

Rock-Paper-Scissors, Viviparity, and Speciation: Links Between Climate Warming and Lizard Extinctions

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Outline

- **Climate Change & Global Warming – the Basics**
- Meet the “Lizards” (clade Squamata; grade “Sauria”)
- The Rock-Paper-Scissors Dynamic, Speciation, & Viviparity
- Early Clues of Unexpected Lizard Declines
- Climate Change as a Driver of Lizard Extinctions

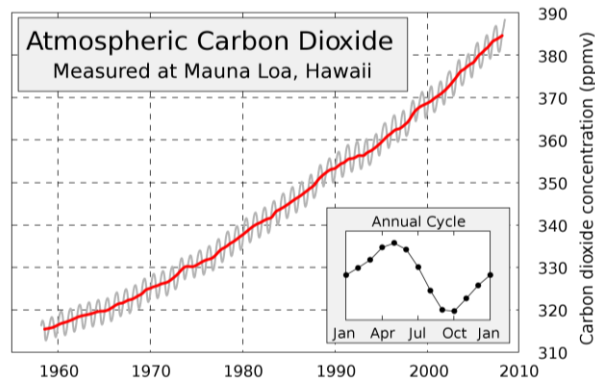


Key Climate Change Questions

1. Is climate changing in unusual ways? If it is, are humans the cause?

2. If climate is changing how large are the expected changes?

3. If climate change will cause significant damage, & is it too late, too difficult, or too expensive to correct or adapt to?



Route to a Scientific Consensus

- Heat-trapping gases & the Greenhouse Effect: Jean Fourier, 1824
- Global Warming: Svante Arrhenius, 1894; doubling CO₂ increases earth temperature 1.5 - 4.0° C
- CO₂ measurement: Roger Revelle, 1957 @ Mauna Loa, Hawaii
- The "Hockey Stick": Mann, Bradley, and Hughes, 1998 and 1999
- IPCC Reports (1990, 1995, 2001, 2007)
- Professional Scientific Societies

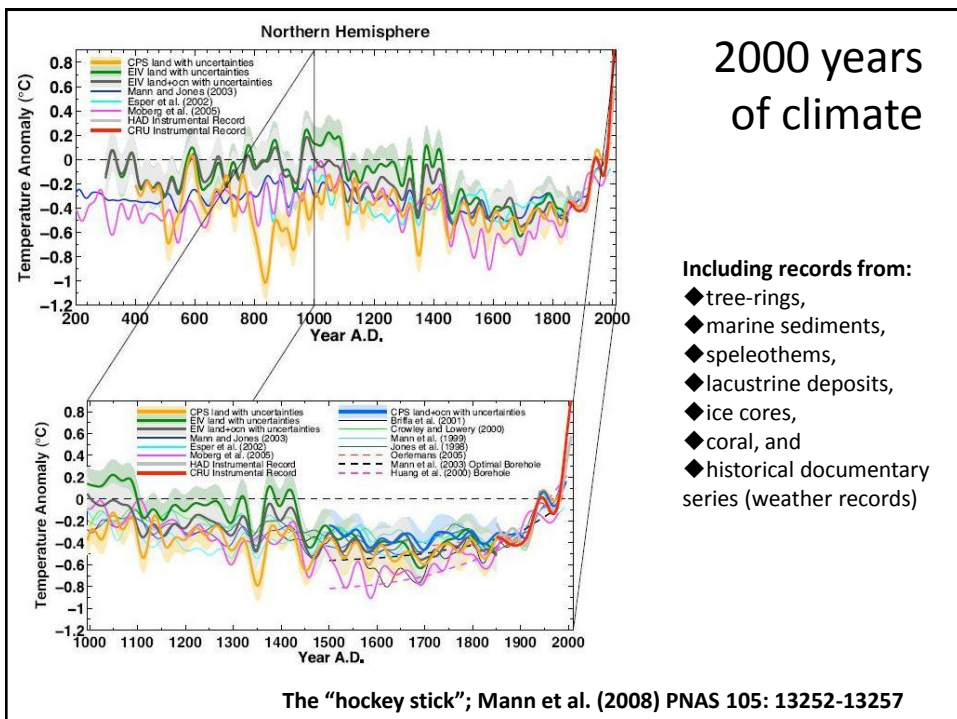
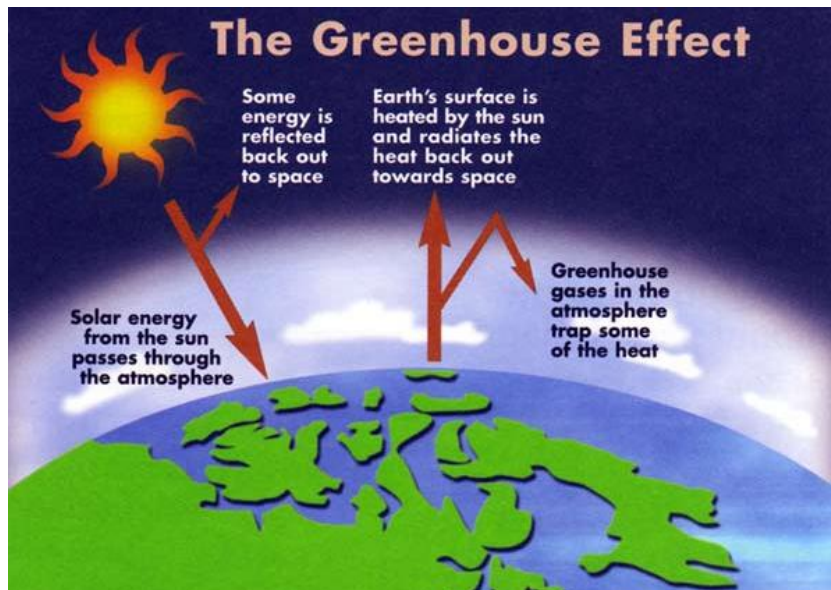
Atmosphere: Gasses With 3 or More Atoms will Trap/Hold Heat

<u>Gas</u>	<u>% (Volume)</u>	<u>Source</u>	<u>Variability</u>
Nitrogen	78.08	Biologic	permanent
Oxygen	20.95	Biologic	permanent
Argon	0.93	Radiogenic	permanent
Water (H ₂ O)	0.4 (1-4% at surface)	Evaporation	variable
Carbon dioxide (CO ₂)	0.039	Biological, industrial	increasing
Methane (CH ₄)	0.00017	Biological	increasing
Helium	0.0005	Radiogenic	escaping
N ₂ O	0.00003	Bio/industrial	increasing

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Greenhouse Effect – warming of earth's surface due to heat trapped by gases

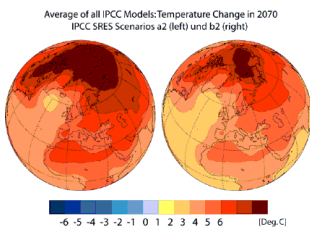


Intergovernmental Panels on Climate Change (IPCC)

- IPCC 1: 1990
- IPCC 2: 1995
- IPCC 3: 2001, Working Group 1: The Scientific Basis
 - 122 lead authors
 - 515 contributing authors
 - 420 independent reviewers
 - Several hundred government and other relevant reviewers
 - Consensus: ~ **65% probability of significant human impact**

IPCC Timeline cont'd.

- IPCC 4: **90% probability of significant human impact**



2500+ SCIENTIFIC EXPERT REVIEWERS
800+ CONTRIBUTING AUTHORS AND
450+ LEAD AUTHORS FROM
130+ COUNTRIES
6 YEARS WORK
1 REPORT

2007

The IPCC 4th Assessment Report is coming out
A picture of climate change
 the current state of understanding

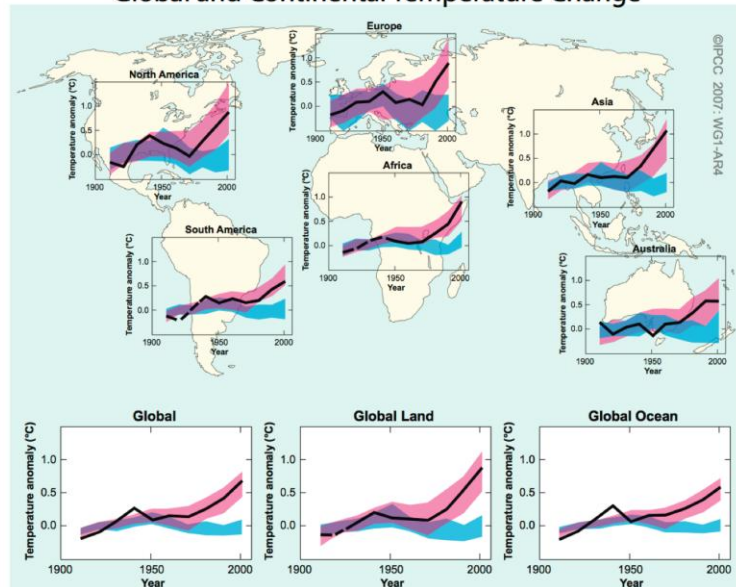


INTERGOVERNMENTAL PANEL ON CLIMATE CHANGE



Climate Models: **Observed**, **Predicted** and **Null model** (no anthropogenic effect)

Global and Continental Temperature Change



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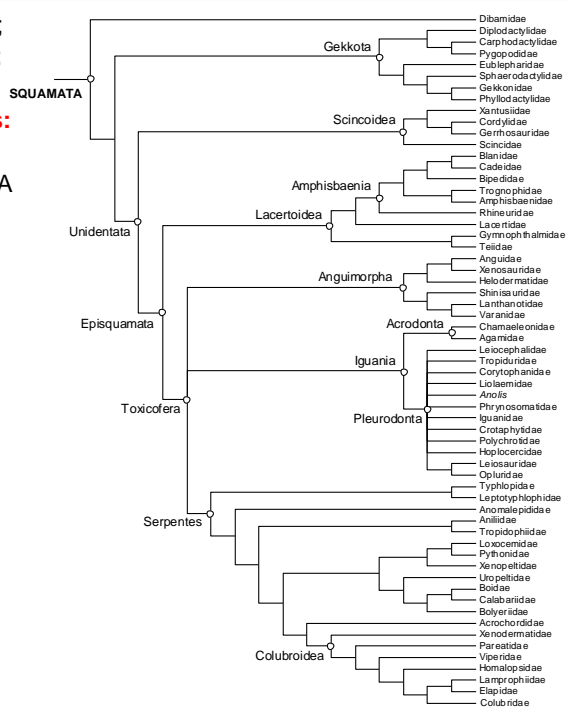
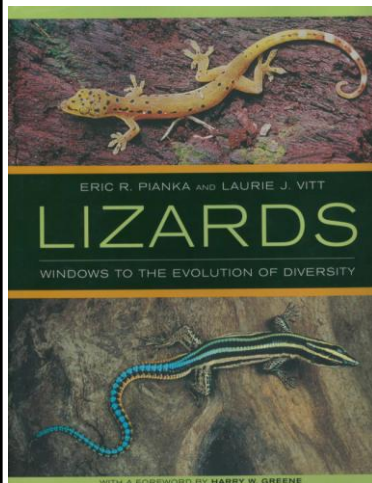
If Greenland Ice
Cap Melts



Clade **SQUAMATA** – “scaly”;
lizards, snakes, & amphisbaenians;

Most recent summary (Uetz et al.,
2011), **> 9,000 species/61 families:**

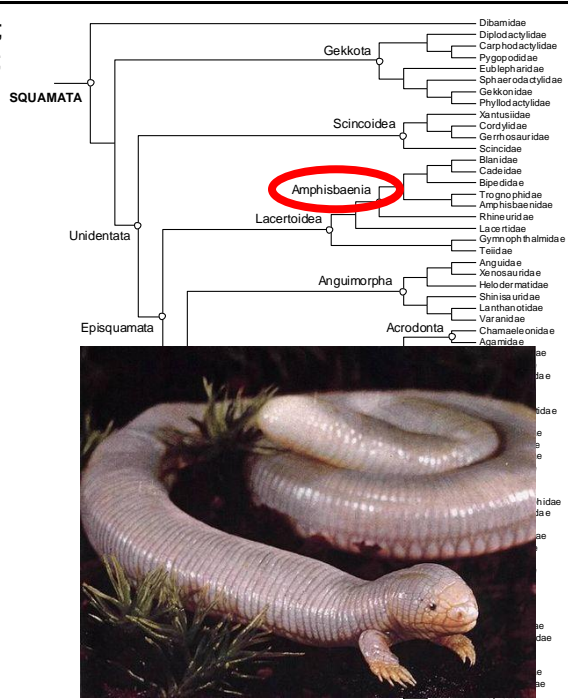
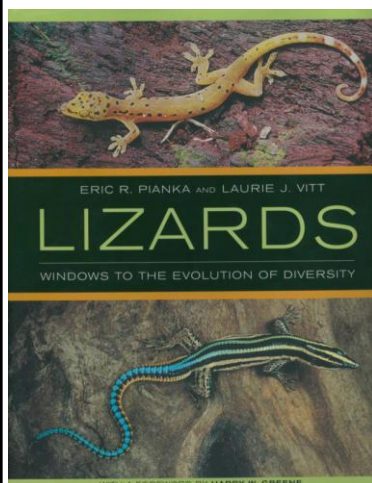
Fig. at right – best estimate of
phylogenetic relationships from DNA



Clade **SQUAMATA** – “scaly”;
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Amphisbaenians (~ 160 sp.)



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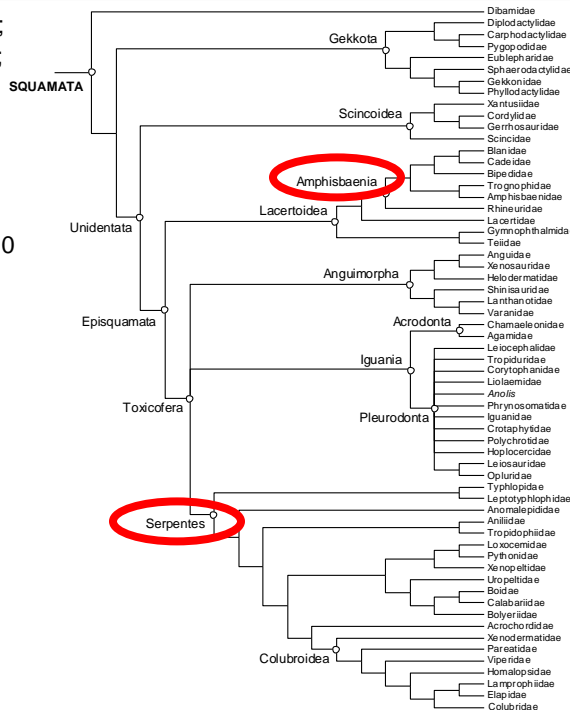
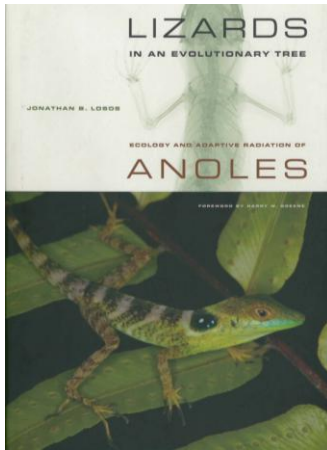
Most recent summary (Uetz et al., 2011) , > 9,000 species/61 families:

Amphisbaenians (~ 160 sp.)

Snakes (3,315 sp.)

Lizards (5,642 sp.); paraphyletic

>100 new species described in 2010



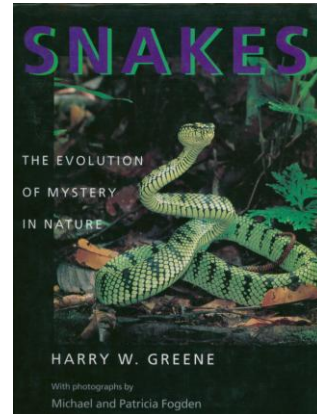
Clade **SQUAMATA** – model systems for studies of many interesting aspects of biology:

- ~150 independent origins of **toe fringes**
- ~ 107 independent origins of **viviparity**
- ~ 40 independent origins of **true parthenogenesis**
- ~ 25 independent transitions from 4-limbed lizard-like body form to **elongate limb-reduced/limbless snake-like body form**



Clade **SQUAMATA** – “scaly”; lizards, snakes, & amphisbaenians; general features of biology:

- multiple origins of **venoms & venom-delivery systems**



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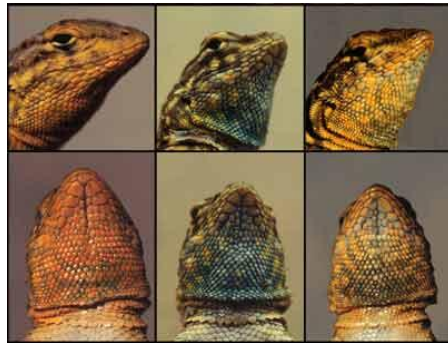
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Family: **PHRYNOSOMATIDAE** - the cast of characters: **10 general/~110+ species**; the games lizards play – the “rock-paper-scissors” mating strategy in *Uta stansburiana*



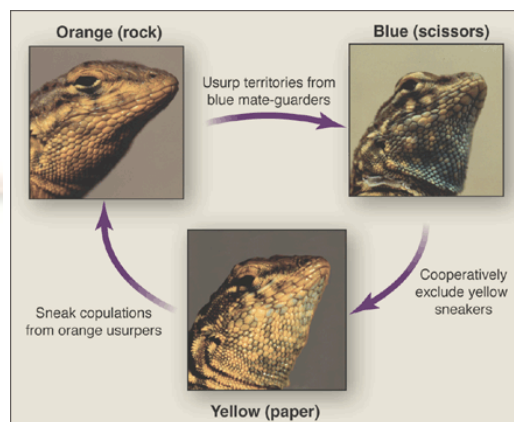
Dr. Barry Sinervo (UCSC) - **orange, blue, and yellow-throated males**; single gene locus (called OBY) controls this trait, with the **O and B alleles co-dominant to each other; both dominant to Y**; the three can co-exist because all have alternative mating strategies . . .



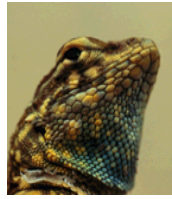
Strategy 1: Have a Lot of Territory – **O males** establish **large territories** occupied by several females, and here the more females present, the more often the male can mate;

Strategy 2: Guard Your Mate – **B males** defend **small territories** holding just a few females, but because the territories are so small, males can more carefully guard their mates; or

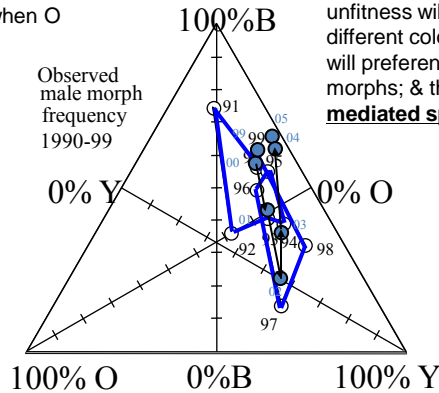
Strategy 3: Be Sneaky – **Y males** **mimic the markings and behavior of females**, and can invade (undetected) the large territories of O males, and ‘sneak’ copulations (= **cuckoldry**)



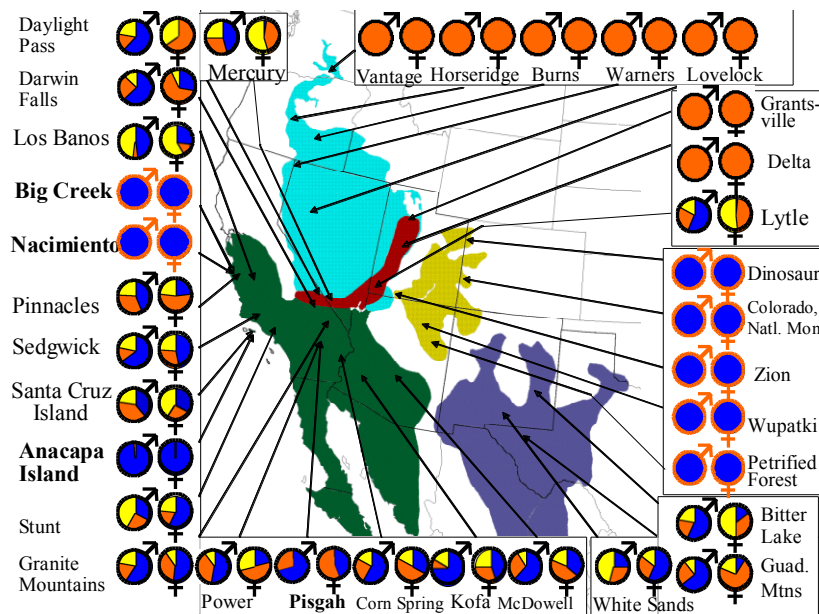
RPS cycle: classic example of **frequency-dependent selection**, each morph has a selective advantage when it is rare; O is common and can take territories from B males, but O is subject to invasion & **cuckoldry** by "sneaker males" when Y is rare; then B can take more territories from Y when O is rare, etc.

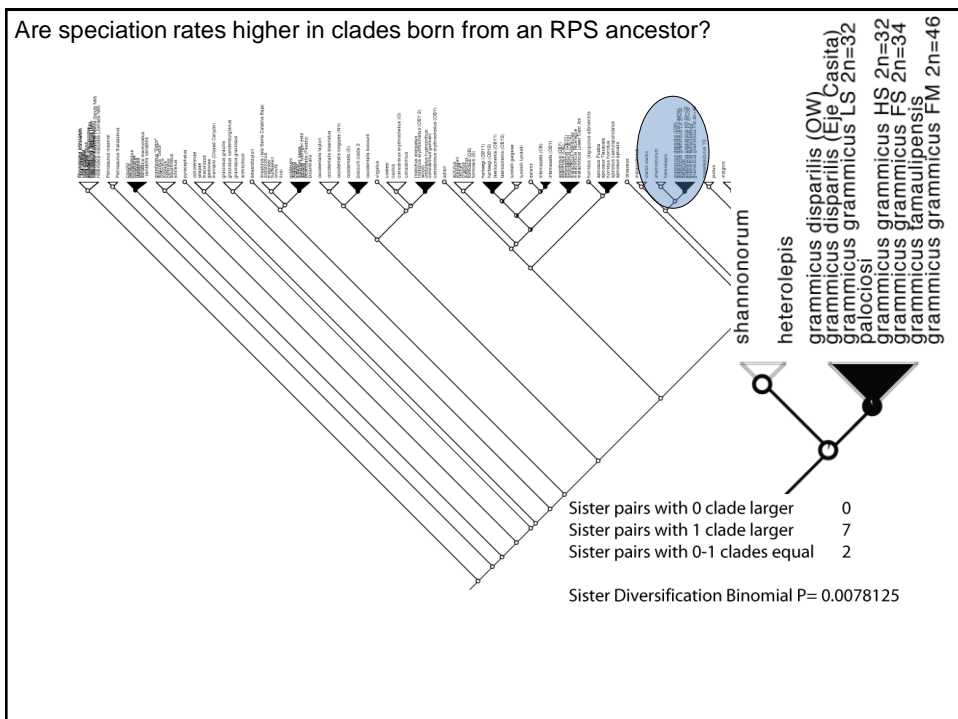


RPS cycle: **reproductive isolation can evolve within a polymorphic system like this**, given 2 other loci: one for mate choice, & one mediating social interactions (altruistic donation) – all present in *Uta* (based on gene mapping and field pedigree studies of > 7,700 lizards at Los Banos site [22 yrs]). Under some conditions hybrid unfitness will be generated when different color morphs meet – morphs will preferentially mate with their own morphs; & this **may lead to socially-mediated speciation**



Ammon Corl et al. PNAS (2010), Evolution (2009); variation in *U. stansburiana* RPS system; perhaps 15 - 20 species? (Mulcahy et al. in prep.)

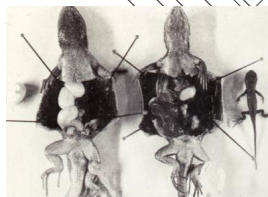




Are speciation rates higher in clades born from an RPS ancestor?

Yes, the rate of speciation is about 2-3 times higher with RPS;

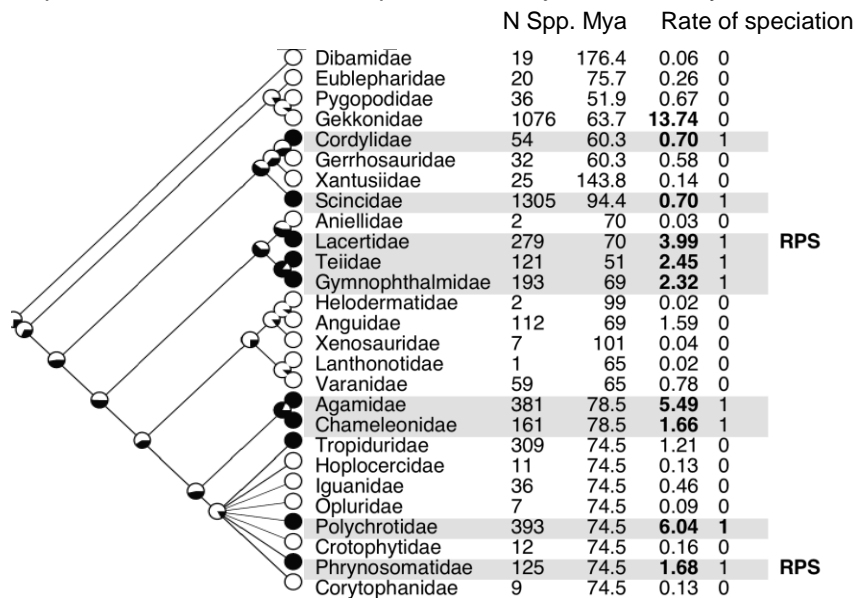
RPS dynamic drives a speciation process, and is linked to the evolution of viviparity



Sister pairs with 0 clade larger 0
Sister pairs with 1 clade larger 7
Sister pairs with 0-1 clades equal 2

Sister Diversification Binomial $P = 0.0078125$

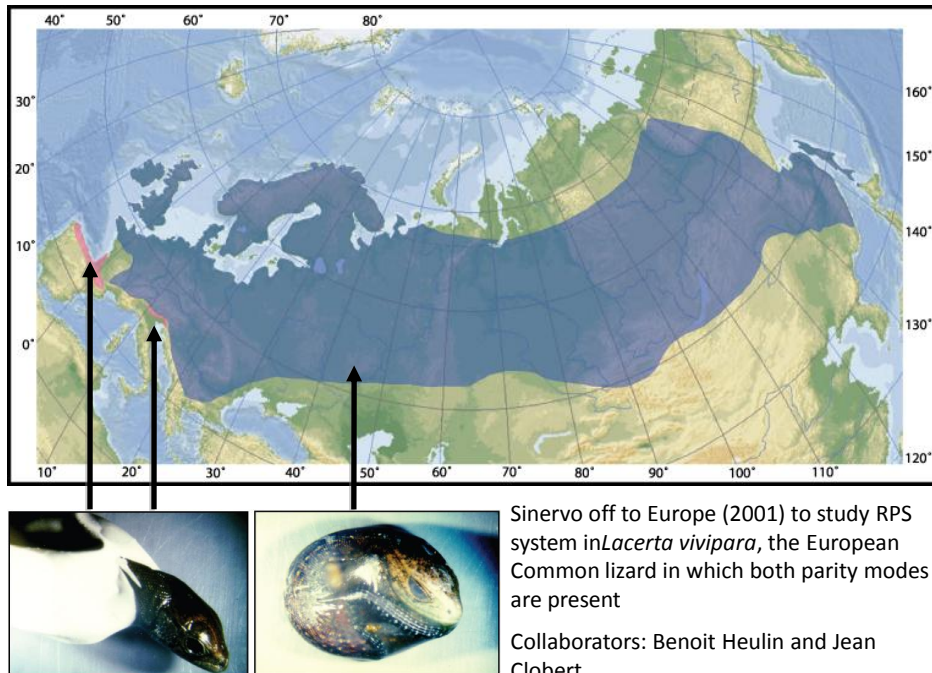
Morphs, lizard families, & RPS speciation: not just *Uta* & Phrynosomatidae




Speciation rate is **5 x higher** in polymorphic compared to monomorphic taxa

Outline

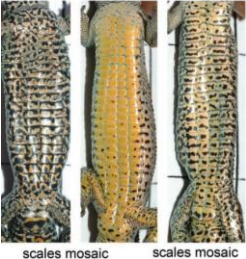
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 - *Lacerta* in Eurasia
 - *Sceloporus* in Mexico
- Climate Change as a Driver of Lizard Extinctions




oo yy ww



ow yo wy


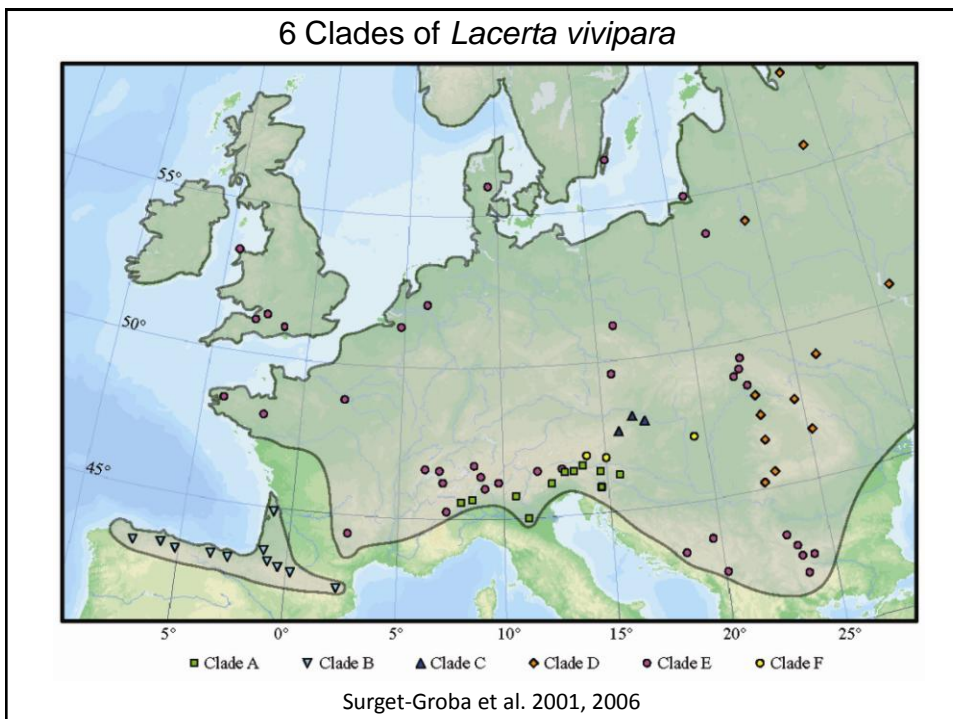


scales mosaic for w and o scales mosaic for w and y



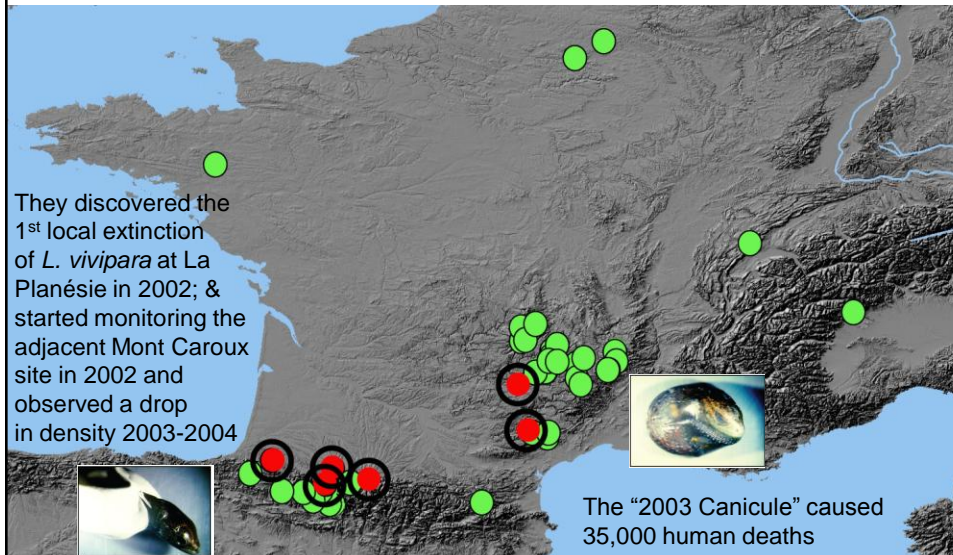
RPS in *Lacerta vivipara* parallel to *Uta stansburiana*

- ◆ Resampled sites visited by B. Heulin (1980s); 100+ sites across Europe (N=117 and counting)
- ◆ J. Clobert and Sinervo had started a global climate change project in France (focused on the Massif Central and the Pyrenees of France); a goal was to quantify adaptation of lizard populations to warming at these sites
- ◆ They immediately discovered 6 “species” (well-supported clades defined by sequence data) likely due to RPS speciation processes

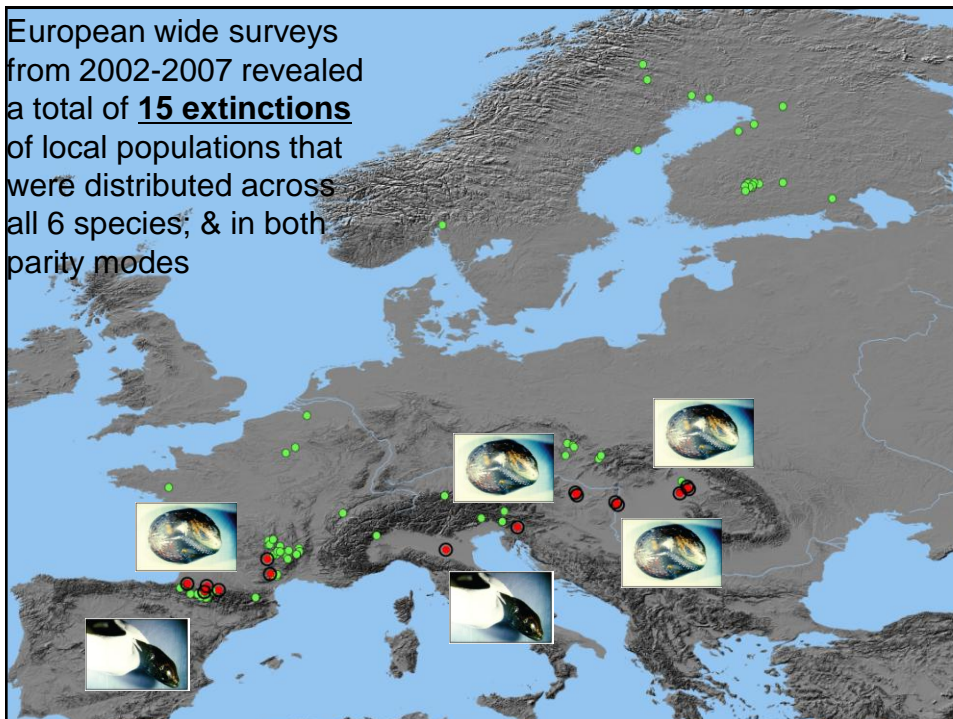



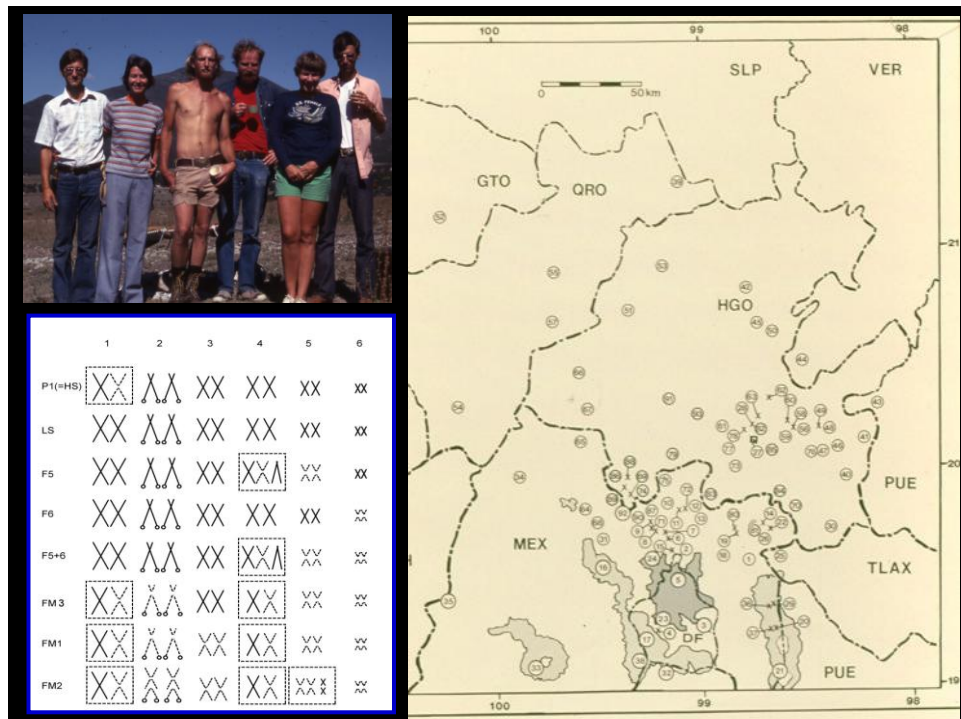
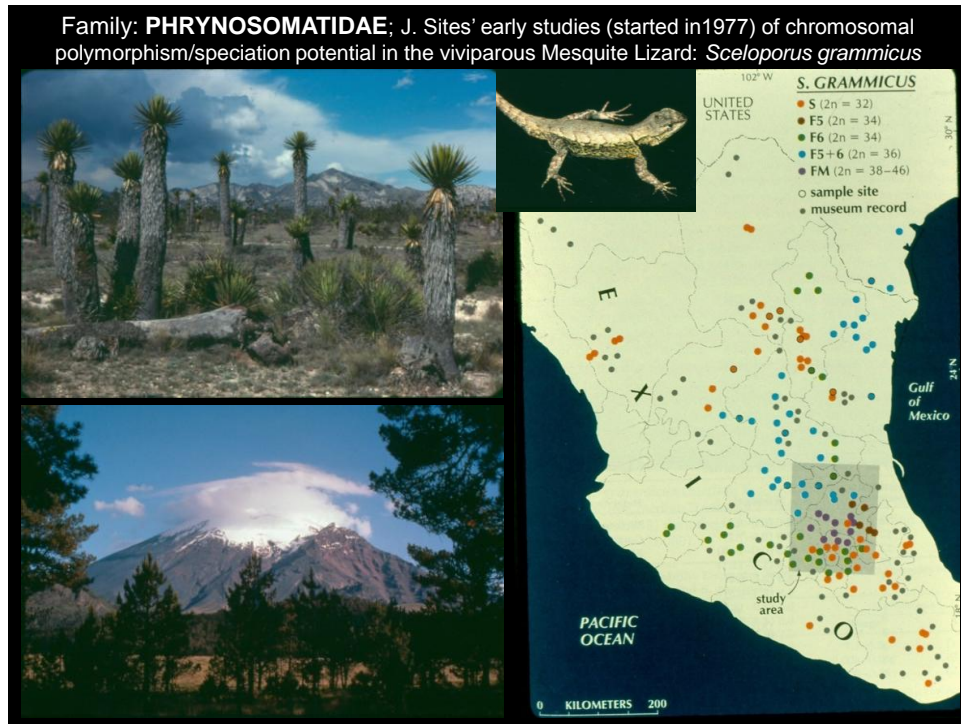
Massif Central: Sinervo located about 20 populations in the Massif Central, which he & J. Clobert started surveying every 2 years (with Virginie Lepetz and Don Miles)

Pyrenees: B. Heulin and Sinervo started detailed demographic studies in 2004 of 5 populations in the Pyrenees, and they have resurveyed 20 populations in total

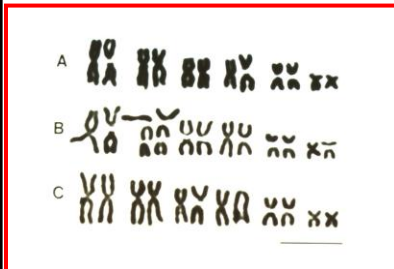


European wide surveys from 2002-2007 revealed a total of **15 extinctions** of local populations that were distributed across all 6 species; & in both parity modes





Sceloporus grammicus complex – a model in speciation research



2N = 32 – 46 (females)

- Hall (1973)
- Sites (1983)
- Porter & Sites (1986)
- Arévalo et al. (1991)



Sceloporus grammicus complex – a case of disappearing populations

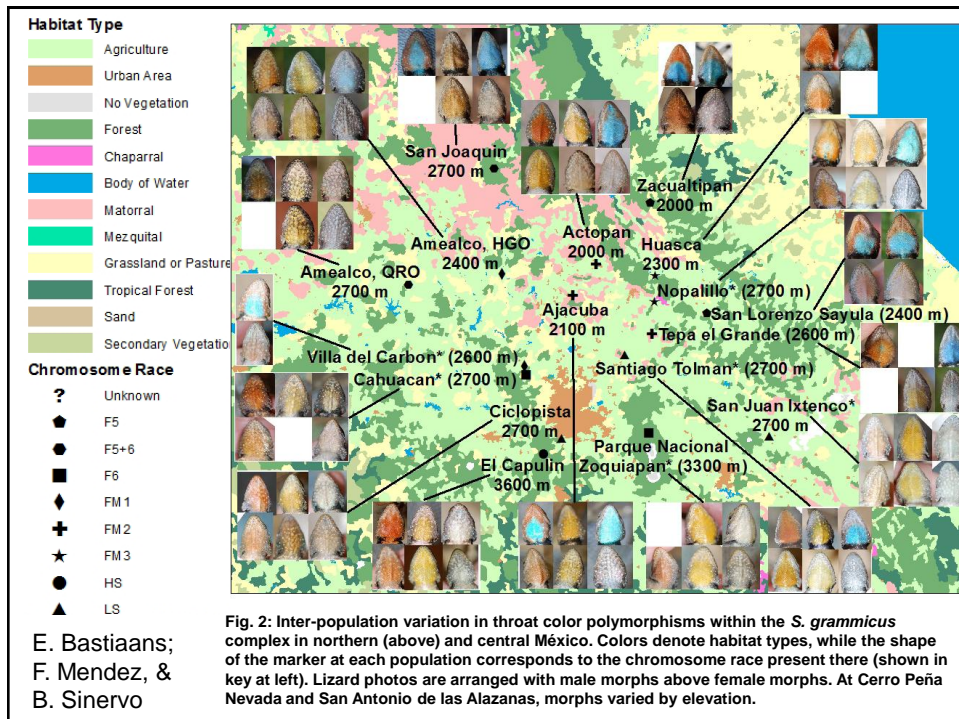
T.W. Sites, Jr.	Catalog	J. Sites, Jr.	Catalog	T.W. Sites, Jr.	Catalog
●	Texas: Hidalgo County, 8.0 mi. S of Alamo 16 + 17 July 1977	●	Mexico: Nuevo Laredo, 6.0 mi. E of San Rectorio 21 May 1978	●	Mexico: Nuevo Laredo, 12.4 mi. N of Dr. Arce on Hwy. 68 2 August 1978
54991 18P	Sceloporus variabilis juv	55405 698	Sceloporus grammicus ♀	54921 1011	Sceloporus grammicus
54990 18S	Sceloporus variabilis ♂	55406 699	"	54922 1012	"
54994 190	Sceloporus variabilis ♂ (AK-587)	55407 700	"	54923 1013	"
54993 191	Sceloporus olivaceus ♀	55408 701	"	54924 1014	"
54994 192	Sceloporus olivaceus ♀	55409 702	"	54925 1015	"
54994 193	Sceloporus olivaceus ♀ (AK-588)	55410 703	"	54926 1016	"
54994 194	Sceloporus grammicus ♀	55411 704	"	54927 1017	"
54994 195	Sceloporus grammicus ♀	55412 705	"	54928 1018	"
54994 196	Sceloporus grammicus ♂ (AK-637)	55413 706	"	54929 1019	"
54994 197	Sceloporus grammicus ♂ (AK-638)	55414 707	"	54930 1020	"
54994 198	Sceloporus grammicus ♂ (AK-639)	55415 708	"	54931 1021	"
54994 199	Sceloporus grammicus ♀ (AK-640)	55416 709	"	54932 1022	"
54994 200	Sceloporus grammicus ♂	55417 710	"	54933 1023	"
54994 201	Sceloporus grammicus ♀	55418 711	"	54934 1024	"
54994 202	Sceloporus grammicus ♀	55419 712	"	54935 1025	"
54994 203	Sceloporus grammicus ♀ (AK-632)	55420 713	"	54936 1026	"
54994 204	Sceloporus grammicus ♂ (AK-633)	55421 714	"	54937 1027	"
54994 205	Sceloporus grammicus ♂ (AK-634)	55422 715	"	54938 1028	"
54994 206	Sceloporus grammicus ♂ (AK-635)	55423 716	"	54939 1029	"
54994 207	Sceloporus grammicus ♂ (AK-636)	55424 717	"	54940 1030	"
54994 208	Sceloporus grammicus ♂ (AK-637)	55425 718	"	54941 1031	"
54994 209	Sceloporus grammicus ♀ (AK-638)	55426 719	"	54942 1032	"
54994 210	Sceloporus grammicus ♀ (AK-639)	55427 720	"	54943 1033	"
54994 211	Sceloporus grammicus ♀ (AK-640)			54944 1034	"
				54945 1035	"

Samples of J. Sites' field catalog for sampling in southern Texas & several localities in north-central Mexico (1977 – 1978)



Documenting extinctions: 2006 – 08; B. Sinervo & F. Mendez re-survey of 48 *Sceloporus* @ 200
localities in Mexico; originally sampled 1975-95; 12% of these showed local extinctions by 2009

J.W. Sites, Jr.	Catalog	* = frozen in liquid N ₂	J.W. Sites, Jr.	Catalog
Mexico: Querétaro, ca. 114 mi. E of La Laguna (W of El Lobo) on Hwy 120.	6 June 1977		MEXICO: Jalisco: Nevado de Colima, dirt road up to radio tower (on Volcan Fuego) 15 May 1989 cont'd.	
52180 1140 <i>Eumeces</i> <i>lynce</i>	adult		Sceloporus "grammicus" (adult ♂) (3500m)	
52131 1141 <i>Chenodromus rubriventris</i> (tail L = 61mm)	♀		" " (adult ♂) (3500m)	
52132 1142 <i>Eleutherodactylus decoratus</i>			" " (adult ♀) (3500m)	
52133 1143 " "			" " (adult ♀) (3500m)	
Mexico: Querétaro, 114 mi. E of La Laguna (W of El Lobo) on Hwy 120.	7 June 1977		39765 1930 " " (adult ♀) (3500m)	
1144 <i>Sceloporus grammicus</i> juv *			39766 1932 " " (adult ♀) (3500m)	
1145 " " " *			39767 1933 " " (adult ♂) (3200m)	
1146 " " " *			39768 1934 " " (adult ♂) (3300m)	
1147 " " " *			1935 " " (adult ♂) (3100m)	
1148 " " " *			1936 " " (adult ♂) (3100m)	
1149 " " " *			39781 1937 " " (adult ♀) (3200m)	
1149 " " " *			39790 1938 " " (adult ♀) (3000m)	
52134 1150 " " adult ♂ + ♀			39791 1939 " " (adult ♀) (3000m)	
52135 1151 " " " " have solid blue throats			39792 1940 " " (adult ♀) (3300m)	
52136 1152 " " " " " " " "			MEXICO: Jalisco: Sierra del Tigre, ~3.0 km N of El Terrero (Carretera Tiquipán-Cd. General)	
Mexico: Querétaro, 2.3 mi. W of El Lobo, Hwy 120	7 June 1977		2,050 m.	
1153 <i>Sceloporus grammicus</i> juv *			16 May 1989	
1154 " " juv *			39780 1941 <i>Sceloporus heterolepis</i> (newborn ♀)	
1155 " " juv *			1942 " " (newborn ♂)	
1156 " " juv *			1943 " " (newborn ♀)	
1157 " " juv *			1944 " " (adult ♂) (39781)	
1158 <i>Sceloporus leucotis</i> (ground ♀)			1945 " " (adult ♀) (39781)	
1159 <i>Sceloporus leucotis</i> (♂ + ♀) (solid blue throat)			1946 " " (adult ♀) (39781)	
Mexico: Querétaro, 1.2 mi. W of El Lobo on Hwy 120	7 June 1977		1947 <i>Sceloporus torquatus</i> (newborn ♀)	
1159 <i>Sceloporus grammicus</i> juv *			1948 " " (adult ♂)	
1160 " " " *				



E. Bastiaans;
 F. Mendez, &
 B. Sinervo

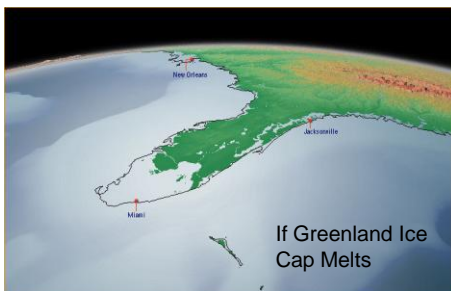
Sceloporus chaneysi (dimorphic with respect to alleles), in *S. aeneus* group



Beth Bastiaans resurveyed many of the 2007 sites for *S. chaneysi*, and has found new extinctions. The related *S. goldmani* (viviparous) will almost certainly be totally extinct within 10 years

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Global climate change (GCC) affects organisms in all biomes & ecosystems; with sufficient time/dispersal capability, species may adjust by:

- shifting distributions to more favorable thermal environments
- adapting to modified local environments by behavioral/physiological plasticity, or via directional selection (given sufficient h^2); or
- failing to do these things → demographic collapse & extinction



Erosion of Lizard Diversity by Climate Change and Altered Thermal Niches

Barry Sinervo,^{1,15*} Fausto Méndez-de-la-Cruz,² Donald B. Miles,^{3,15} Benoit Heulin,⁴ Elizabeth Bastiaans,¹ Maricela Villagrán-Santa Cruz,⁵ Rafael Lara-Resendiz,² Norberto Martínez-Méndez,⁶ Martha Lucía Calderón-Espinosa,⁶ Rubi Nelsi Meza-Lázaro,² Héctor Gadsden,⁷ Luciano Javier Avila,⁸ Mariana Morando,⁹ Ignacio J. De la Riva,⁹ Pedro Victoriano Sepúlveda,¹⁰ Carlos Frederico Duarte Rocha,¹¹ Nora Ibargüengoytia,¹² César Aguilar Puntriano,¹³ Manuel Massot,¹⁴ Virginie Lepetz,^{15*} Tuula A. Oksanen,¹⁶ David G. Chapple,¹⁷ Aaron M. Bauer,¹⁸ William R. Branch,¹⁹ Jean Clobert,¹⁵ Jack W. Sites Jr.²⁰

It is predicted that climate change will cause species extinctions and distributional shifts in coming decades, but data to validate these predictions are relatively scarce. Here, we compare recent and historical surveys for 48 Mexican lizard species at 200 sites. Since 1975, 12% of local populations have gone extinct. We verified physiological models of extinction risk with observed local extinctions and extended projections worldwide. Since 1975, we estimate that 4% of local populations have gone extinct worldwide, but by 2080 local extinctions are projected to reach 39% worldwide, and species extinctions may reach 20%. Global extinction projections were validated with local extinctions observed from 1975 to 2009 for regional biotas on four other continents, suggesting that lizards have already crossed a threshold for extinctions caused by climate change.

Current forecasting models are not calibrated with actual extinction events (range shifts, species/area relationships, etc.); **empirical validation of global extinction forecasts requires three forms of evidence:**

- **1 – actual extinctions should be linked to macroclimate patterns and validated to biophysical thermal causes arising from microclimate.**
- 2 – the pace of climate change should compromise thermal adaptation, such that evolutionary rates lag behind global warming due to constraints on thermal physiology.
- 3 – extinctions due to climate should be global in extent spanning continents, *but the models should also be able to predict extinctions at precise local scales.*



Erosion of Lizard Diversity by Climate Change and Altered Thermal Niches

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1 – actual extinctions should be linked to macroclimate and validated to biophysical thermal causes arising from microclimate

Many lizards are **heliotherms**, but **preferred body temperature** (T_b) cannot \gg **critical thermal maximum** (T_{max}); = **lethal**

lizards retreat to cool places, but **hours of restriction** (h_r) limit foraging, etc., \rightarrow constrain growth, maintenance, & reproduction \rightarrow undermining population growth rates



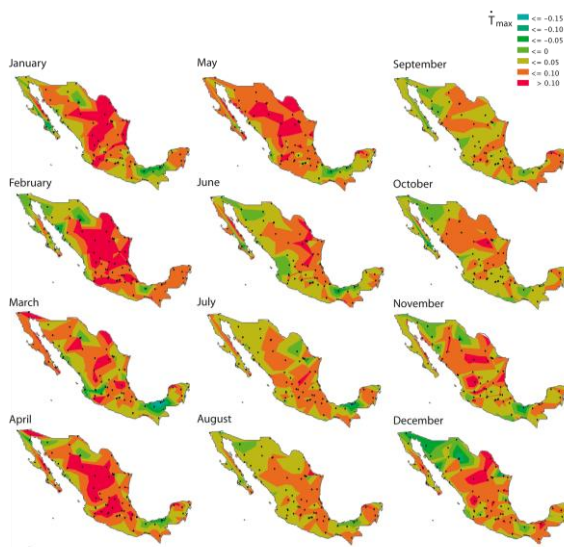
Rate of change in maximum ambient air temperature (ΔT_{max})
data from 99 Mexican weather stations, to construct
climate surfaces for 1973 – 2008; results:

Red = areas where $\Delta T = 3.5^\circ\text{C}$ from 1975 – 2010, but at some sites during breeding ΔT has increased by 4 - 7°C;

Note that:

- ΔT highest for Jan – May
- fastest rates in central & northern MX, & high elevation
- significant **correlation** between ΔT_{max} during W/S breeding, & **extinctions** of local *Sceloporus* populations

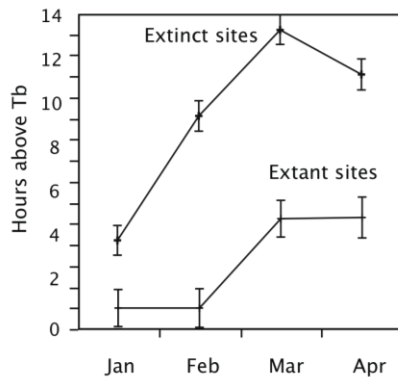
= “**macroclimate pattern**”



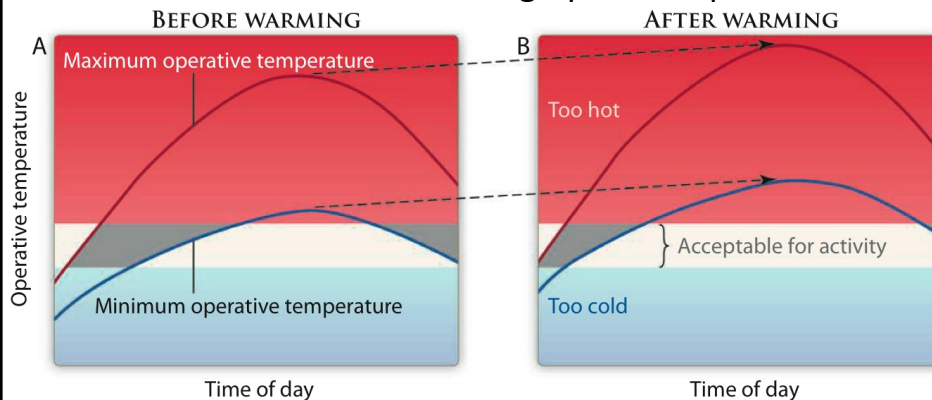
Yucatan ground truth study (Dec 2008-Apr 2009):

- Deploy lizard models hooked up to Hobotemps
- Measure the number of hours T_b is above $T_{\text{preferred}}$ at two extinct and two persistent sites (*Sceloporus serrifer*, viviparous, Yucatan)
- Linear relationship can be used to predict $h_{\text{restriction}}$:

$$h_r [T_b > T_{\text{preferred}}] = \text{slope} \times (T_{\text{max}} - T_{\text{preferred}}) + \text{intercept}$$
- h_r significantly higher in Mar-April at extinction vs extant sites; ($t = 9.26$; $P < 0.0001$)



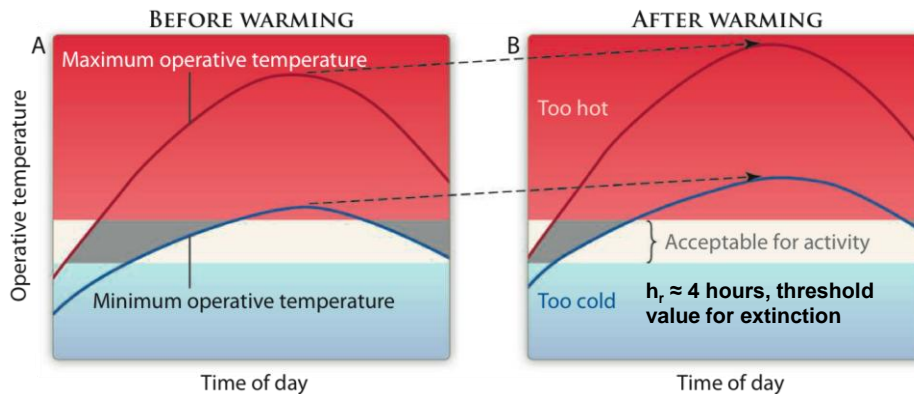
Hypothesis: Hours of restriction (h_r) During Breeding is the cause of demographic collapse



With warming, **restriction on activity is so severe** that lizards have to retreat shortly after emergence; **insufficient time to accumulate enough calories to develop a clutch**

Figure from: Huey et al. (2010) Are Lizards Toast? Science 325. Photo: Dr. Fausto Mendez De la Cruz

S. serrifer: relationship between h_r as a function of T_{\max} provides a **general formula** [$h_r = 6.12 \times 0.74(T_{\max} - T_b)$] **for predicting extinctions**



Best fit: observed vs predicted extinctions for Mexican *Sceloporus*; $h_r > 3.85$ hours

Significant for both parity modes: **Oviparous** – $\chi^2 = 49.0$; $P < 0.001$

Viviparous – $\chi^2 = 4.2$; $P < 0.04$

What approach should be used? Empirical validation of global extinction forecasts requires three forms of evidence:

- 1 – actual extinctions should be linked to macroclimate and validated to biophysical thermal causes arising from microclimate.
- **2 – The pace of climate change should compromise thermal adaptation, such that evolutionary rates lag behind global warming due to constraints on thermal physiology.**
- 3 – extinctions due to climate should be global in extent spanning continents, *but the models should also be able to predict extinctions at precise local scales.*



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Lizards cannot evolve rapidly enough to track pace of regional climate change:

- PIC test shows that T_b and T_{max} are correlated, which constrains adaptation;
- Phrynosomatidae – **shift in T_b of 1°C yields only a 0.5°C correlated response in T_{max}** ;
- the latter attribute (T_{max}) **will not evolve fast enough to keep pace with selection for higher T_b**



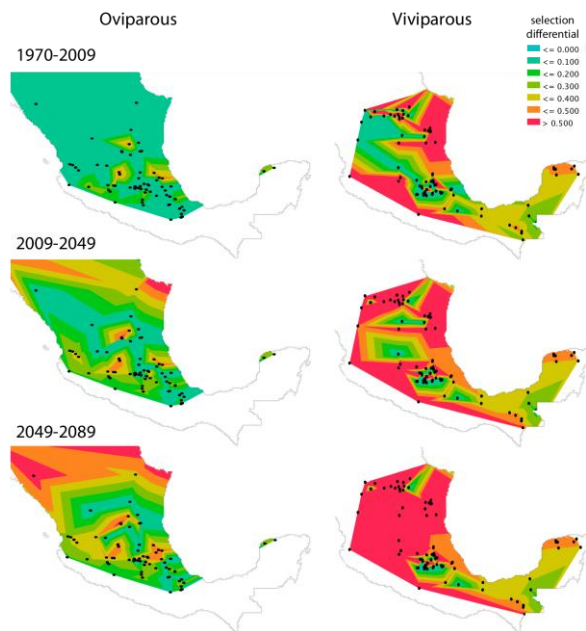
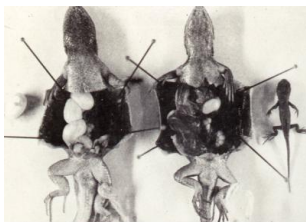
Selection: $R = h^2s$; where **$h^2 = \sim 0.17$** in *S. occidentalis* (T_b heritability)

Maps - **sustained selection differentials** needed per year for T_b to keep pace with warming;

adaptation in T_b is not likely due to low h^2

oviparous – warming alters hydric soil environments, clutches overheat

viviparous – heat loads to adults compromises behavioral strategies for maintaining body temperature suitable for embryonic development



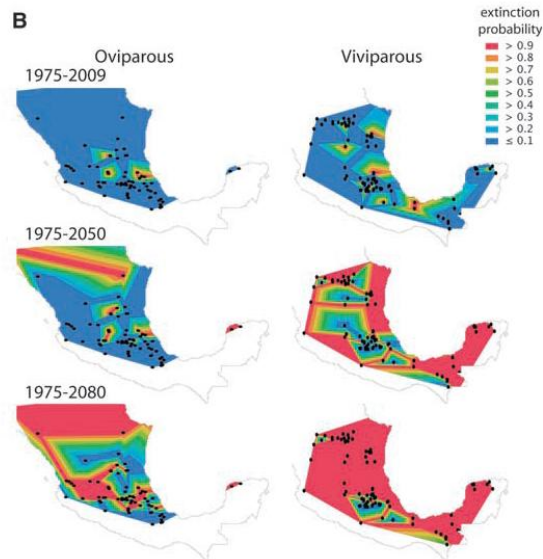
Selection: $R = h^2s$; where $h^2 = \sim 0.17$ in *S. occidentalis* (T_b heritability)

Maps – extinction surfaces:

Viviparous: 56% population extinctions by 2050; 66% by 2080

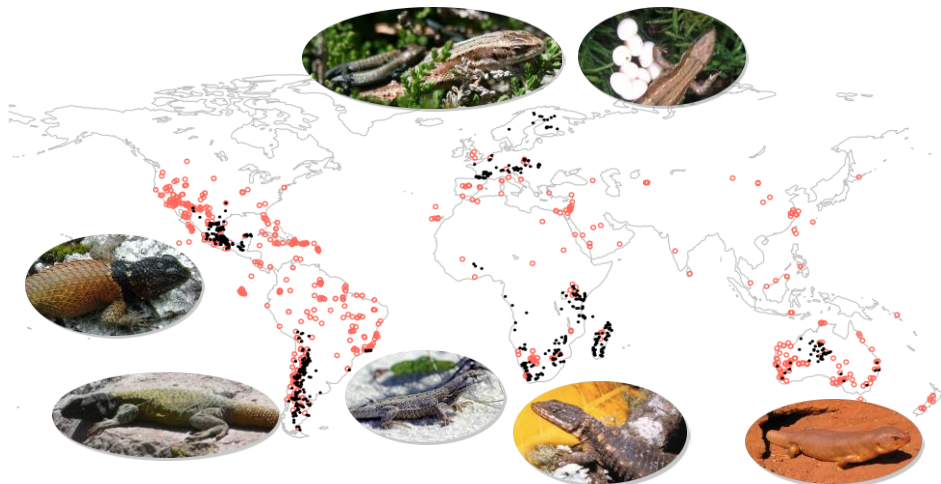
Oviparous: 46% and 61% population extinctions by 2050 and 2080, respectively

Combined: 58% population extinctions of Mexican *Sceloporus* by 2080



The Global Extinction Model:

- 1) T_{max} from WorldClim.org at 10 arc-sec resolution
- 2) T_b parameterized from **N=1216 georeferenced sites, red dots**
- 3) T_b (simulated, climate envelope for high range)
- 4) $h_r = 4.55$, average across all lizard families
- 5) h_r fitted for each of **34 families from 1975 distribution limits**
- 6) local extinctions validated for 6 families on 5 continents, **black dots**



Reliability of predicted vs. observed contemporary extinctions: **72 % accurate** (weighted R²)

Region/Taxon	R-Squared	N sites
Africa (167 sites)+Madagascar (2 sites): Gerrhosauridae, Cordylidae, Chamaeleonidae, Scincidae, Gekkonidae	0.98	169
Europe: Lacertidae, <i>Lacerta vivipara</i>	0.53	46
South America: Liolaemidae, <i>Liolaemus</i> and <i>Phymaturus</i> spp.	0.53	128
Australia: Scincidae, <i>Liopholis</i> spp.	1.00	23
Australia: Scincidae, <i>Liopholis kintorei</i>	0.19	29

1. Where we see errors, drought due to climate warming has caused extinctions not explained by temperature
1. Also ¼ of Mexican extinctions were not predicted by thermal extinction, but in 6 of 8 cases a competitor had expanded its range upwards due to climate warming

What approach should be used? Empirical validation of global extinction forecasts requires three forms of evidence:

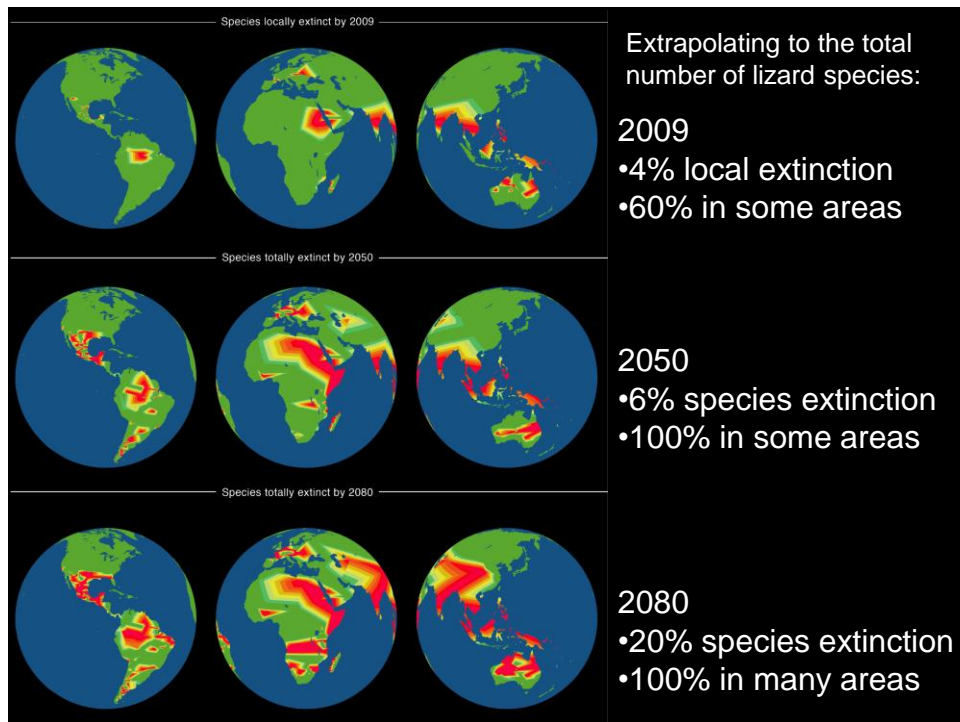
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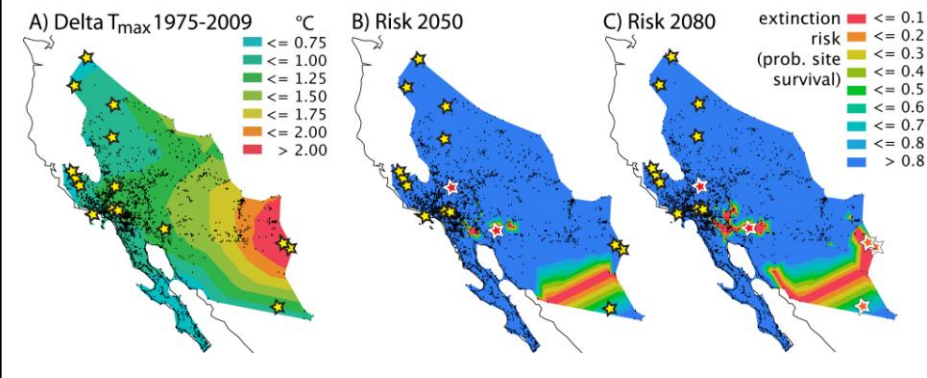
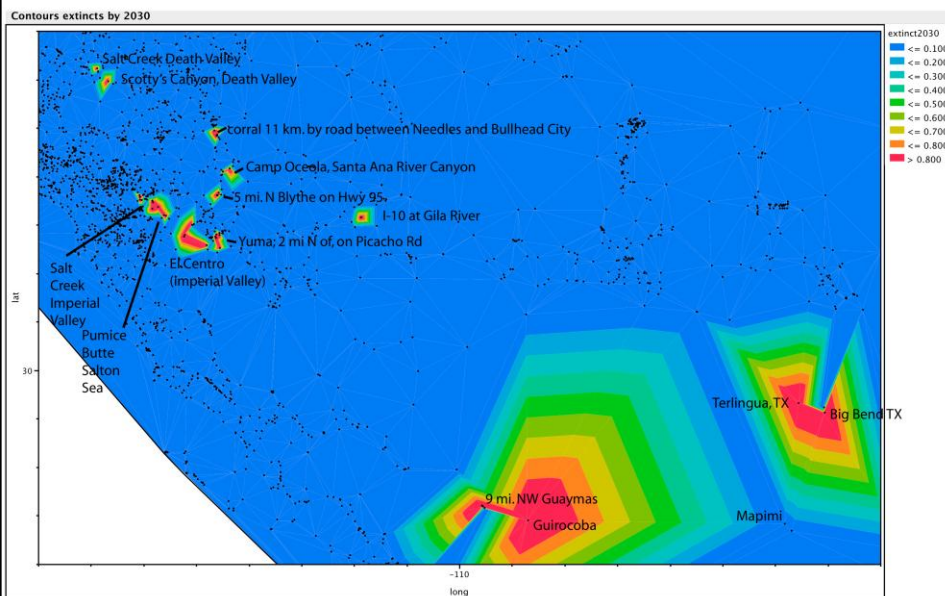


Research at demographic timescales

- *Uta stansburiana*: Can lizards evolve out of the frying pan: Evolution of T_b unlikely, maybe habitat preference?
- *Urosaurus graciosus*: Very high local extinction in 2010, Donald Miles, Sinervo unpub. data.
- *Lacerta vivipara*: tracking the progress of extinctions (periodic resurveys from 1992-2010)
- *Iberolacerta* spp in Europe: ongoing extinctions in endemic montane species that we are tracking
- We are also tracking the extinctions in Mexico

Uta stansburiana:

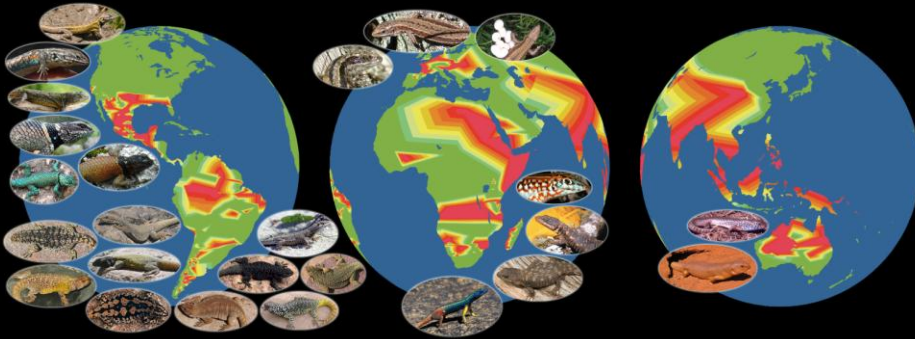
1. Sample 15 sites across the west
2. Estimate climate change evolution at Los Banos (N=32,000, pedigree 1989-2010) to estimate $R=h^2s$
3. Survey extinctions. We have verified predicted extinctions at Death Valley and Pilot Knob. Others: TX and Mexico are currently being surveyed

*Uta stansburiana* predicted extinctions 2010-2030

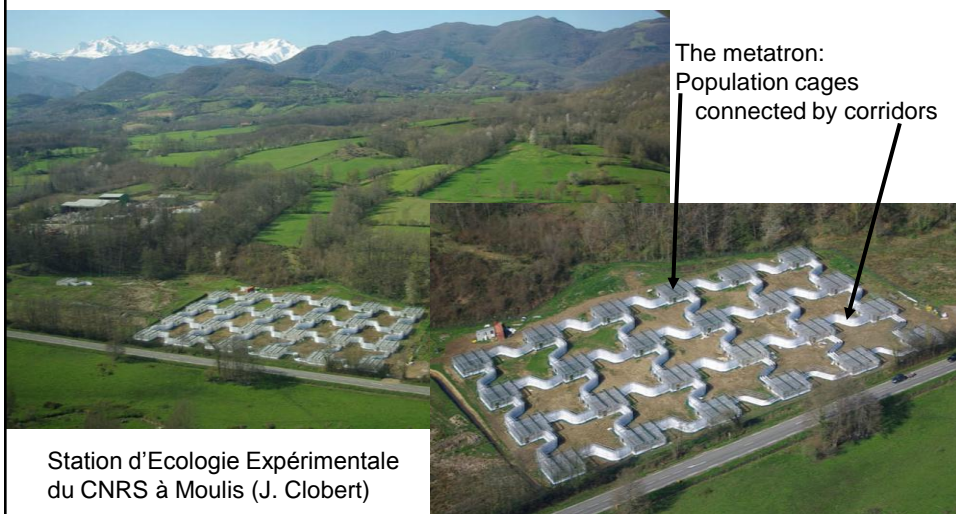
More detailed Global surveys are underway:

- 1) a, Luciano Javier Avila, Mariana Morando
- 2) Brazil: Carlos Frederico Duarte Rocha
- 3) Australia: David G. Chapple; Steve McAlpin, during my sabbatical
- 4) Namibia, South Africa: Christy Hipsely, D. Miles, Aaron Bauer, Bill Branch
- 5) Madagascar Bauer & Branch and during sabbatical year
- 6) Malaysia and Indonesia survey during my sabbatical next year

Species totally extinct by 2080

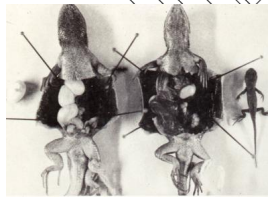


A proposed experimental set up to study the interplay between temperature, humidity and extinctions due to demographic collapse



Are speciation rates higher in clades born from an RPS ancestor?

Yes, the rate of speciation is about 2-3 times higher; if linkage to the origin of viviparity is further corroborated, does socially-mediated speciation lock lizards into inflexible heat tolerance if they are viviparous?



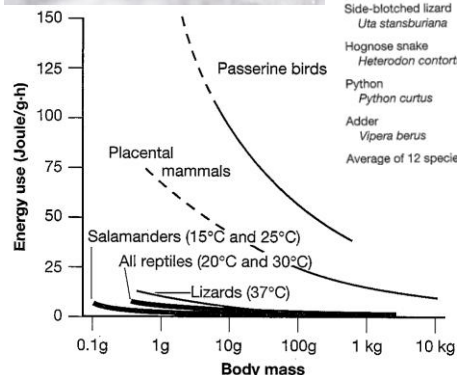
Sister pairs with 0 clade larger 0
Sister pairs with 1 clade larger 7
Sister pairs with 0-1 clades equal 2

Sister Diversification Binomial $P = 0.0078125$

Ecological Consequences of Lizard Declines

Lizards are “energy capacitors” – they can attain very high densities relative to endotherms due to:

- (1) much lower metabolic rates (ectotherms);
- (2) ability to exploit very tiny prey (arthropods) that for energetic reasons birds & mammals cannot eat



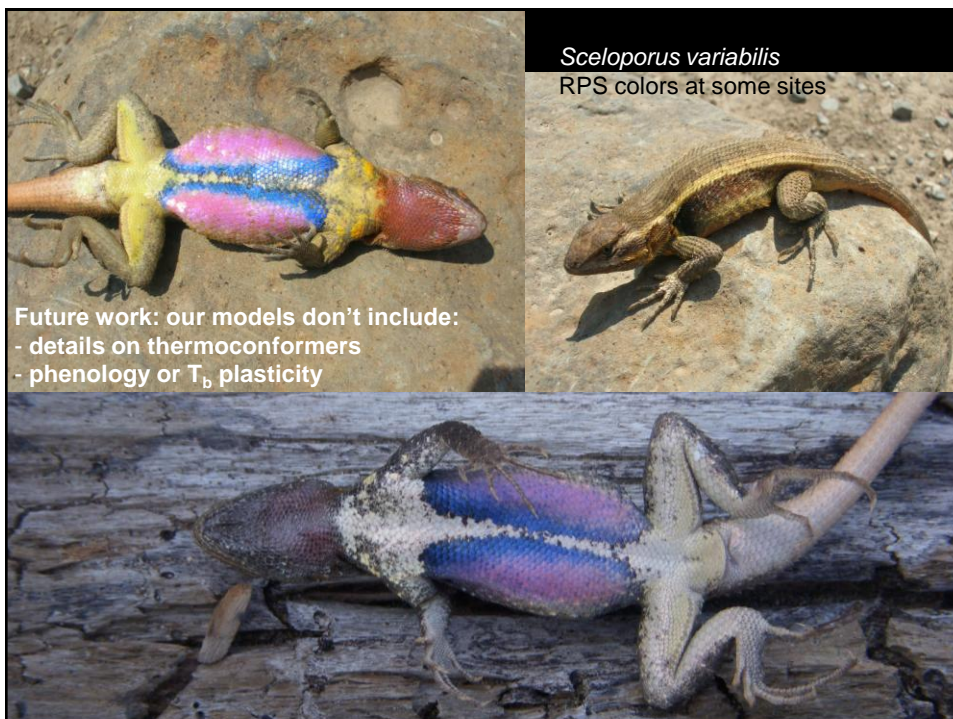
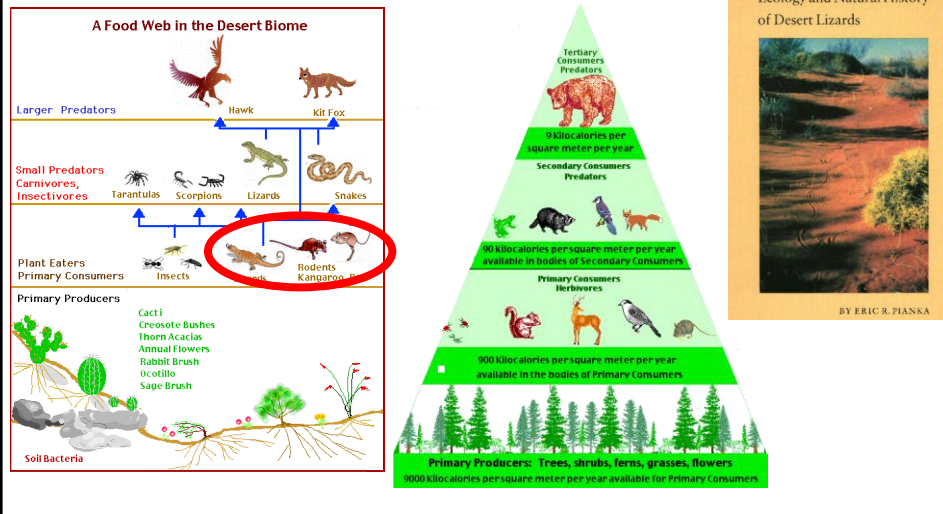
Ectotherms		Endotherms	
Species	Efficiency	Species	Efficiency
Red-backed salamander <i>Plethodon cinereus</i>	48	Kangaroo rat <i>Dipodomys merriami</i>	0.8
Mountain salamander <i>Desmognathus ochrophaeus</i>	76-98	Field mouse <i>Peromyscus polionotus</i>	1.8
Panamanian anole <i>Anolis limifrons</i>	23-28	Meadow vole <i>Microtus pennsylvanicus</i>	3.0
Side-blotched lizard <i>Uta stansburiana</i>	18-25	Red squirrel <i>Tamiasciurus hudsonicus</i>	1.3
Hognose snake <i>Heterodon contortrix</i>	81	Least weasel <i>Mustela erminea</i>	2.3
Python <i>Python curtus</i>	6-33	Savanna sparrow <i>Passerculus sandwichensis</i>	1.1
Adder <i>Vipera berus</i>	49	Marsh wren <i>Telmatochlamys palustris</i>	0.5
Average of 12 species	50	Average of 19 species	1.4



Ecological Consequences of Lizard Declines

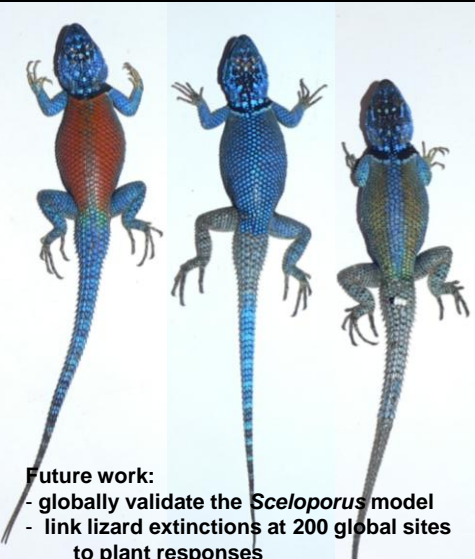
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- (1) much lower metabolic rates (ectotherms);
- (2) ability to exploit very tiny prey (arthropods) that for energetic reasons birds & mammals cannot eat
- (3) more efficient at conversion of plant or insect biomass into new lizard biomass (up to 50% vs < 2% in a bird or mammal of equal body mass; vs 2-3%)




Sceloporus minor
(Barry Stephenson et al.)

oo bb by

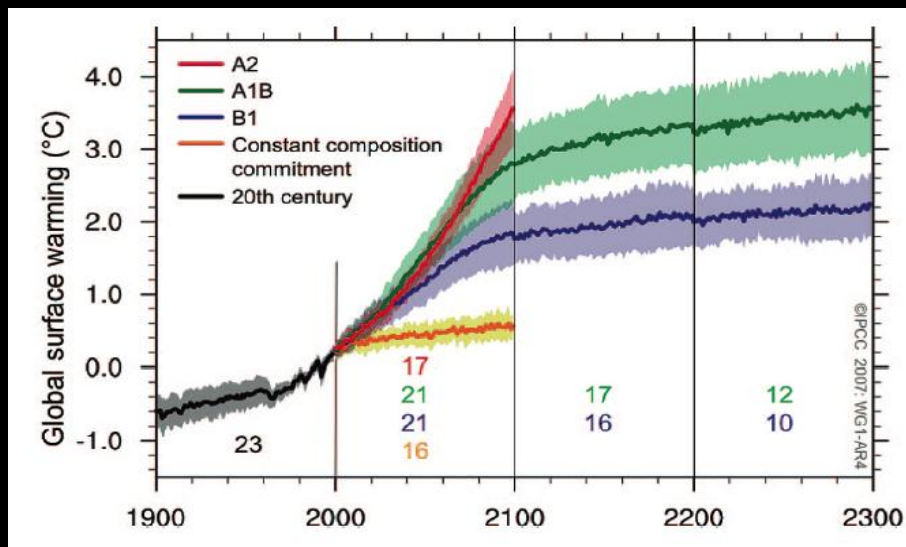



Future work:
 - globally validate the *Sceloporus* model
 - link lizard extinctions at 200 global sites to plant responses

Liolaemus sp.; Santa Cruz Province, Argentina (2007)


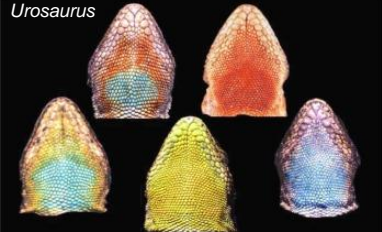


If we bend the CO₂ curve (B1) we limit extinctions to 5% (2050 levels), not the 20% of species extinctions predicted for 2080 (A2).





Sceloporus ornatus

Urosaurus

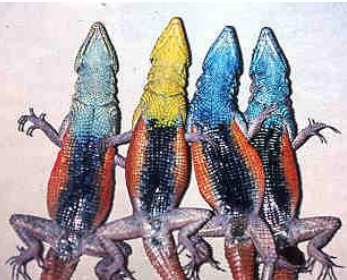
ACKNOWLEDGMENTS

NSF: "Assembling the Tree of Life - The Deep Scaly Project: Resolving Higher Level Squamate Phylogeny Using Genomic and Morphological Approaches" (EF 0334966);

"Partnership for International Research and Education: Establishing Sustainable International Collaborations in Evolution, Ecology, and Conservation Biology" (OISE 0530267);

National Evolutionary Synthesis Center: "Perspectives on the Origin and Conservation of Biodiversity in Patagonia" – Catalysis Group meeting at Duke University (EF 0423641).

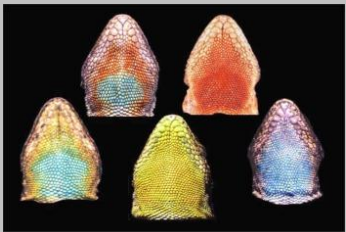
Brigham Young University: Dept. of Biology, Bean Life Science Museum, Office of Research & Creative Activities (student mentoring awards)







Platysaurus broadleyi,
Africa

Urosaurus ornatus mating system - males

- Throat color morphs
 - Orange
 - Blue
 - Yellow
- Social Status
 - Territorial
 - Satellite
 - Floater/Sneaker
- Mating System
 - Polygynous
 - Monogamous





Change in Operative Temperature between wet and drought years

