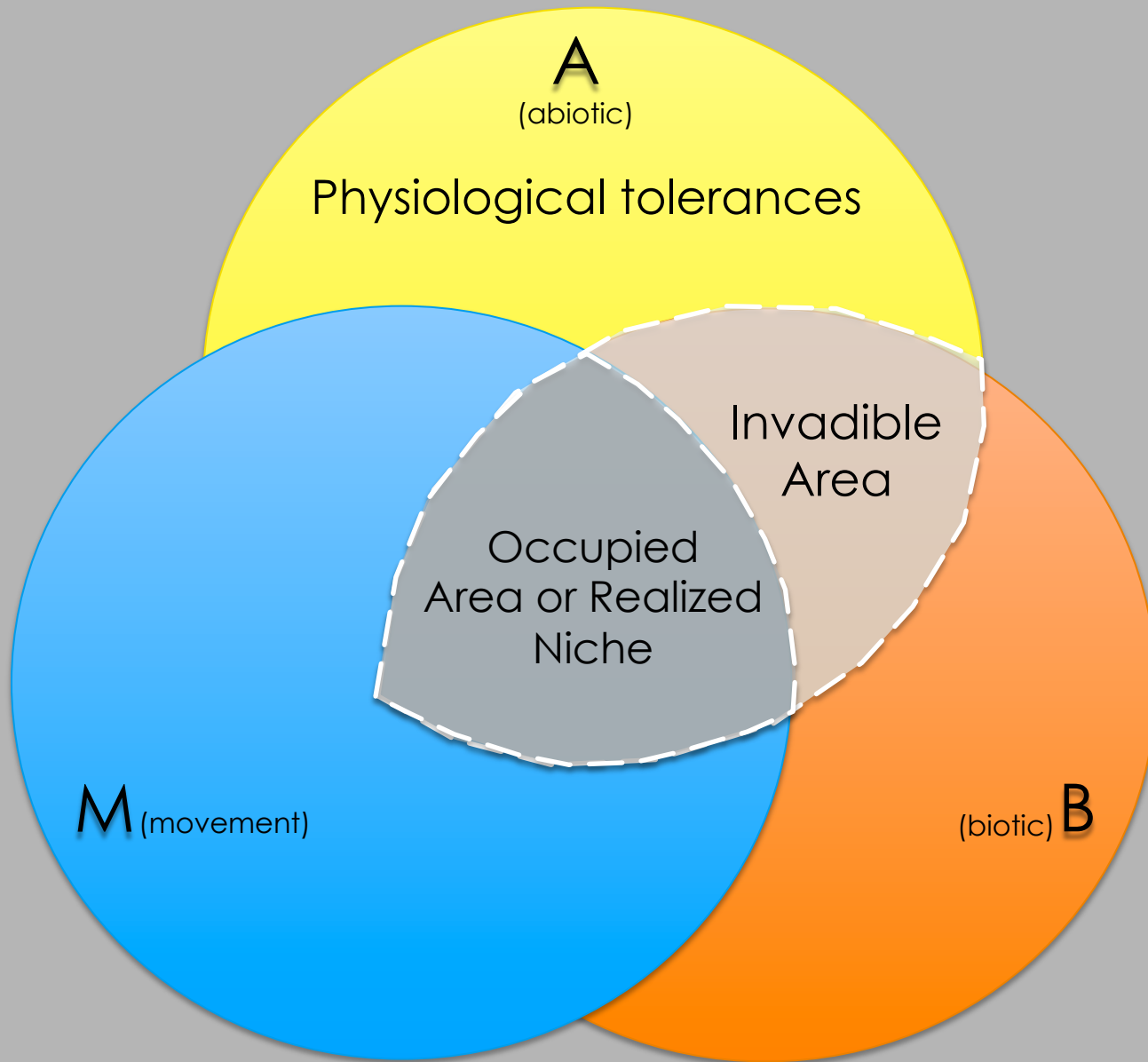
Virginia Museum of Natural History



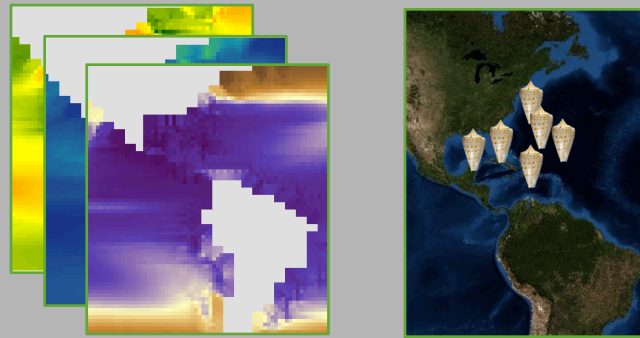
Florida Museum of Natural History

Testing whether species that survived:

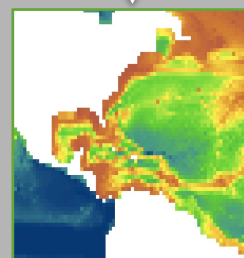
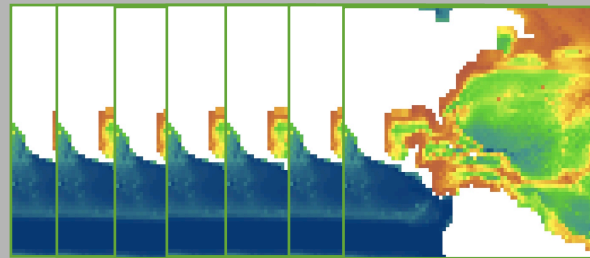
- (i) had larger geographic ranges than now-extinct species
- (ii) had broader fundamental niche breadths than now-extinct species
- (iii) occupied broader realized environmental space than now-extinct species
- (iv) had more suitable area remaining during the LGM than now-extinct species



Model algorithm



Maxent

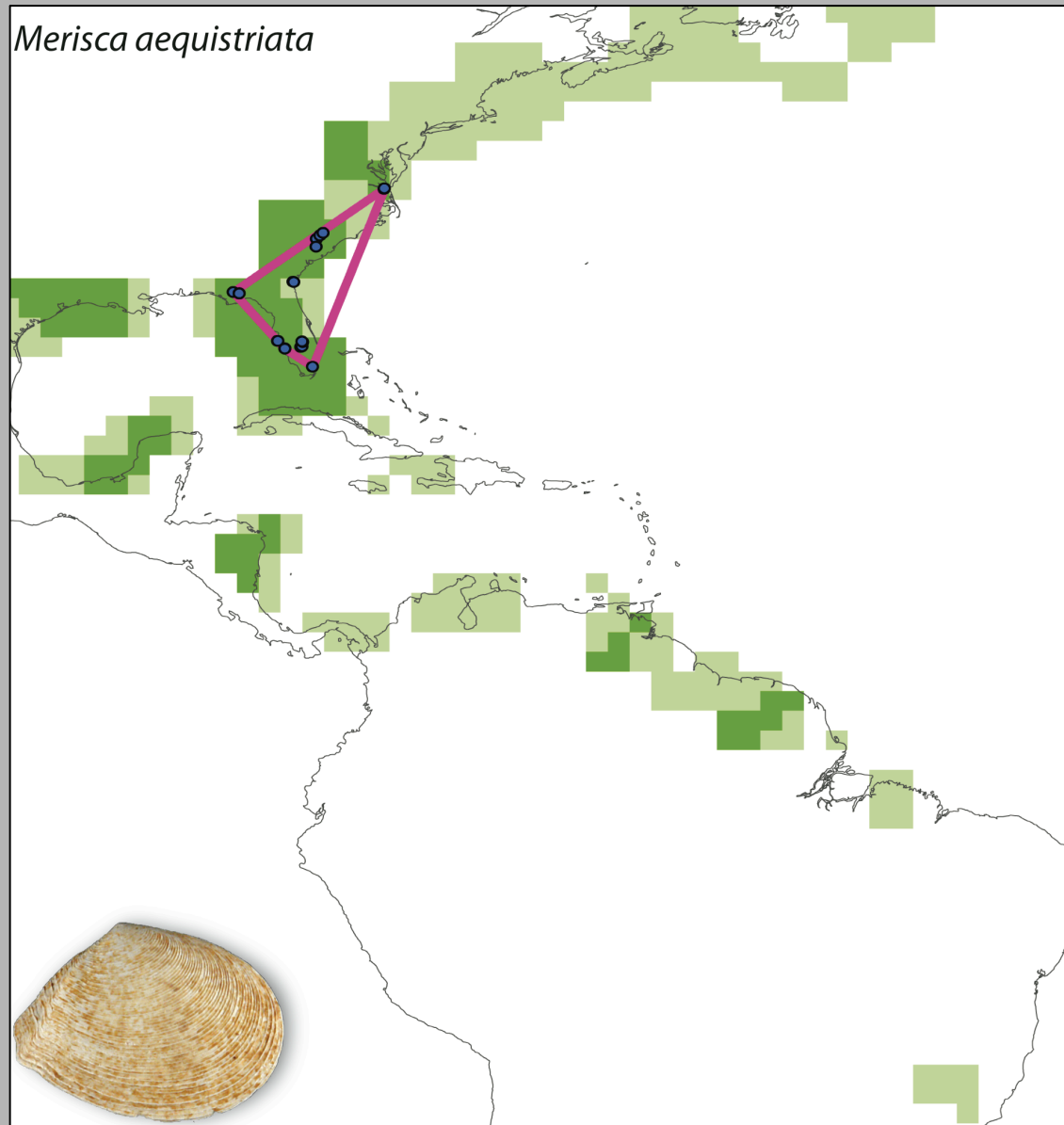


Median model

Testing whether species that survived:

- (i) had larger geographic ranges than now-extinct species
- (ii) had broader fundamental niche breadths than now-extinct species
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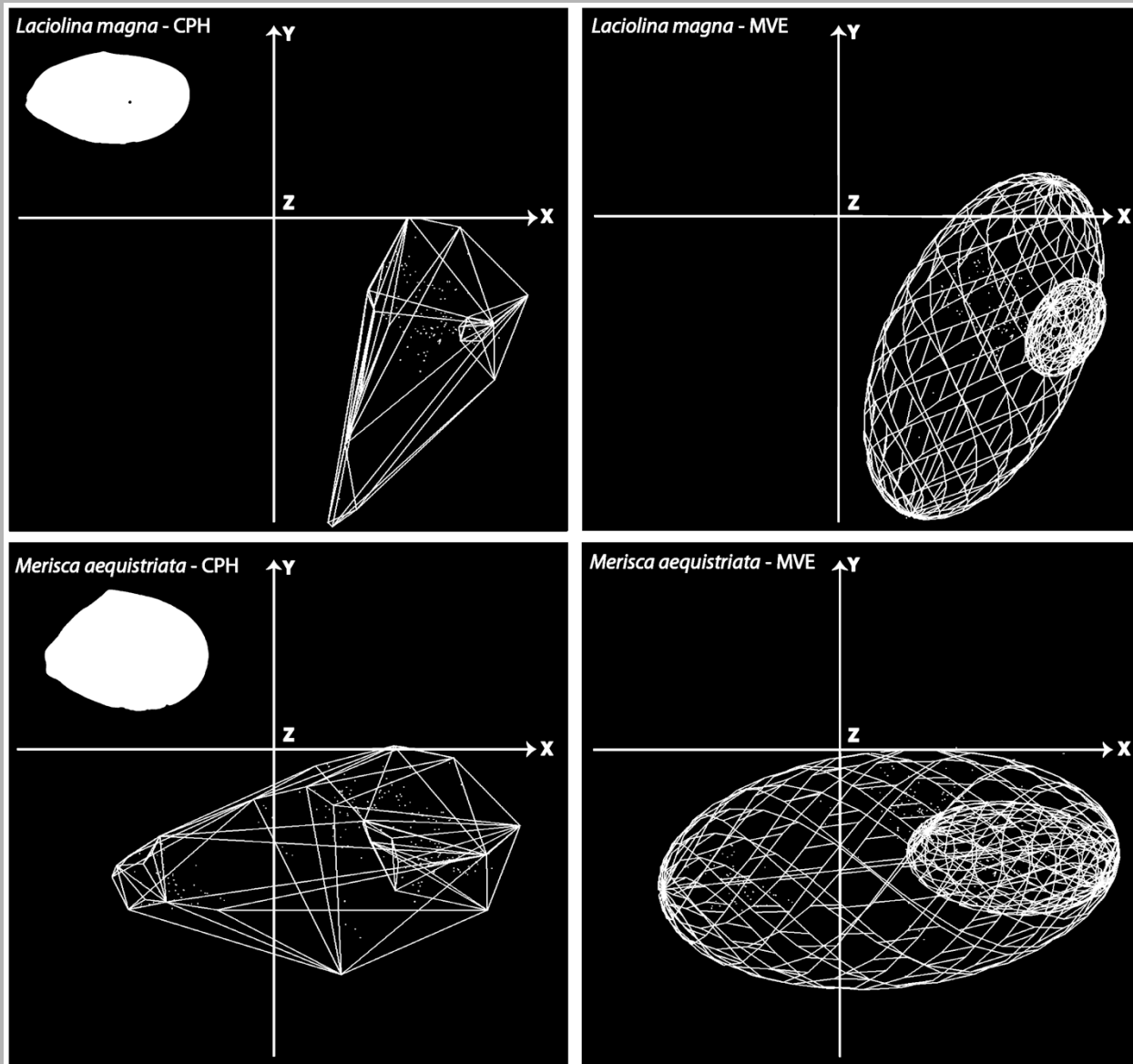
(i) Geographic range



Testing whether species that survived:

- (i) had larger geographic ranges than now-extinct species
- (ii) had broader fundamental niche breadths than now-extinct species
- (iii) occupied broader realized environmental space than now-extinct species
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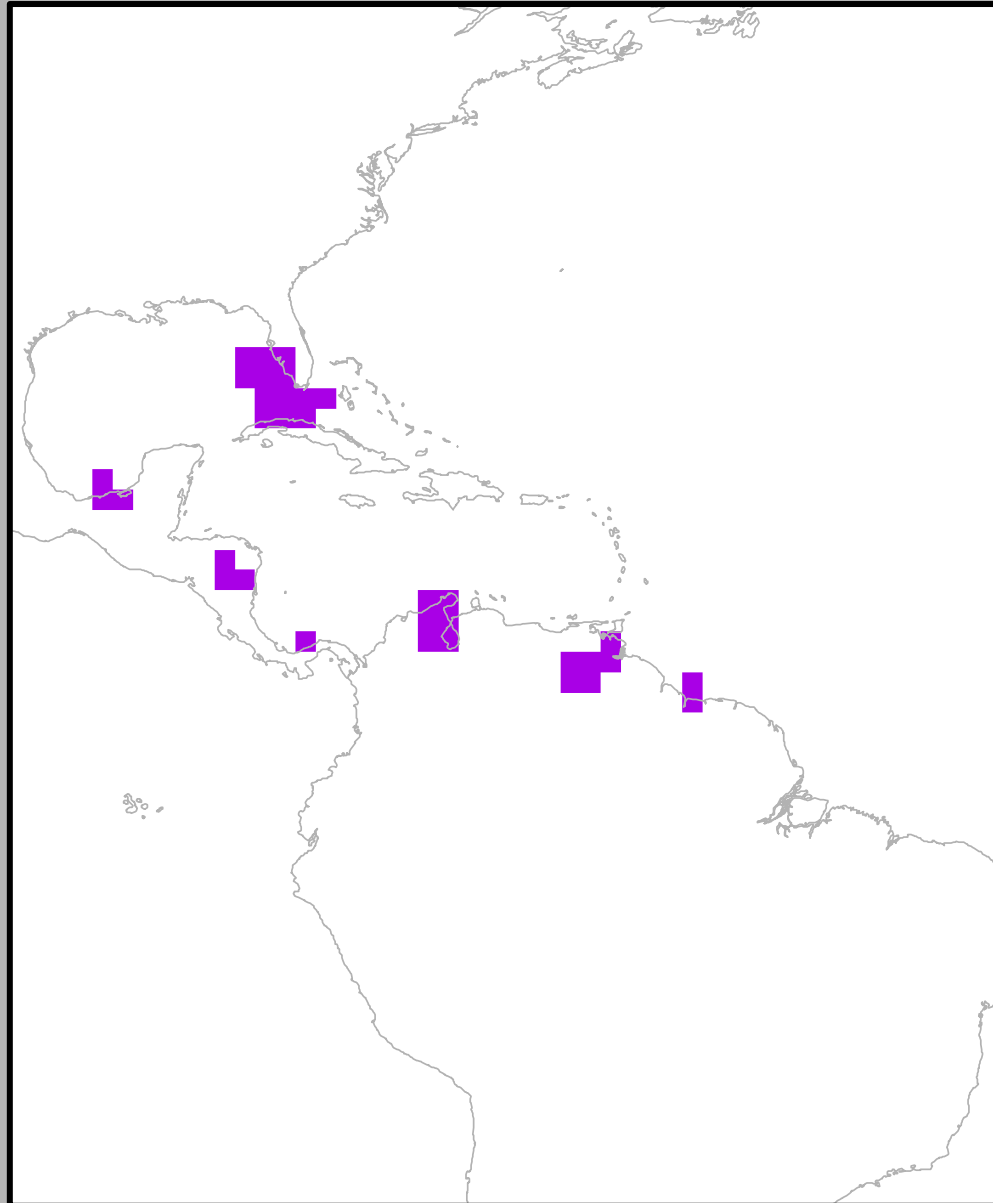
(ii-iii) Niche breadths



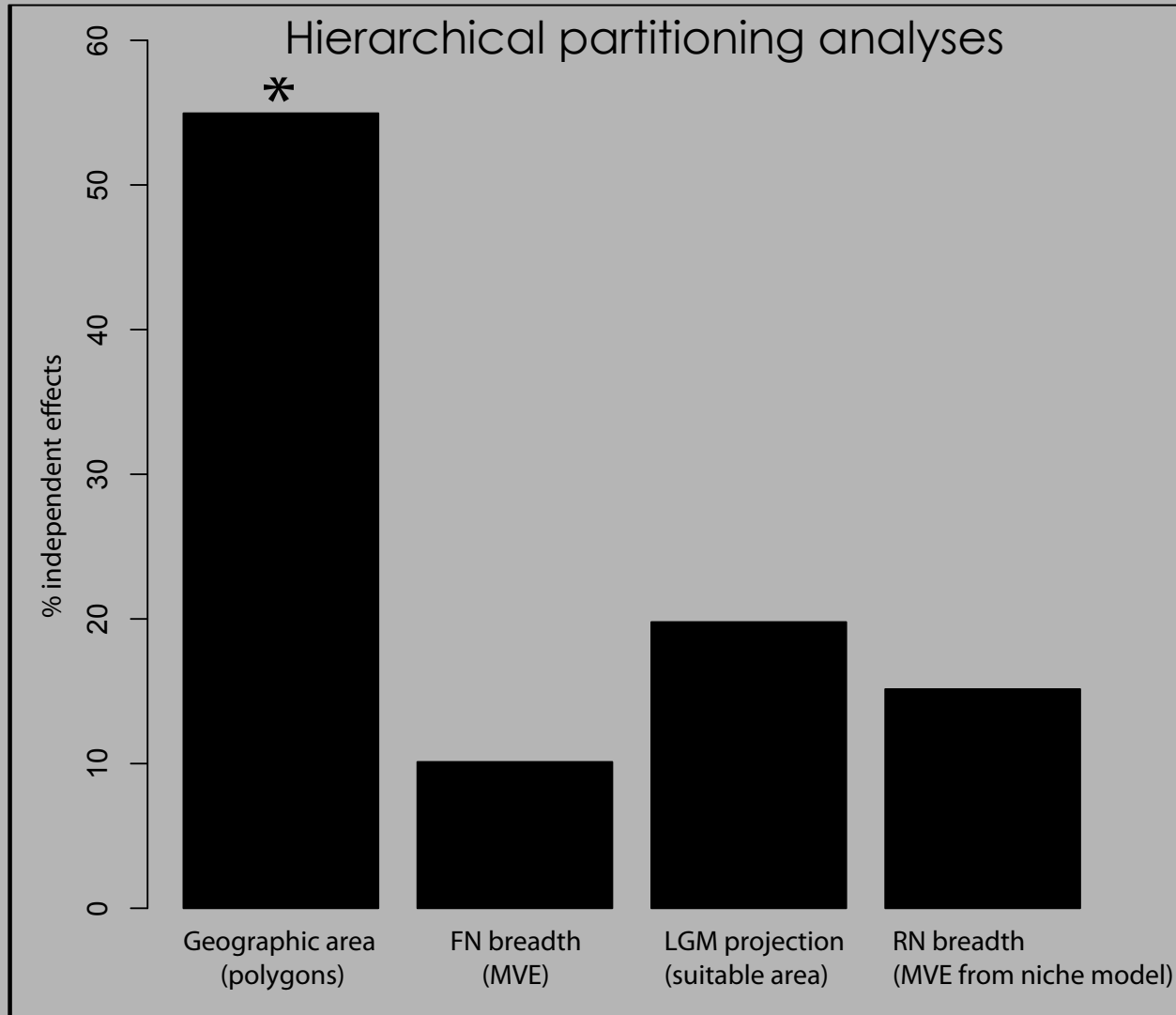
Testing whether species that survived:

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(iv) LGM Area



Extinction selectivity results

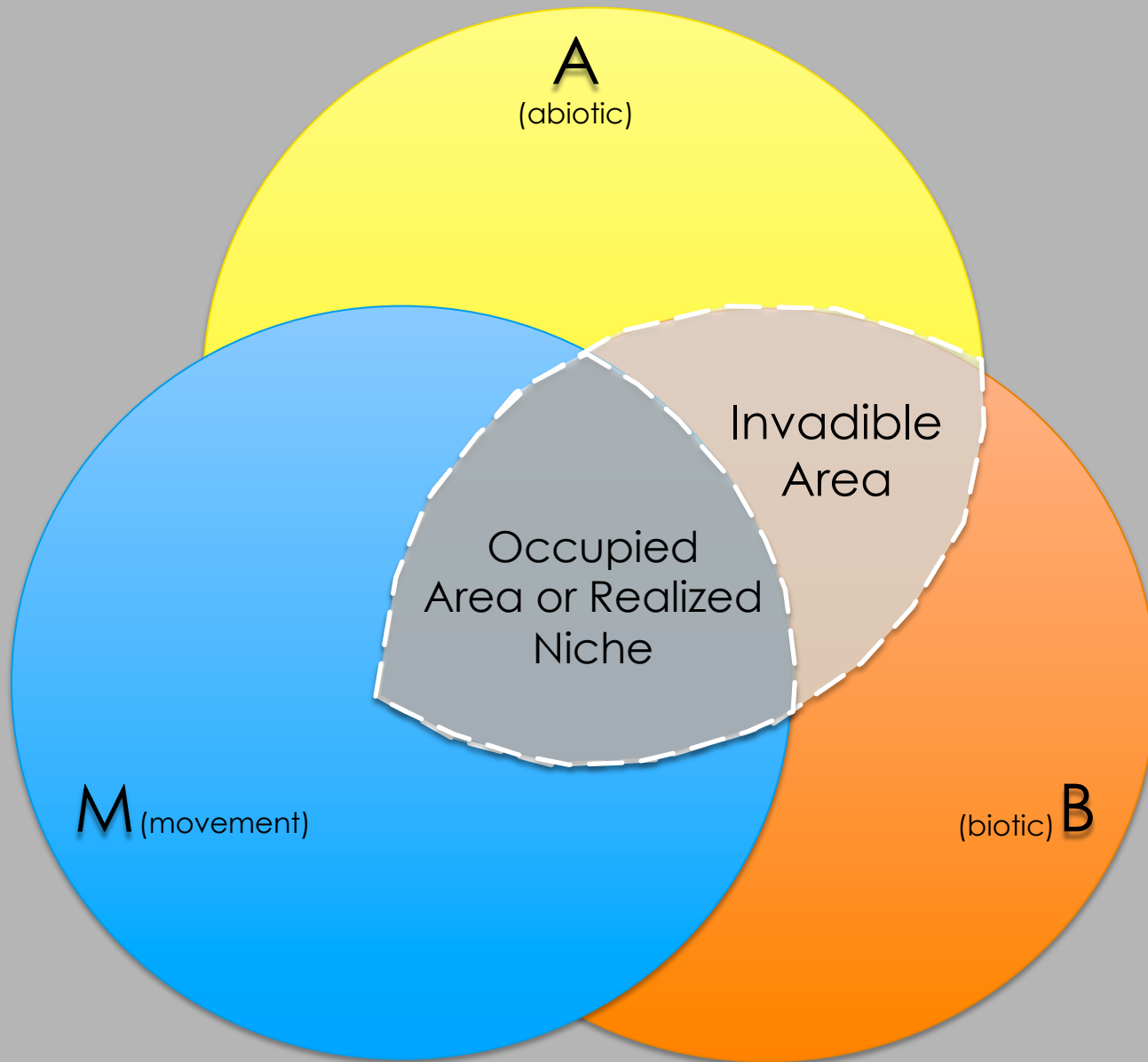


Hypothesis ii: still-extant species have larger geographic ranges than now-extinct species
Hypothesis ii: Supported

Hypothesis i: still-extant species have greater floristic richness than now-extinct species
Hypothesis i: NOT supported

Hypothesis iii: still-extant species occupied broader realized niches than now-extinct species
Hypothesis iii: Supported

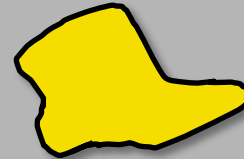
Hypothesis iv: still-extant species had more suitable area remaining during the LGM than now-extinct species
Hypothesis iv: Supported



Geography



Environment



Conclusions



Thanks to...

- Yale Institute for Biospheric Studies
- NSF Emerging Frontiers
- NSF Advancing the Digitization of Biological Collections
- NSF Systematic Biology
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- Huijie Qiao, Chinese Academy
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- Town Peterson, U of Kansas



