

# MACROEVOLUTION

Lecture 13

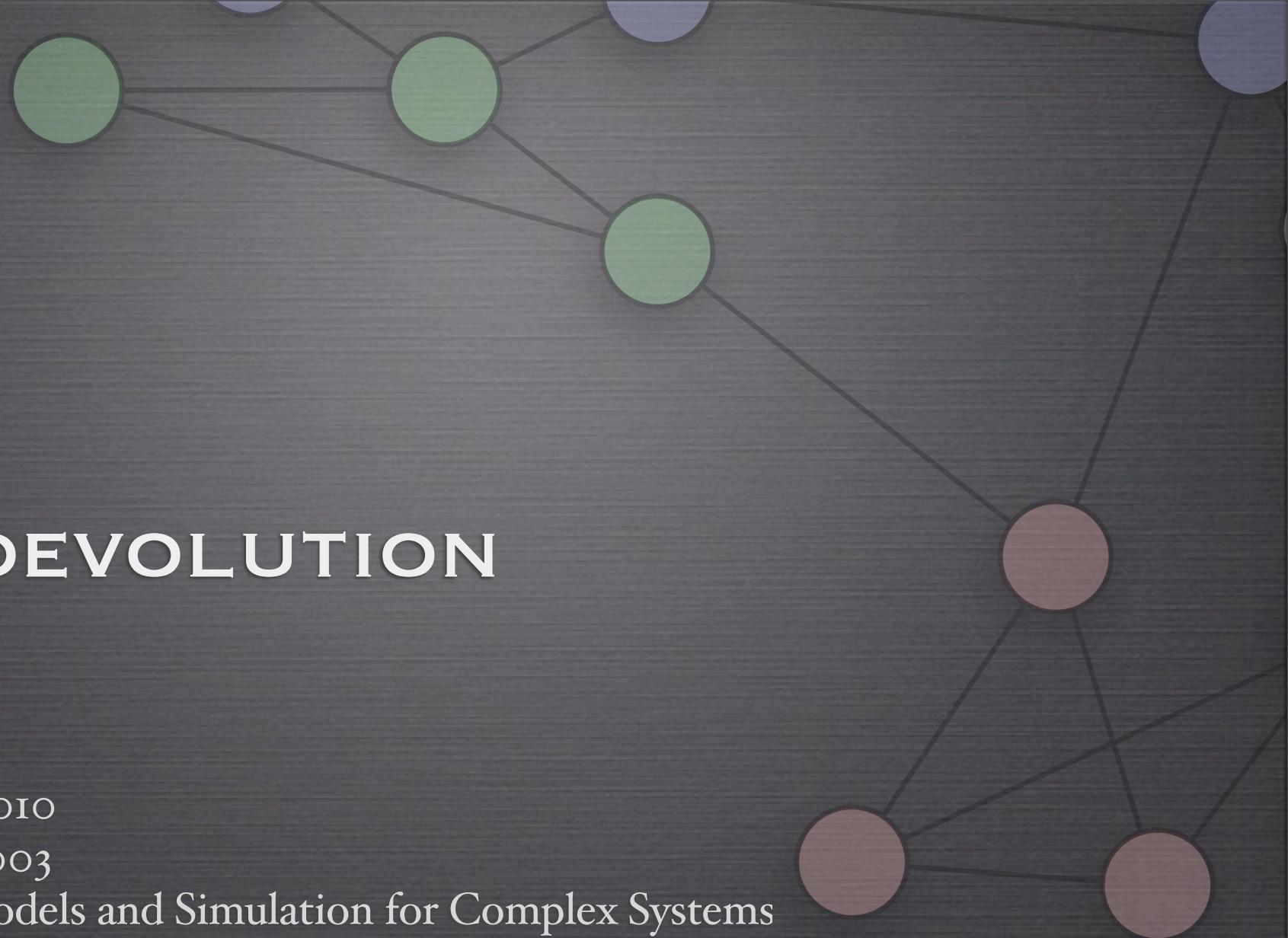
19 October 2010

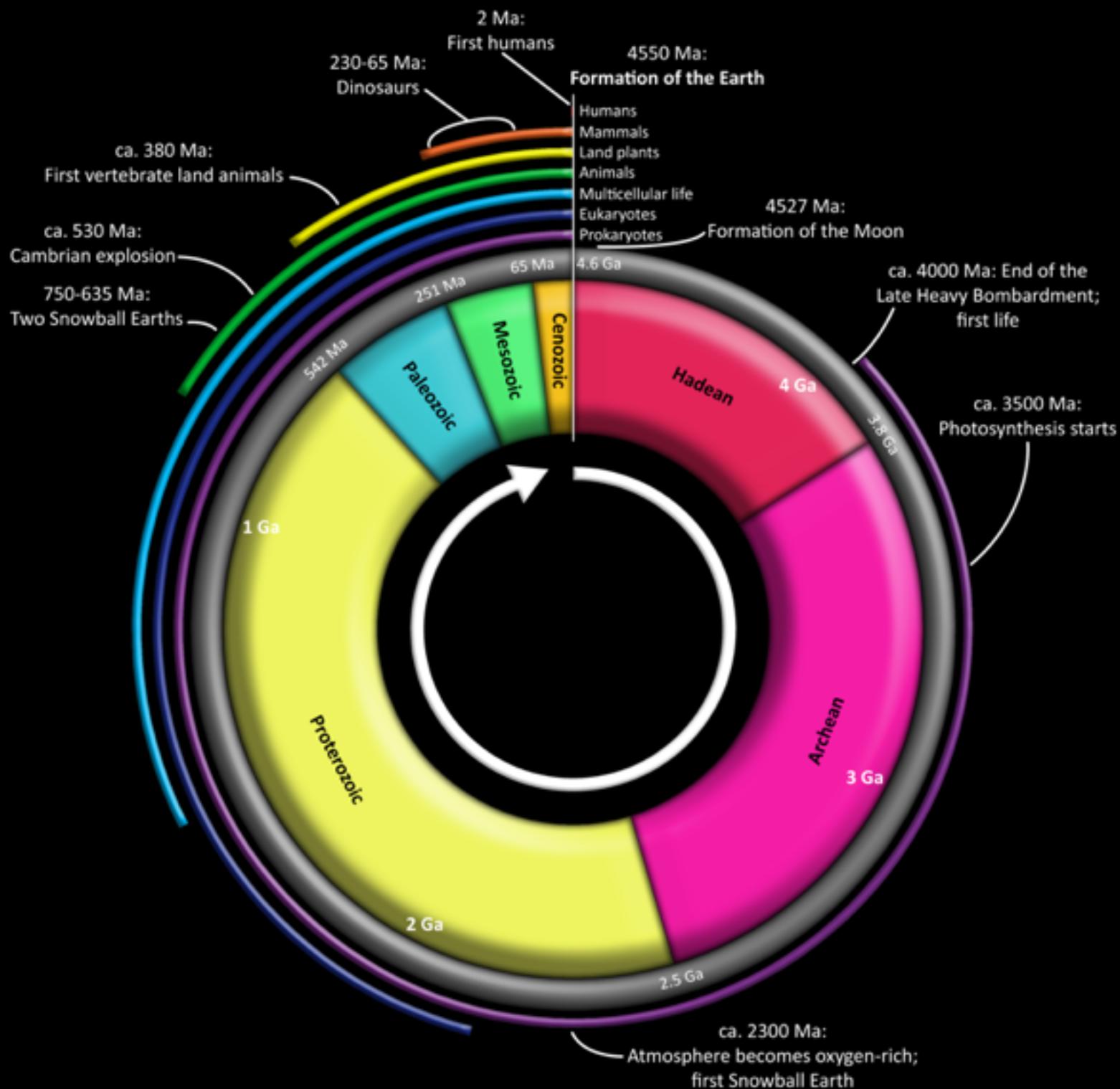
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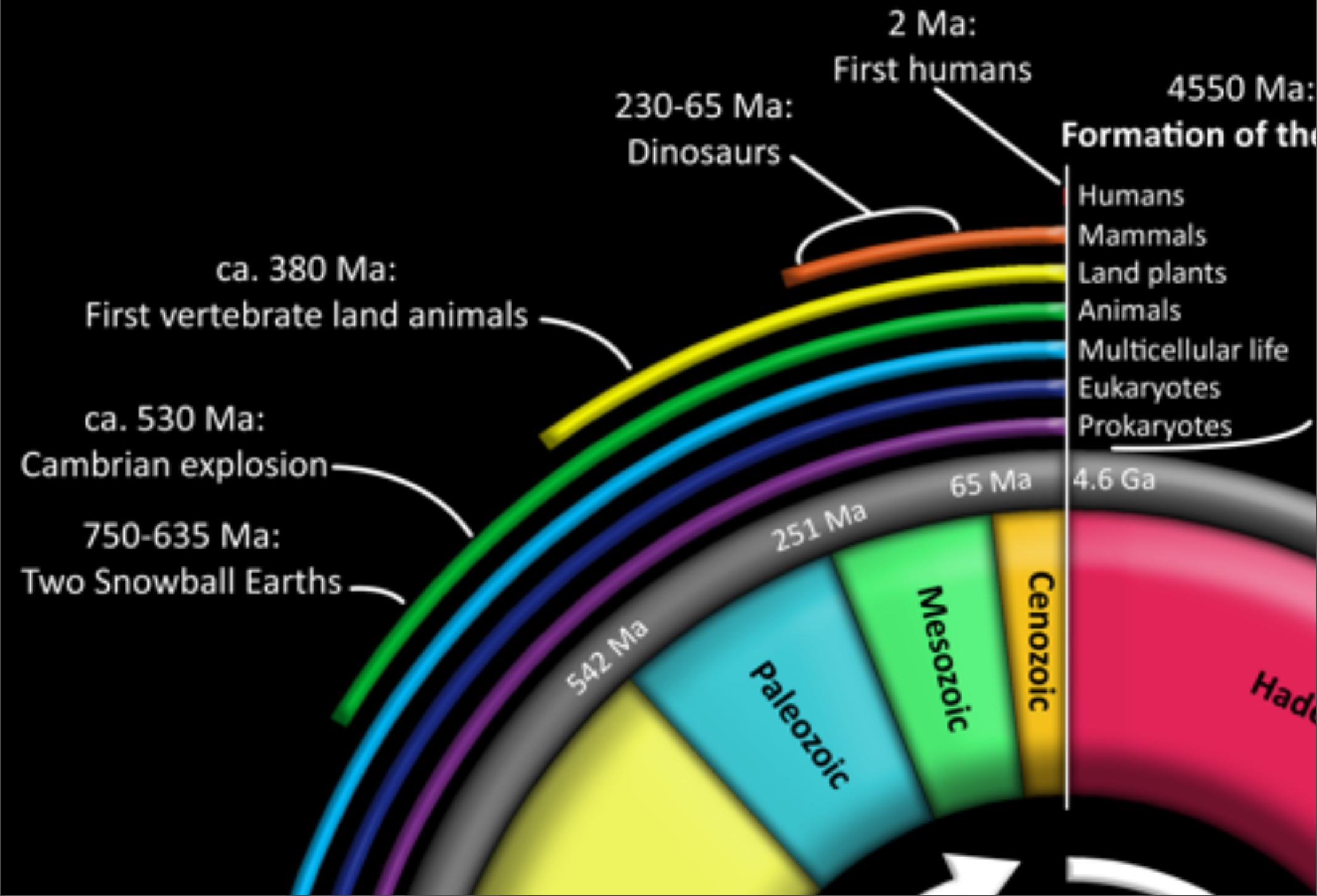
Inference, Models and Simulation for Complex Systems

Prof. Aaron Clauset

University of Colorado

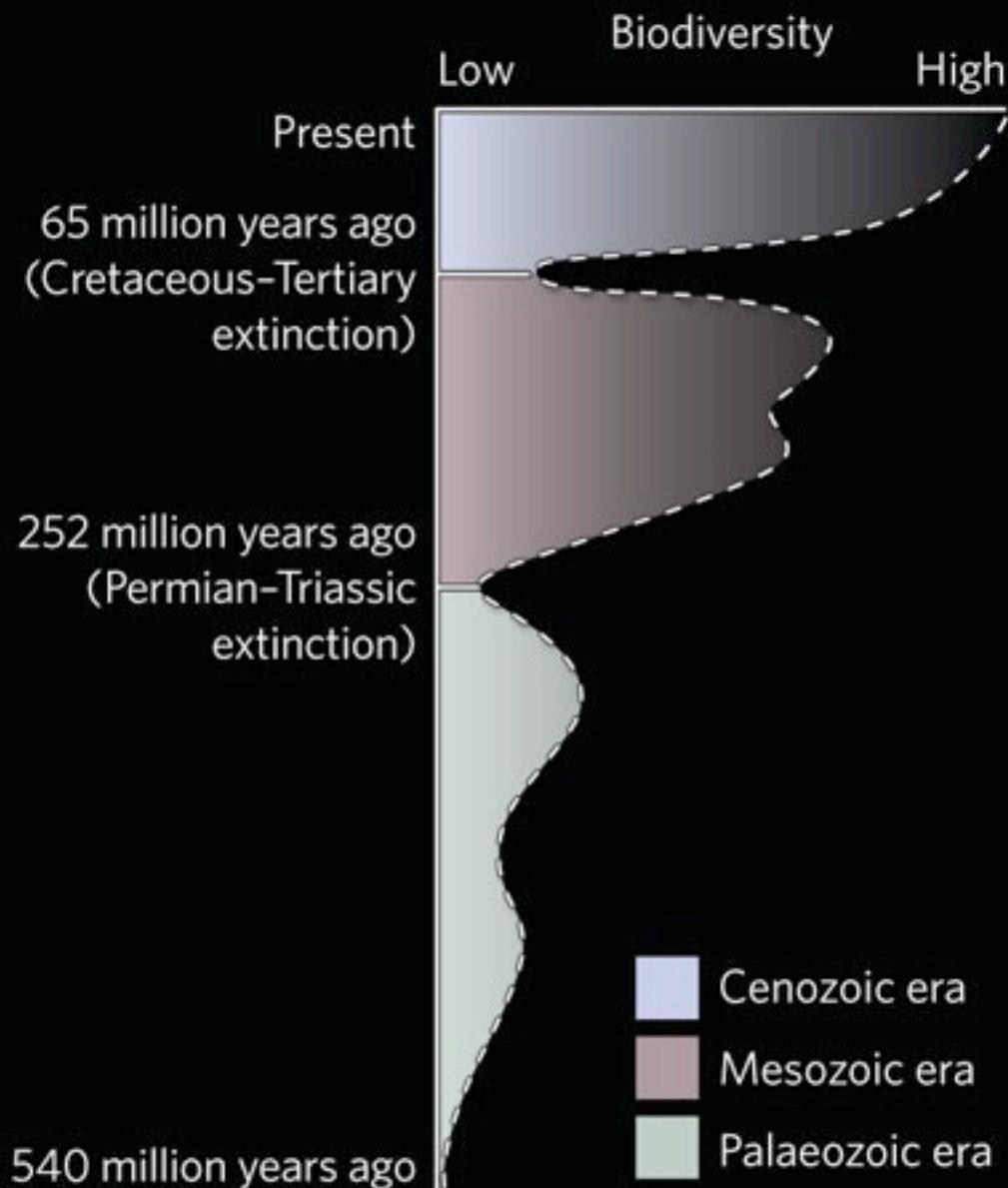






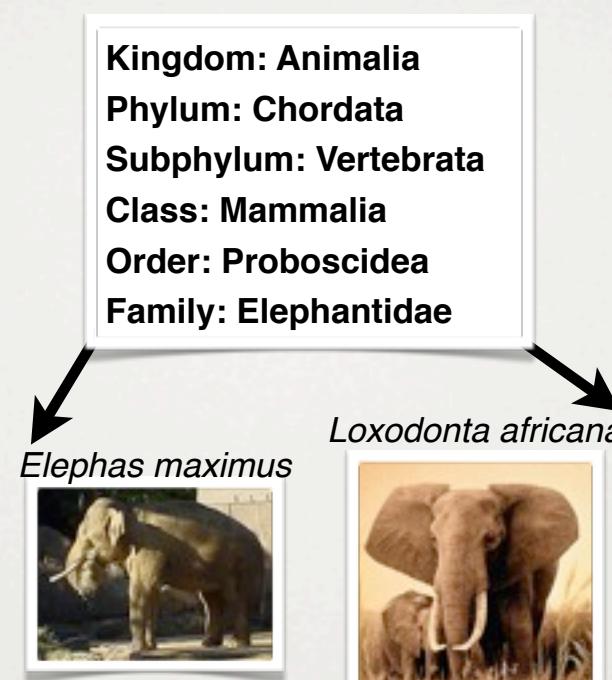
# AN 1860 RECONSTRUCTION OF THE FOSSIL RECORD

The first of many, hotly-debated, estimates  
of Phanerozoic biodiversity.



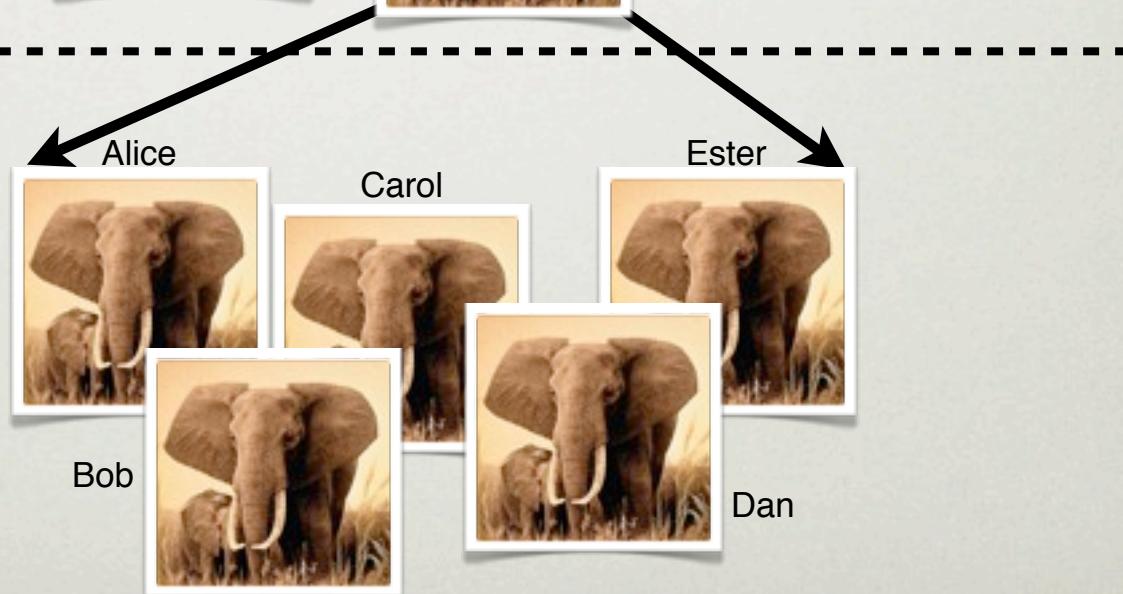
## Macroevolution

*species level and above*



## Microevolution

*below species level*



# MACROEVOLUTIONARY QUESTIONS

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- body size (or mass)
- metabolism
- diversity [number of species, morphology, taxonomy, behavior]
- complexity [prokaryote → eukaryote → multi-cellularity → sociality]
- carrying capacities
- mass extinctions & recoveries
- adaptive radiations

# SPECIES SIZE DIVERSITY

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2g



40g



75,000g



2g



288g



165,000g



5,500,000g



250,000g

# SPECIES SIZE DIVERSITY



2g



2g



40g



288g



75,000g



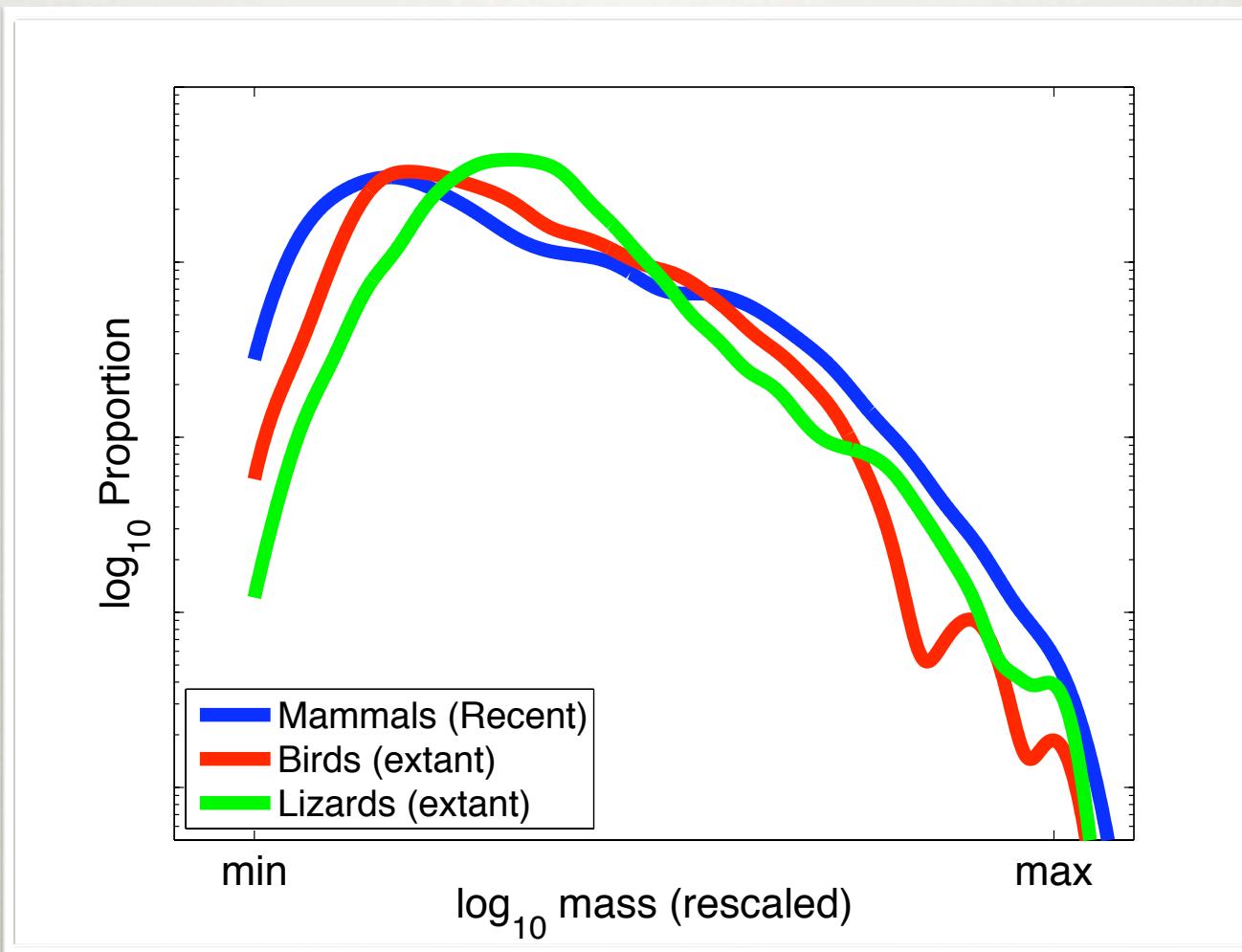
165,000g



250,000g



5,500,000g



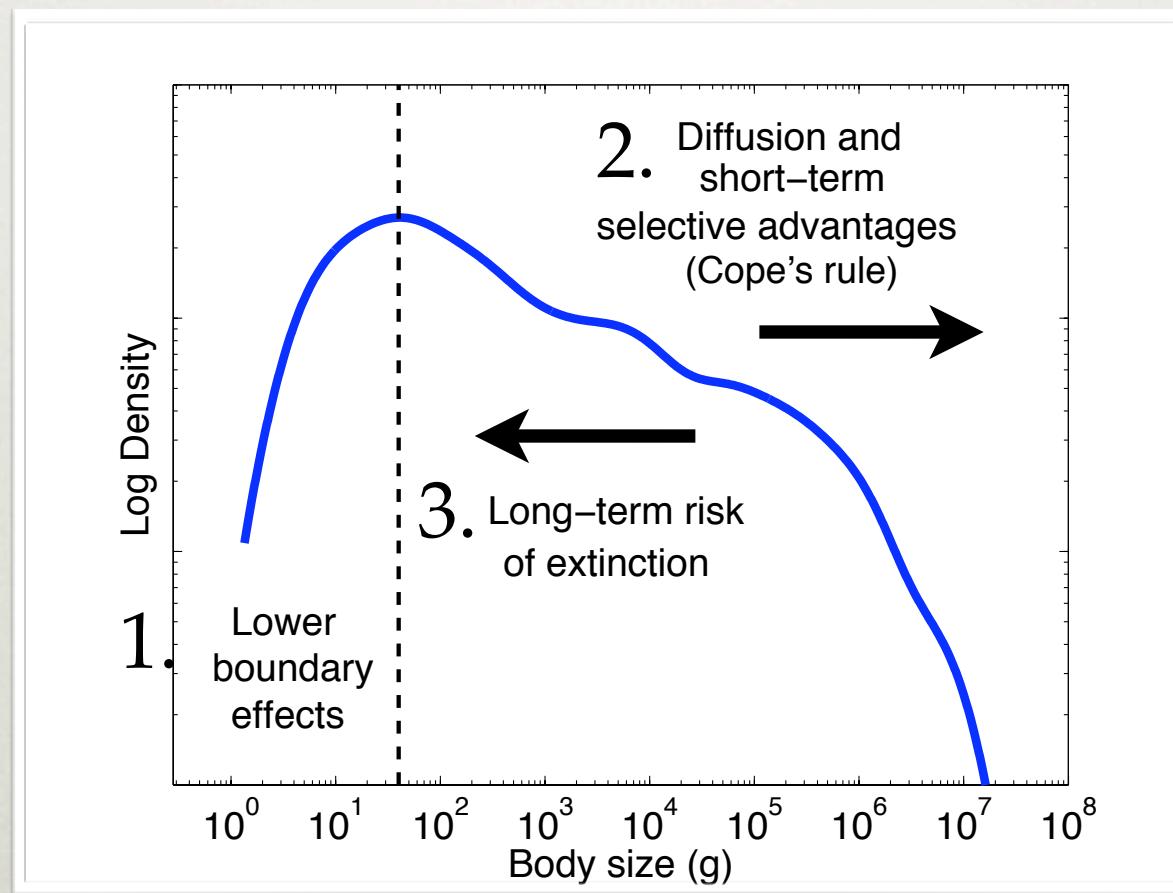
# PAST WORK

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- focus on right-skew and mode's location
  - mainly terrestrial mammals (best data)
1. energetic optima + competition
    - Lomolino (1985)
    - Sebens (1987)
    - Brown, Marquet, Taper (1993, 1996)
  2. minimum size + cladogenetic diffusion (+ Cope's rule)
    - Stanley (1973)
    - McShea (1994)

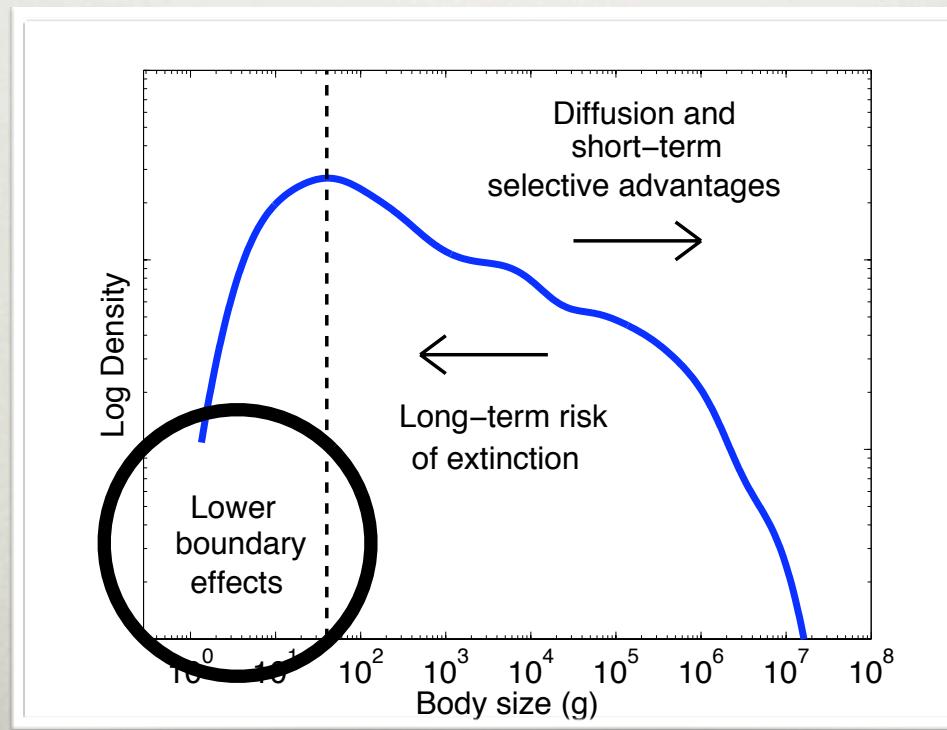
# MACROEVOLUTIONARY MODEL

three mechanisms



# MACROEVOLUTIONARY MODEL

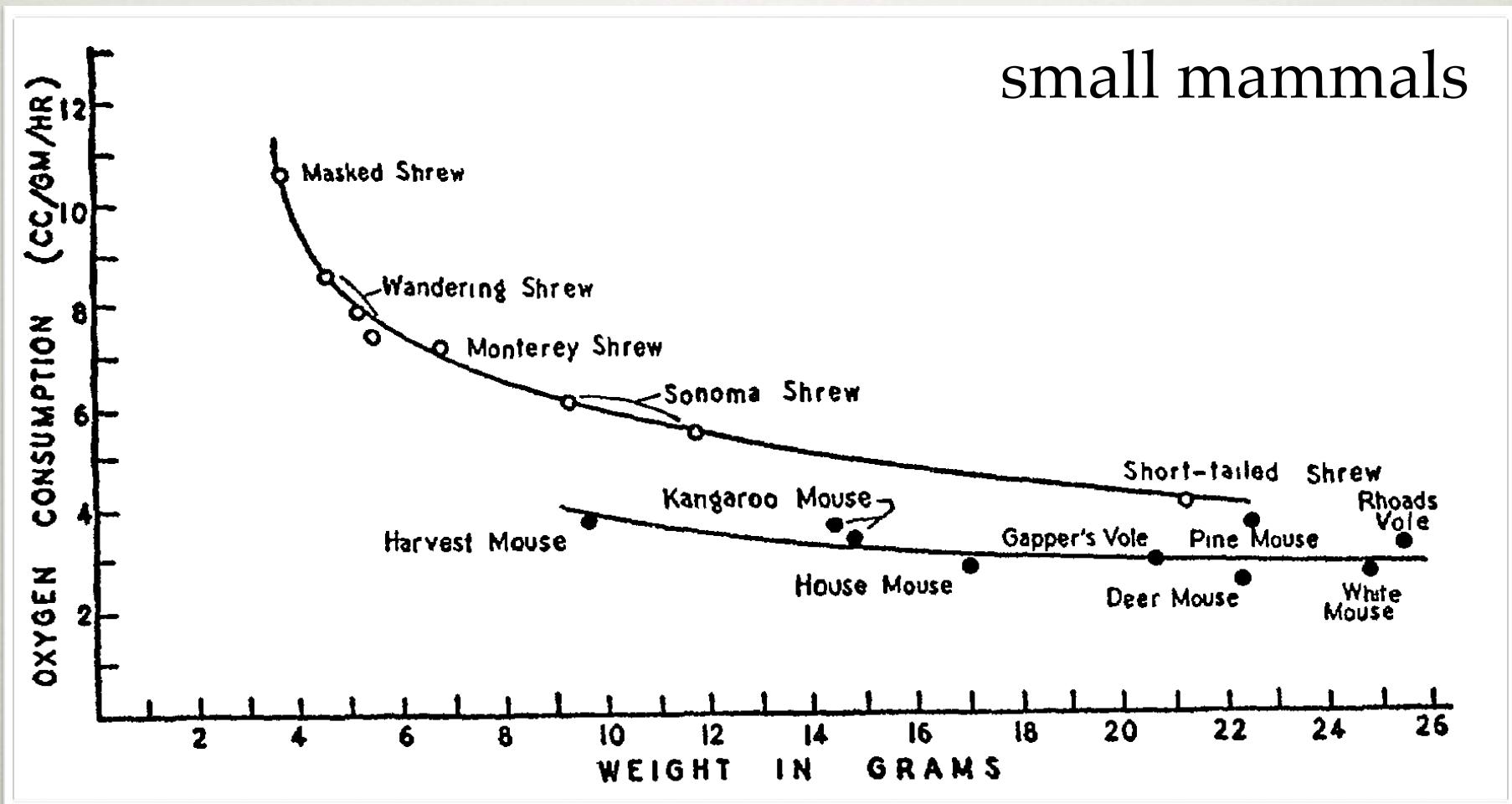
1. hard lower boundary = physiology, metabolism
2. biased diffusion = short-term advantages
3. size-dependent extinction rates = long-term risks



data from Smith *et al.* (2003)

# HARD LOWER LIMIT

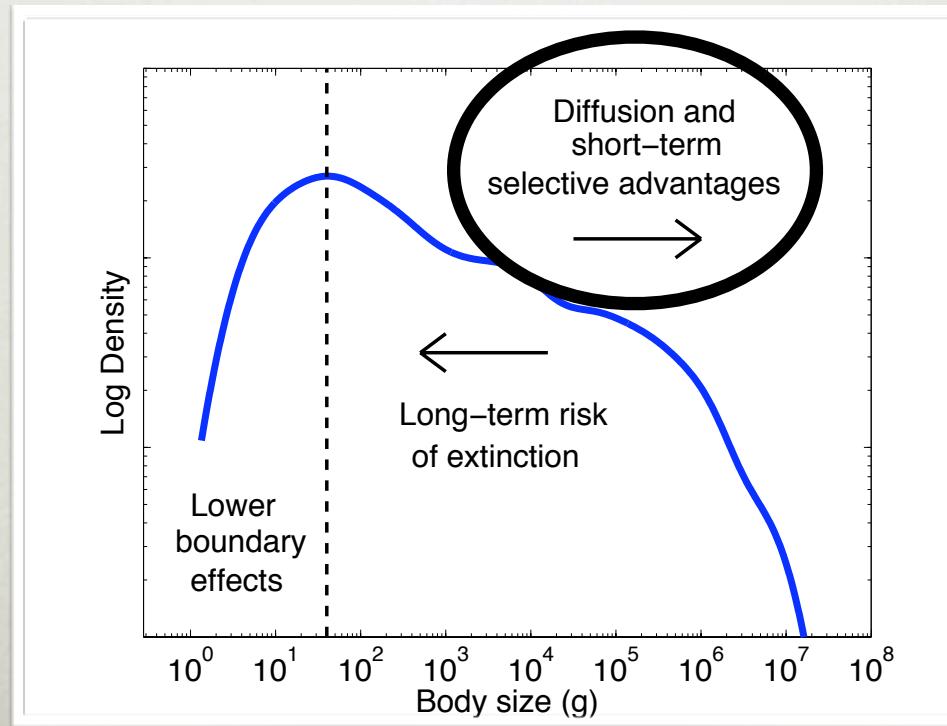
metabolic constraints  $x_{\min} \approx 2g$



# MACROEVOLUTIONARY MODEL

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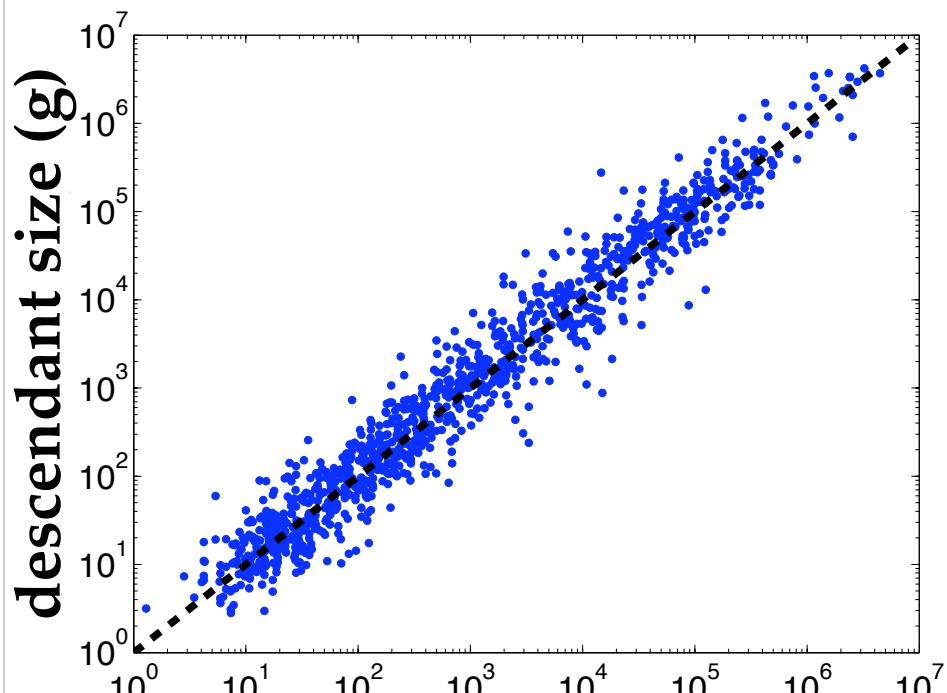
1. hard lower boundary = physiology, metabolism
2. biased diffusion = **short-term advantages**
3. size-dependent extinction rates = **long-term risks**



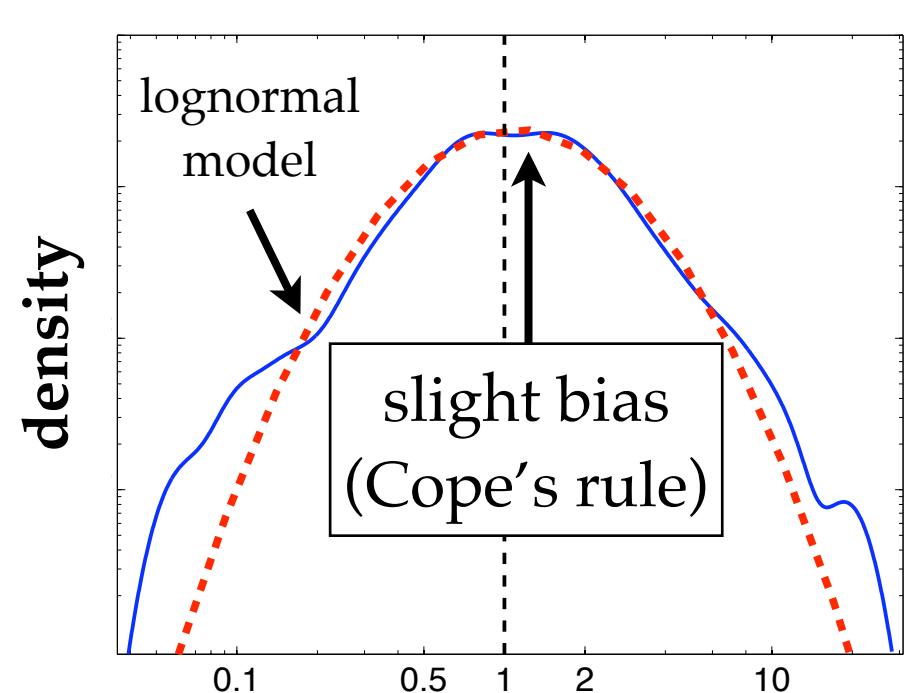
# BIASED DIFFUSION

ancestor-descendant data

[North Am. mammals]



ancestor size (g)

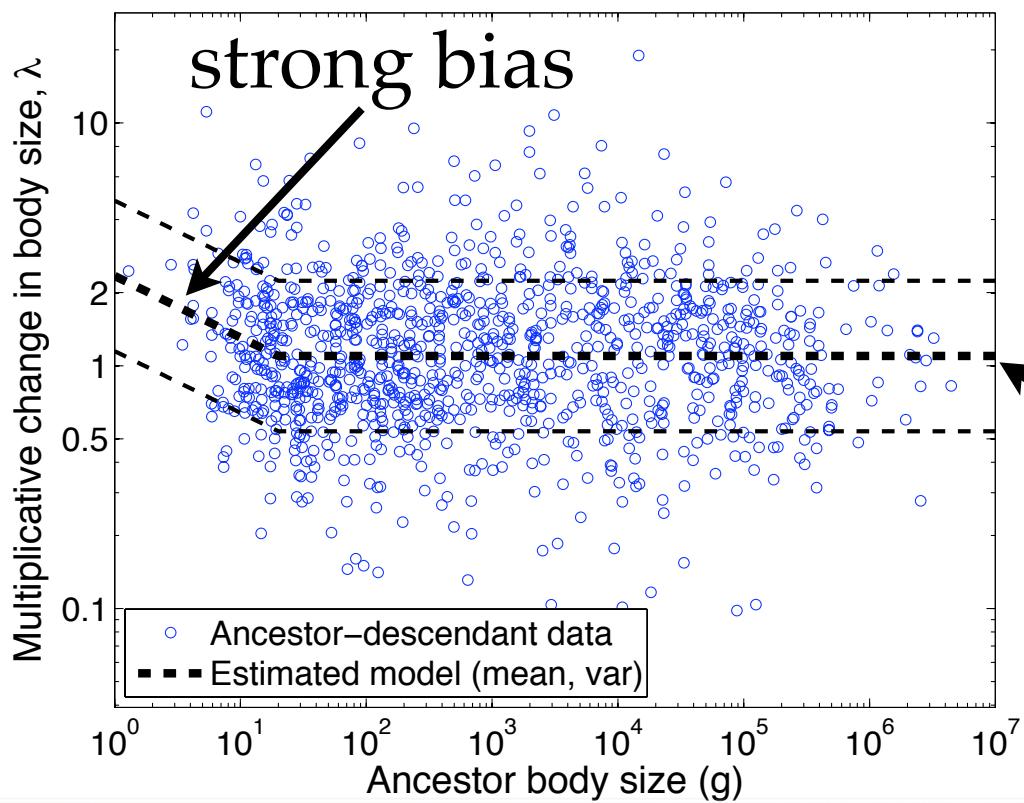


proportional change

# COPE'S RULE

ancestor-descendant data

[North Am. mammals]



$$F(\lambda) = \mathcal{N}(\mu(x_A), \sigma)$$

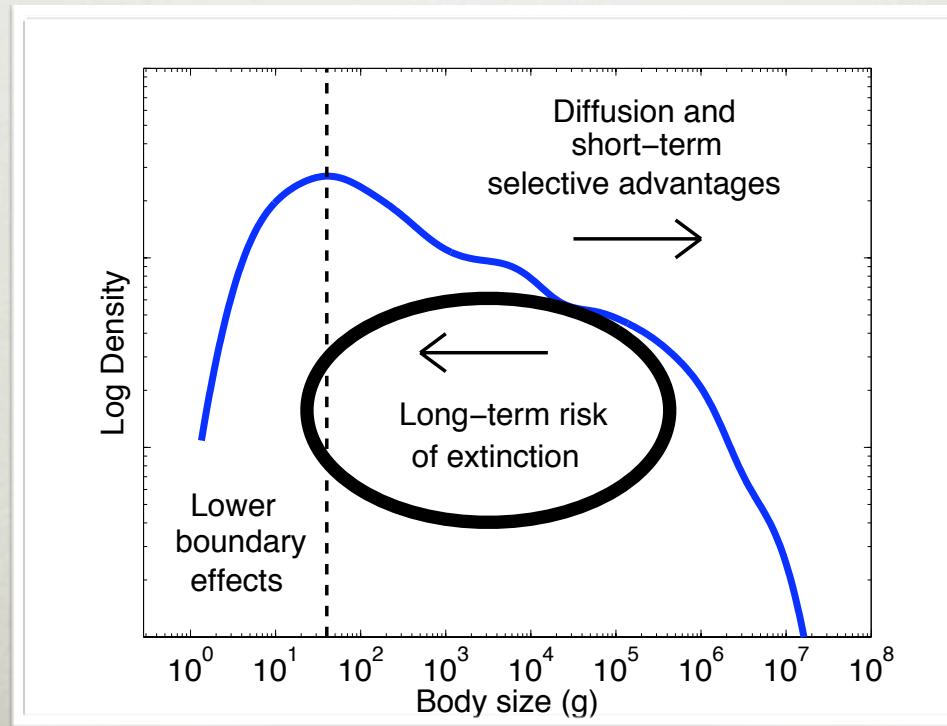
$\mu(x)$  is piece-wise  
log-linear

slight bias (4.1%)

# MACROEVOLUTIONARY MODEL

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1. hard lower boundary = physiology, metabolism
2. biased diffusion = short-term advantages
3. size-dependent extinction rates = **long-term risks**



# EXTINCTION / SPECIATION

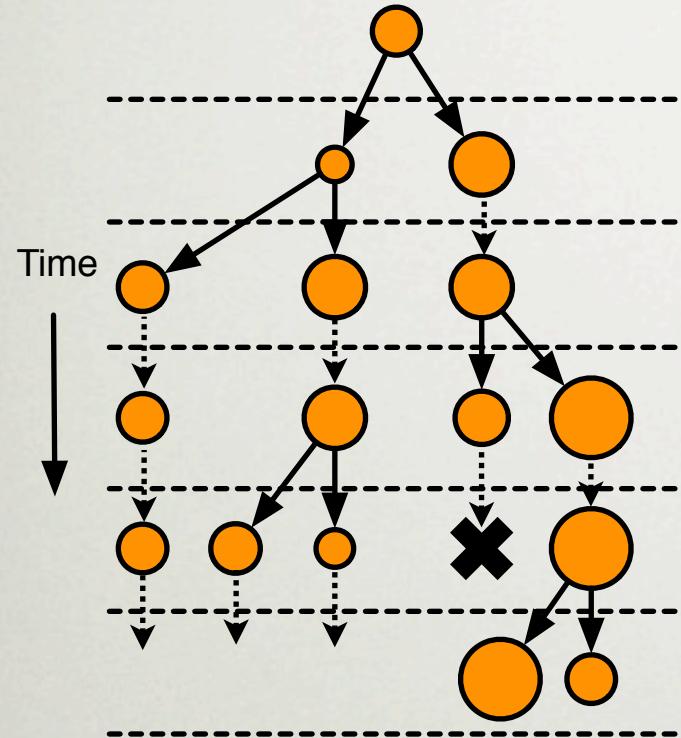
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1. extinction rates difficult to estimate from data
2. simple parametric model
  - rate depends on size  $dp_{\text{extinct}}/dx > 0$   
[Van Valen (1973), Van Valkenburgh et al. (2004), Liow et al. (2008),  
Alroy (unpublished)]
  - speciation rate independent of size\*
  - fitted to data

\*only ratio of extinction-to-speciation rate matters

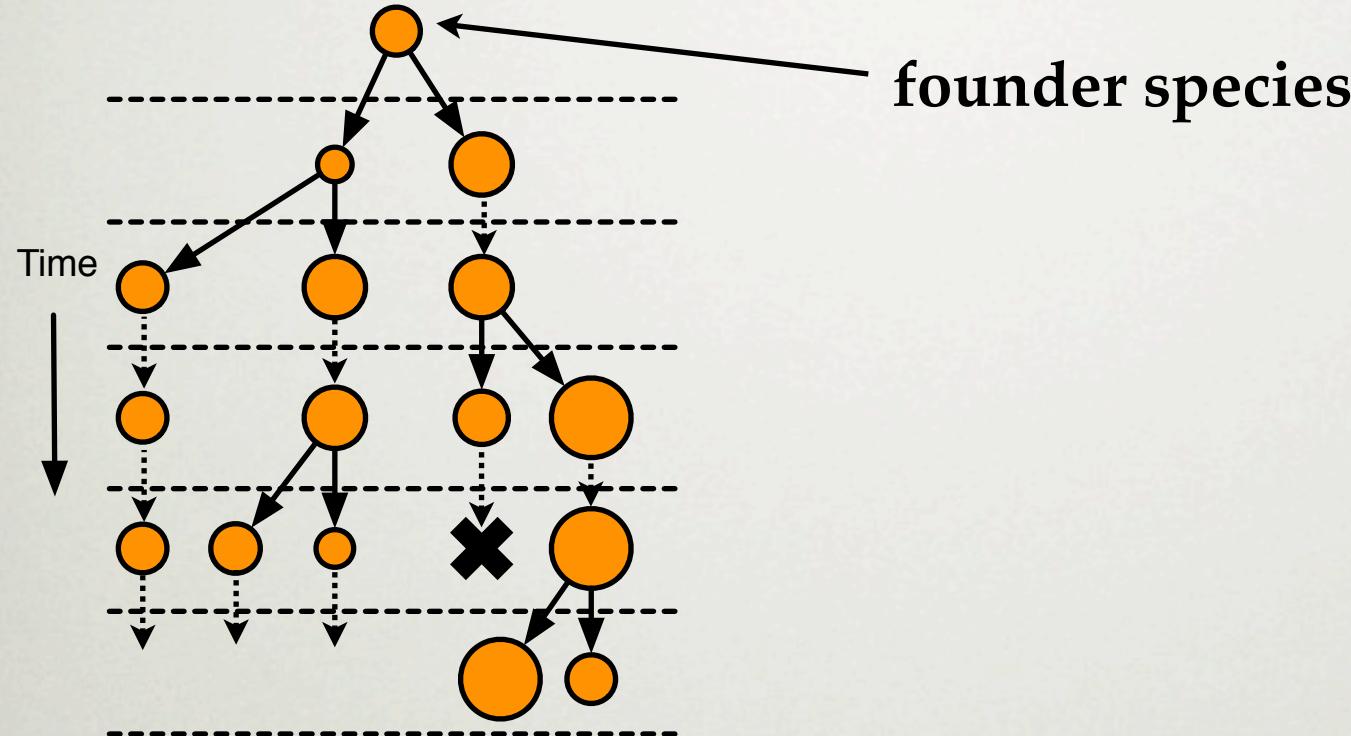
# CLADOGENESIS

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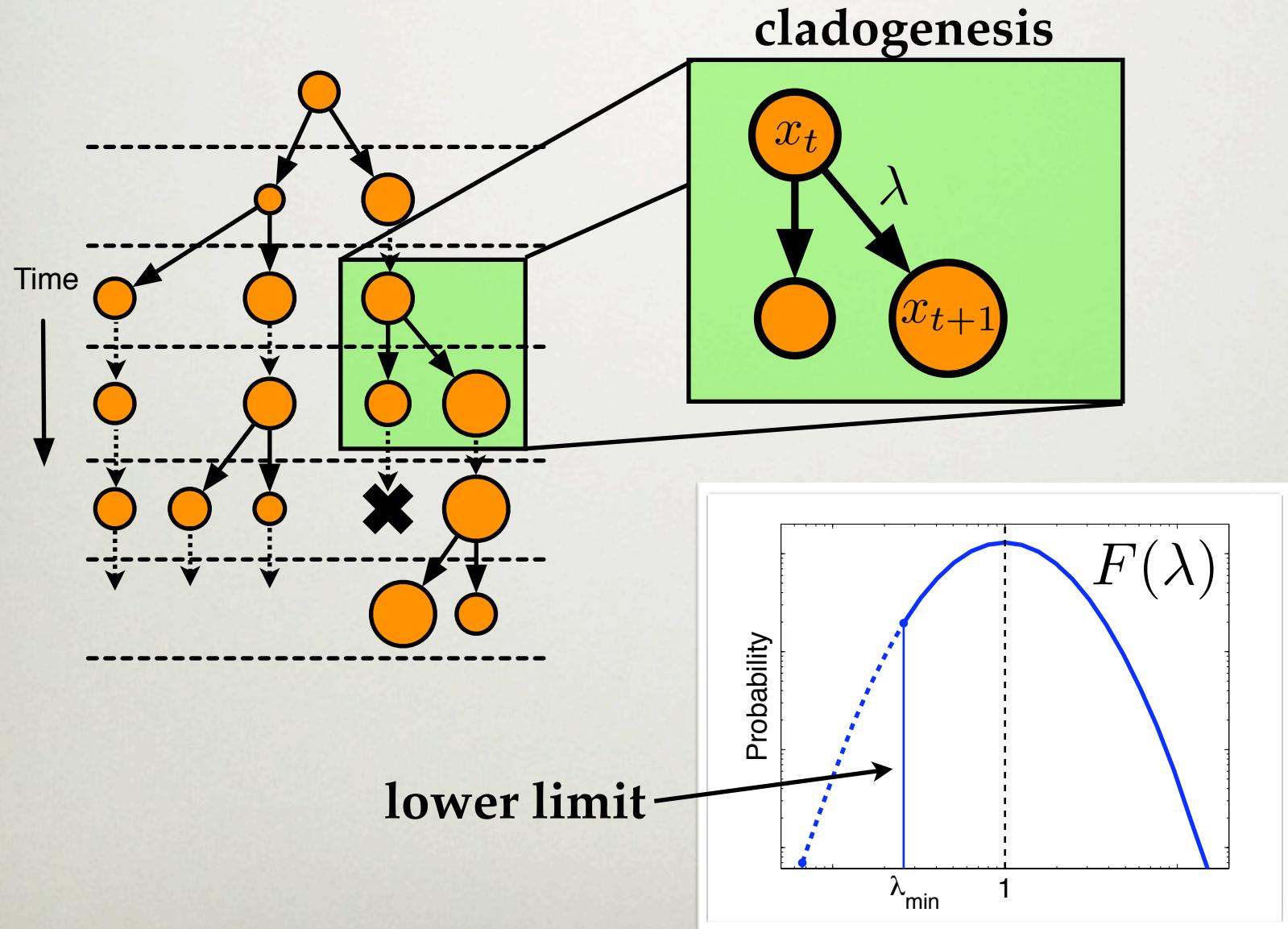


# CLADOGENESIS

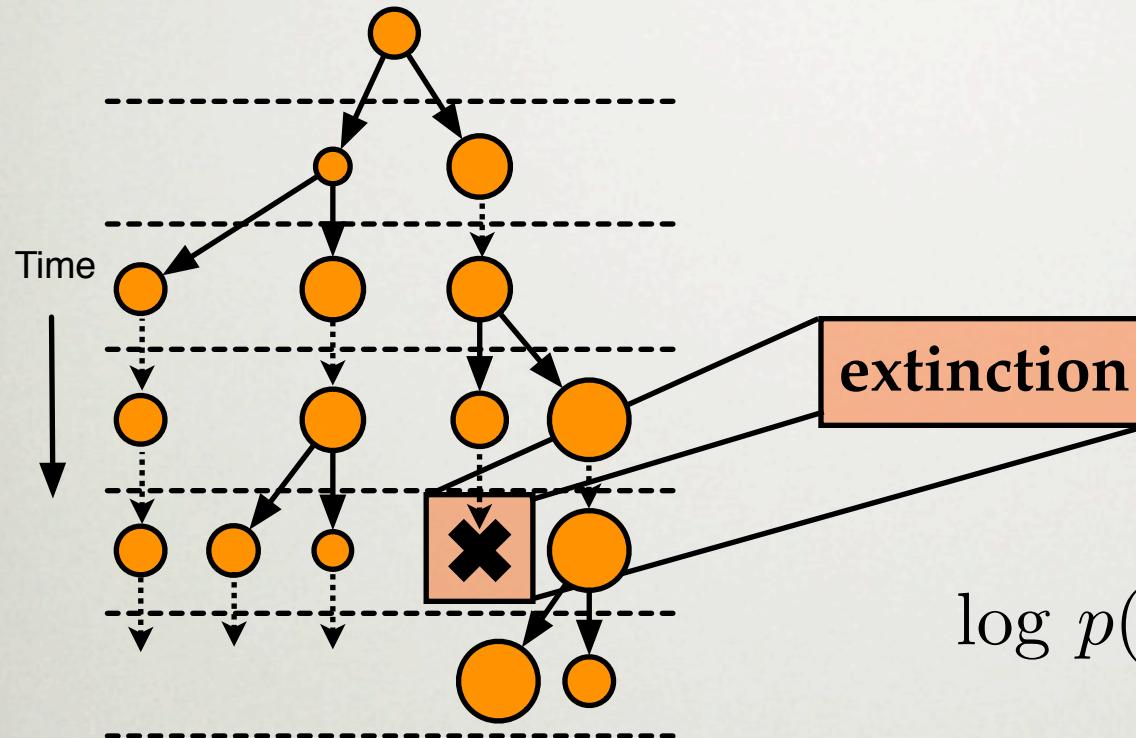
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# CLADOGENESIS



# CLADOGENESIS



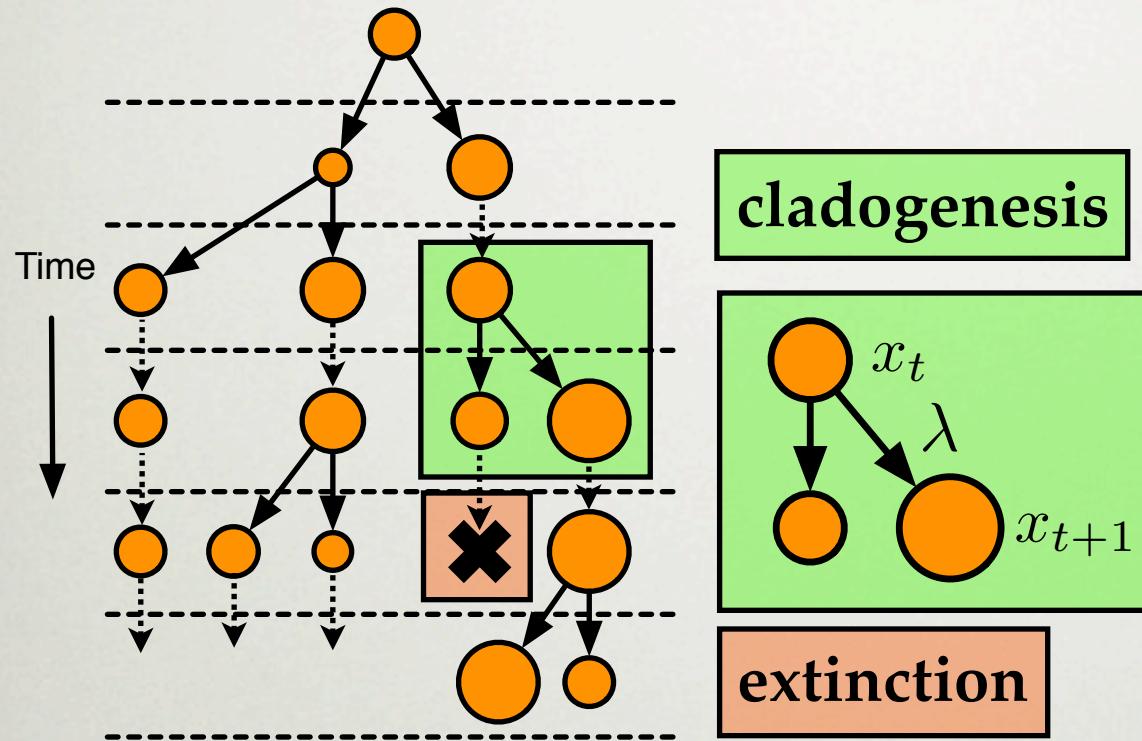
independent events  
prob. depends on size

$$\log p(x) = \rho \log x + \log \beta$$

size dependence

basal rate

# CLADOGENESIS



## model features

- size-dependent fluctuations
- Cope's rule
- size-dependent extinction rate
- lower limit  $x_{\min}$

# COMMENTS

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## Ecology-free model

- no inter-specific interaction  
(competition, predation, etc.)
- no population dynamics
- no environment
- no mass extinctions
- no anagenetic variation in body size

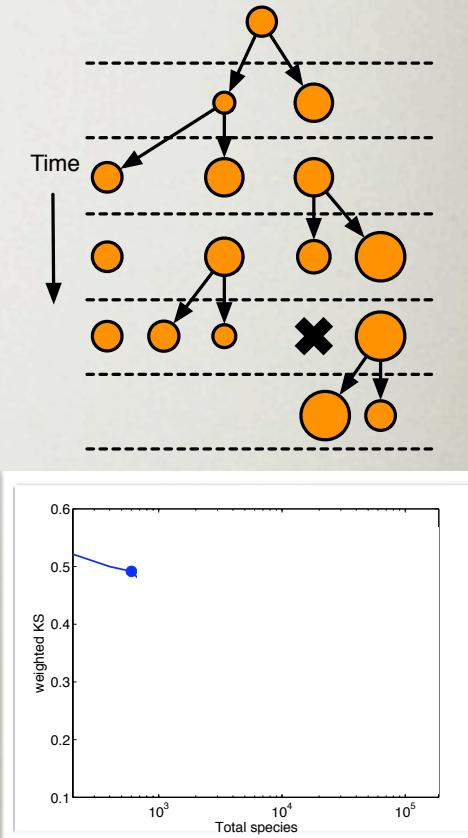
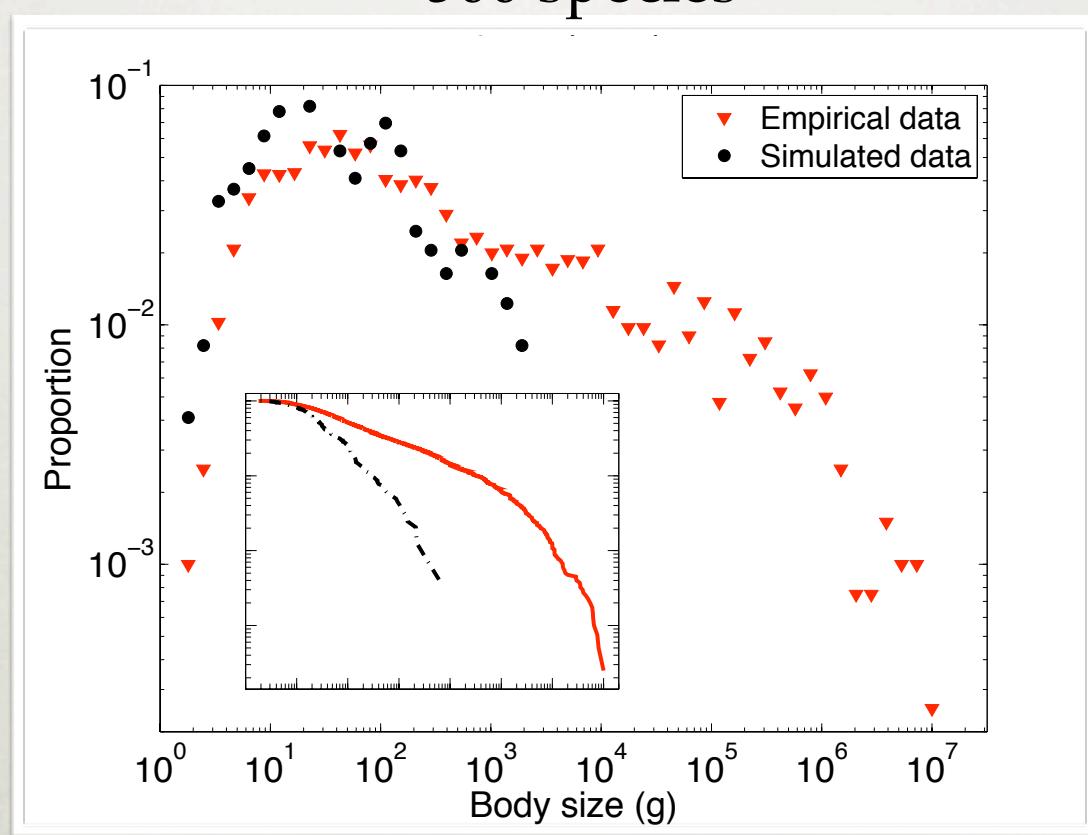
# PARAMETERS

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	founder body size	$x_0$	40g
initialization	mean species lifetime	$\nu$	1.60(1) My
length of sim.	years in equilibrium	$\tau$	60 My
boundary	lower bound	$x_{\min}$	1.8g
diffusion	log $\lambda$ -intercept	$c_1$	0.33
	log $x$ -intercept	$c_2$	1.30
	systematic bias	$\delta$	0.04
	variance	$\sigma$	0.63
extinction	species at equilibrium	$n$	5000
	baseline extinction rate	$\beta$	$1/n$
	rate of extinction increase	$\rho$	0.025

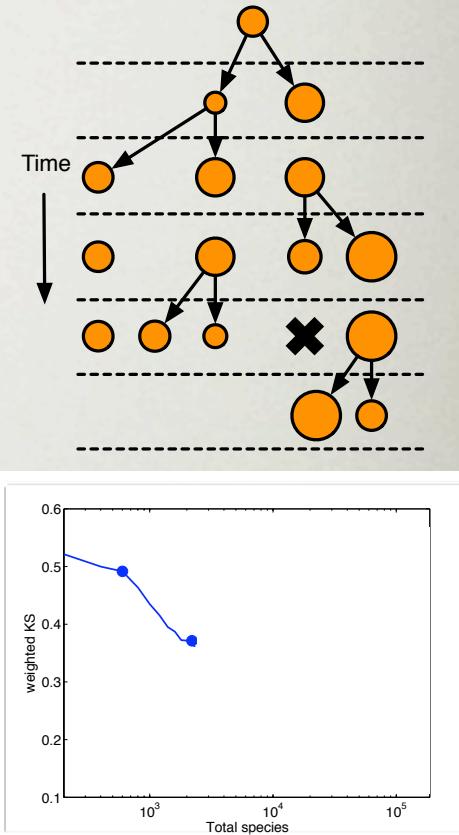
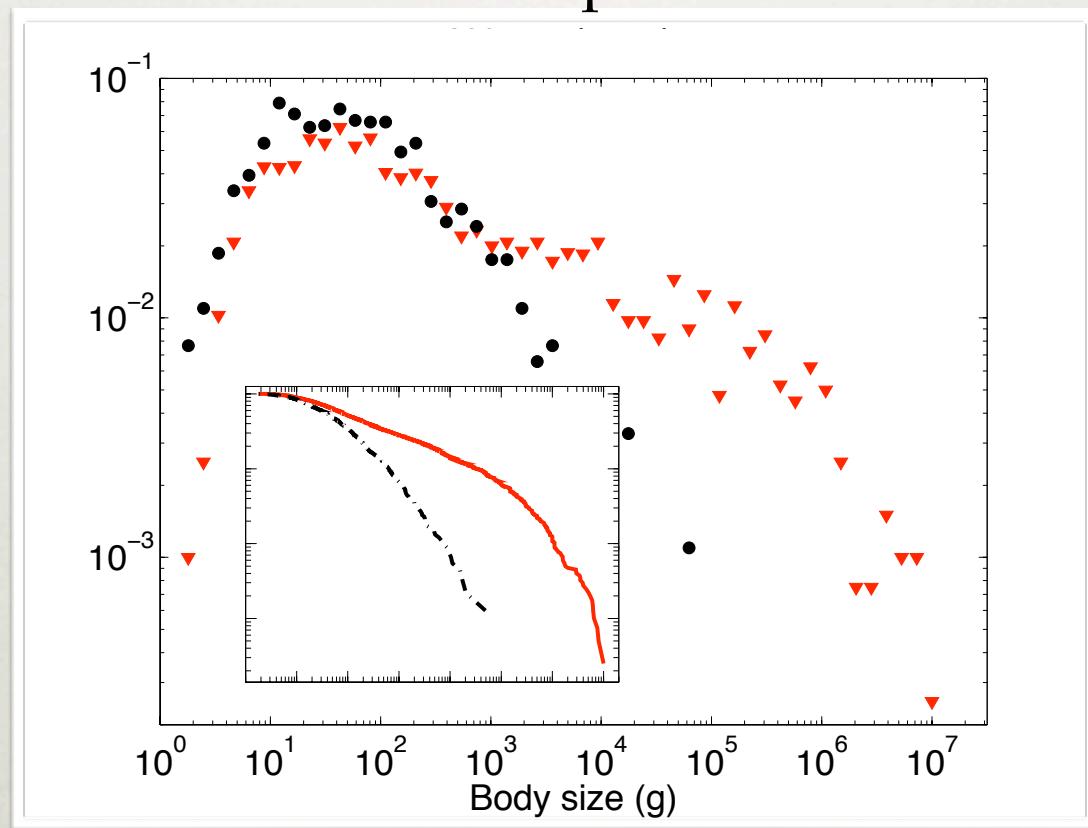
# SIMULATION

- ▼ empirical
- simulated



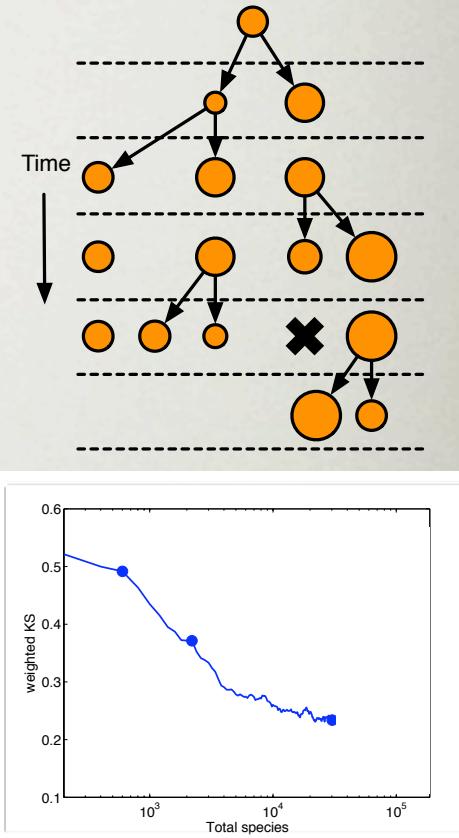
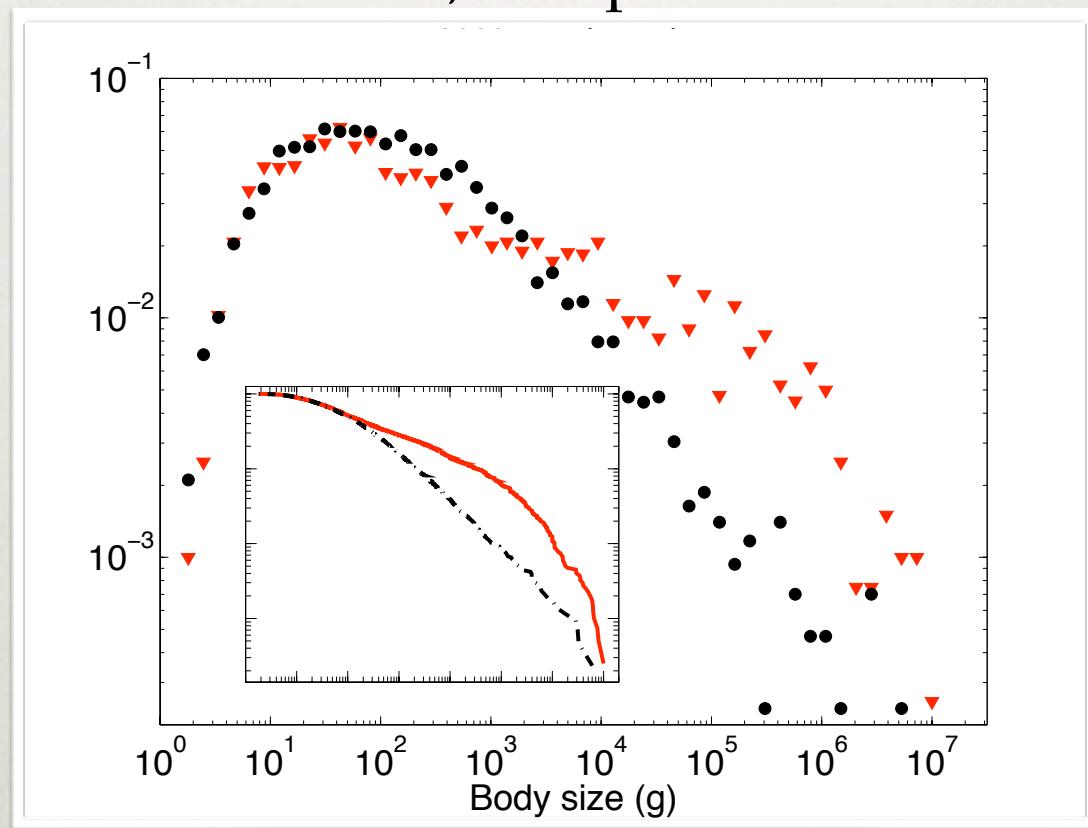
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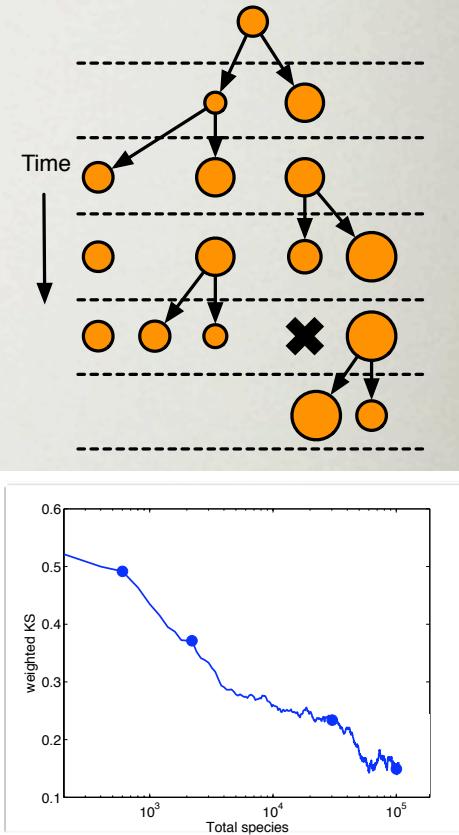
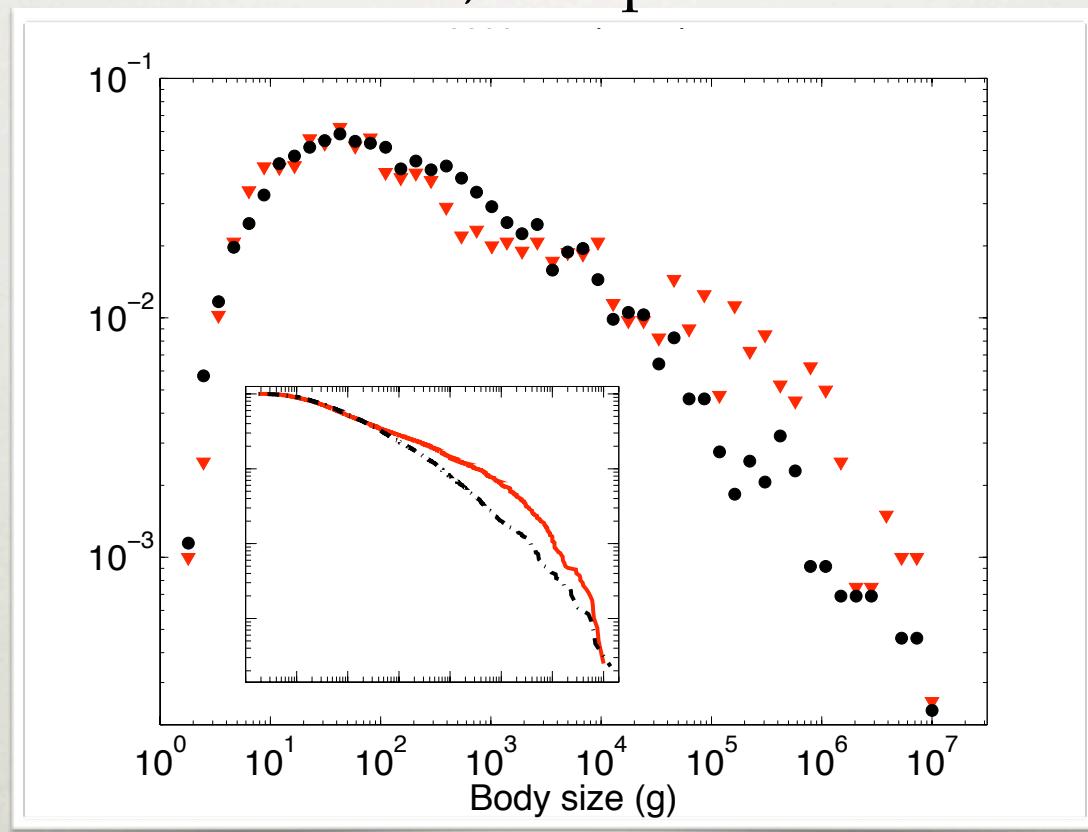
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# SIMULATION

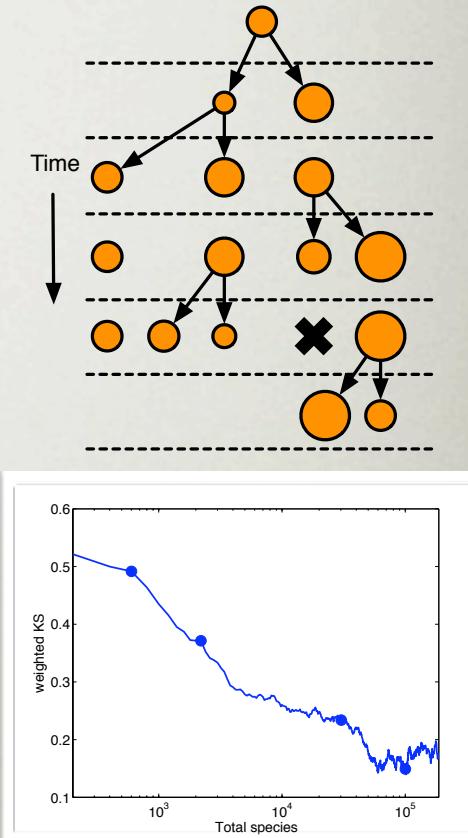
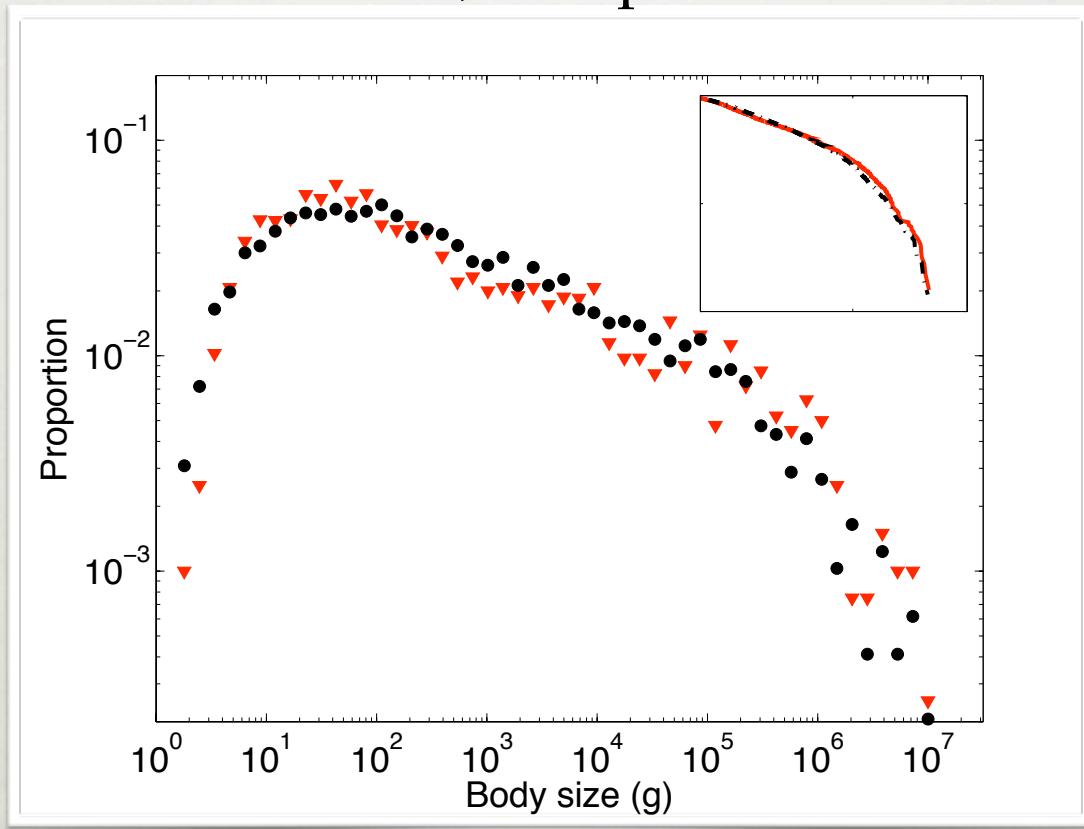
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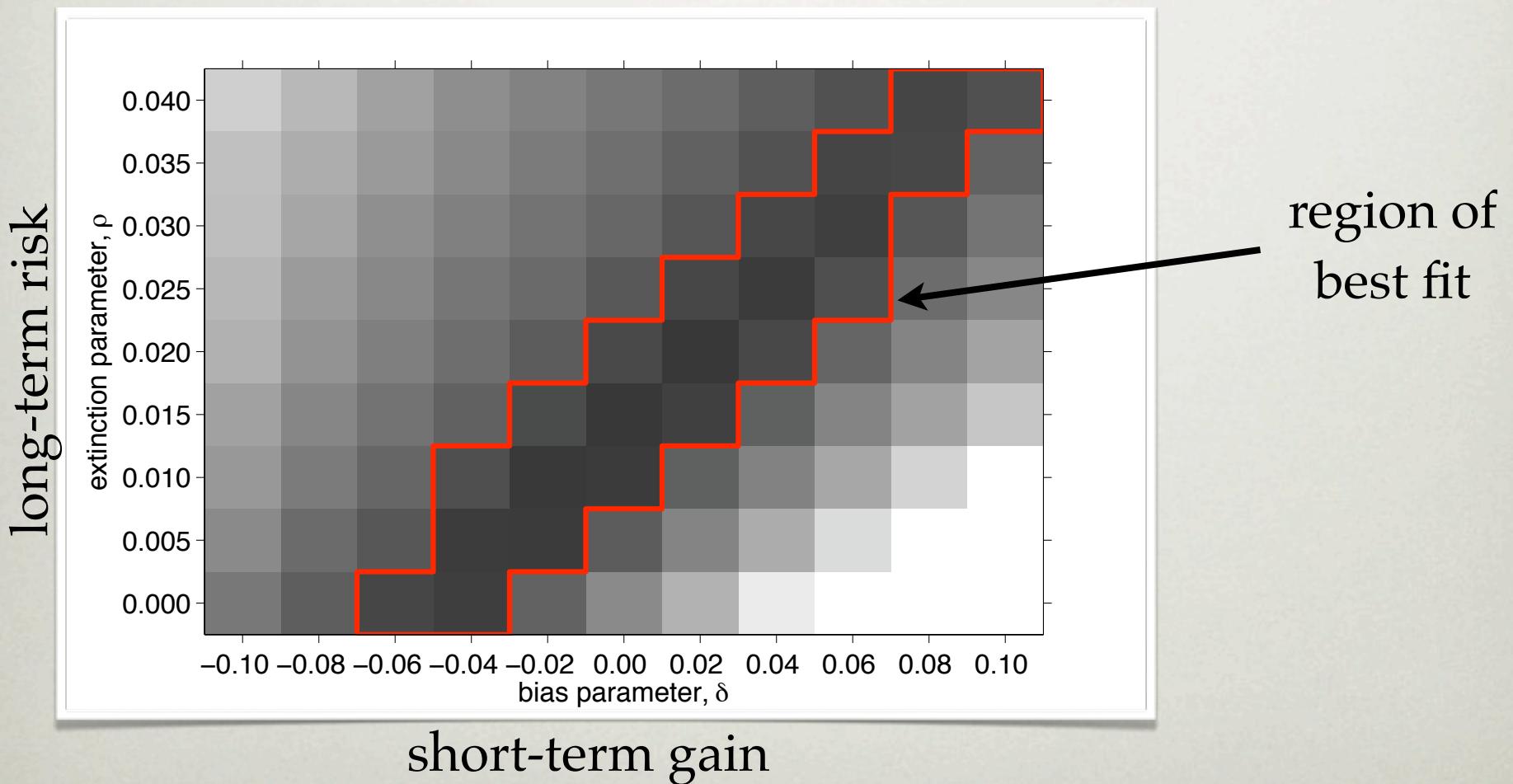
# SIMULATION

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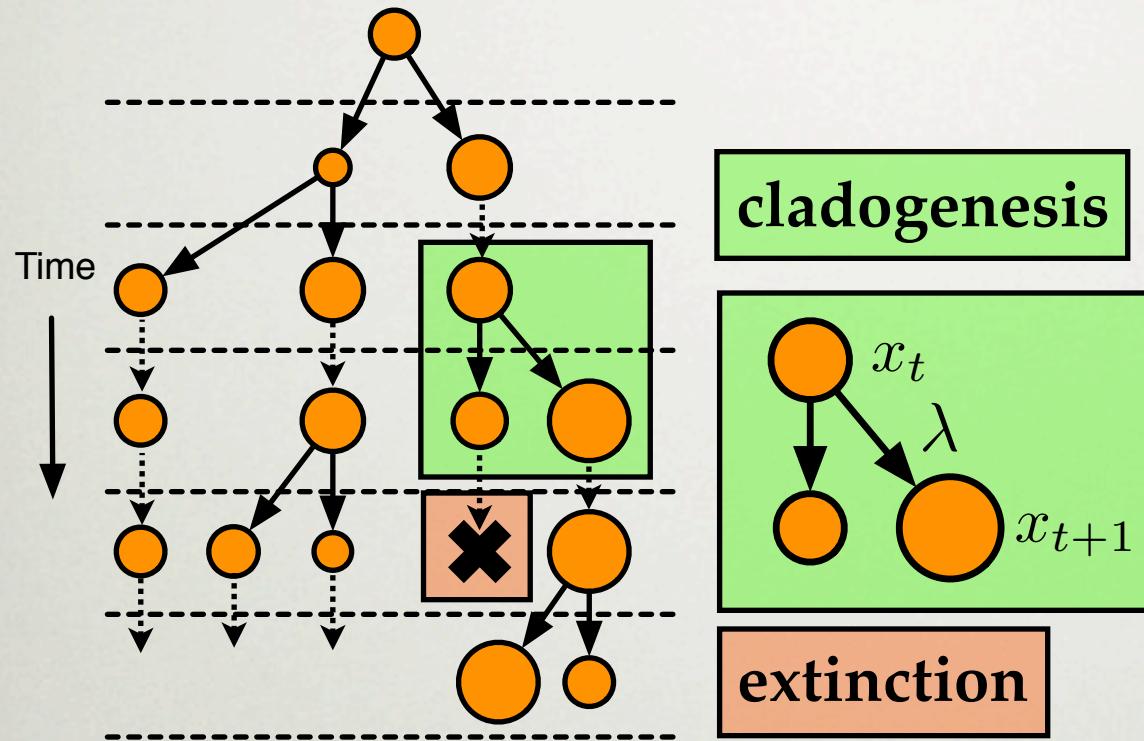
60 million years  
180,000 species



# SHORT-TERM GAIN VS. LONG-TERM RISK



# CLADOGENETIC MODEL



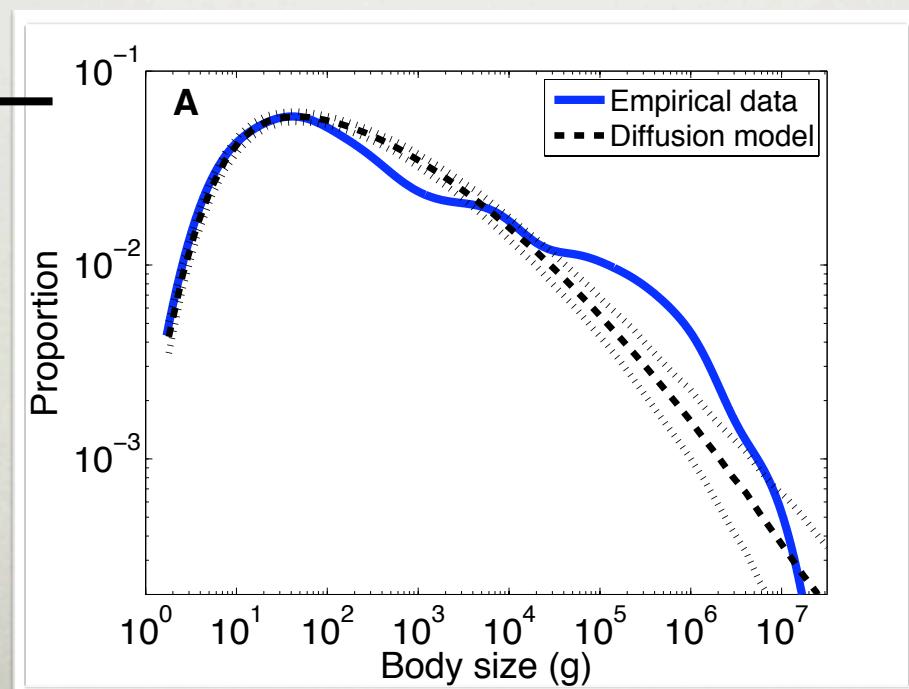
## model features

- size-dependent fluctuations
- Cope's rule
- size-dependent extinction rate
- lower limit  $x_{\min}$

are all these features necessary?

# MODEL COMPARISON

model features	size-dependent extinction	lower limit $x_{\min}$	$\langle wKS \rangle$	support
Cope's rule	█	█	0.181	good
		█	2.97	poor
	█	█	11.72	none
		█	10.37	none



# ANALYTIC VERSION

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reaction-diffusion-convection equation:

$$\frac{\partial c}{\partial t} + v \frac{\partial c}{\partial x} = D \frac{\partial^2 c}{\partial x^2} + (k - A - Bx)c$$

↑                    ↑                    ↑  
drift term        diffusion term        speciation  
(Cope's rule)      term                  and extinction

3 parameters:  $\beta = B/D$

$$\left. \begin{array}{l} \mu = v/D \\ x_{\min} \end{array} \right\}$$

estimated from  
fossil data

# ANALYTIC VERSION

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**Reaction-diffusion-convection equation<sup>\*</sup>:**

$$\frac{\partial c}{\partial t} + v \frac{\partial c}{\partial x} = D \frac{\partial^2 c}{\partial x^2} + (k - A - Bx)c$$

**Solving yields size distribution:**

$$c(x, t) = \sum_n \mathcal{C}_n(x_0) \mathcal{C}_n(x) e^{-\gamma_n t}$$

where

$$\mathcal{C}_n(x) \propto e^{\mu x / 2} \text{Ai} \left[ z_n + (B/D)^{1/3} (x - x_{\min}) \right]$$

$$z_n = z_0 - \frac{\gamma_n}{D(B/D)^{2/3}}$$

<sup>\*</sup> no “small-size bias” here

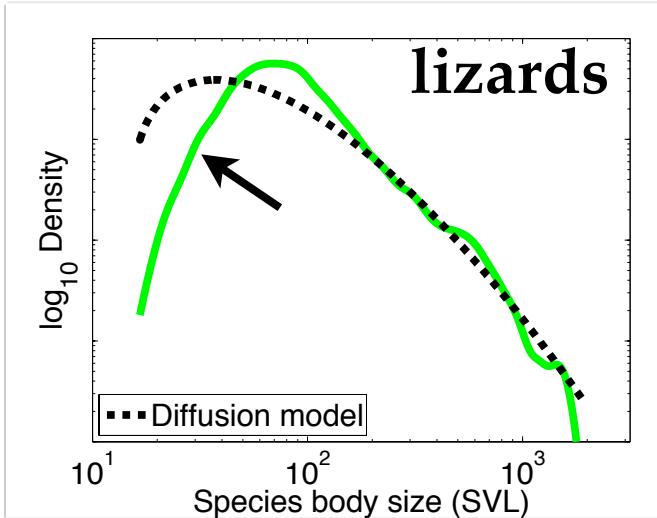
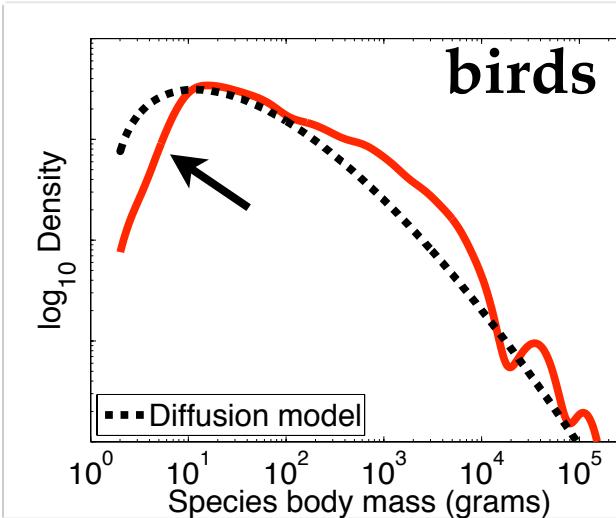
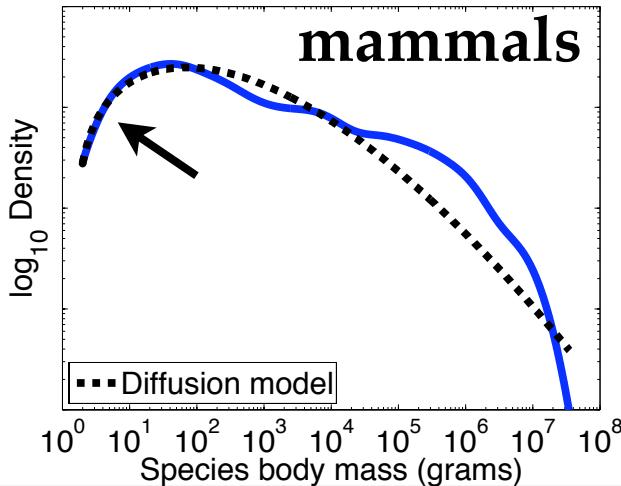
# ANALYTIC VERSION

Reaction-diffusion-convection equation\*:

$$\frac{\partial c}{\partial t} + v \frac{\partial c}{\partial x} = D \frac{\partial^2 c}{\partial x^2} + (k - A - Bx)c$$

Solving yields size distribution:

$$C_n(x) \propto e^{\mu x/2} \text{Ai} \left[ z_n + \beta^{1/3}(x - x_{\min}) \right]$$



\* no “small-size bias” here

# SIZE DIVERSIFICATION

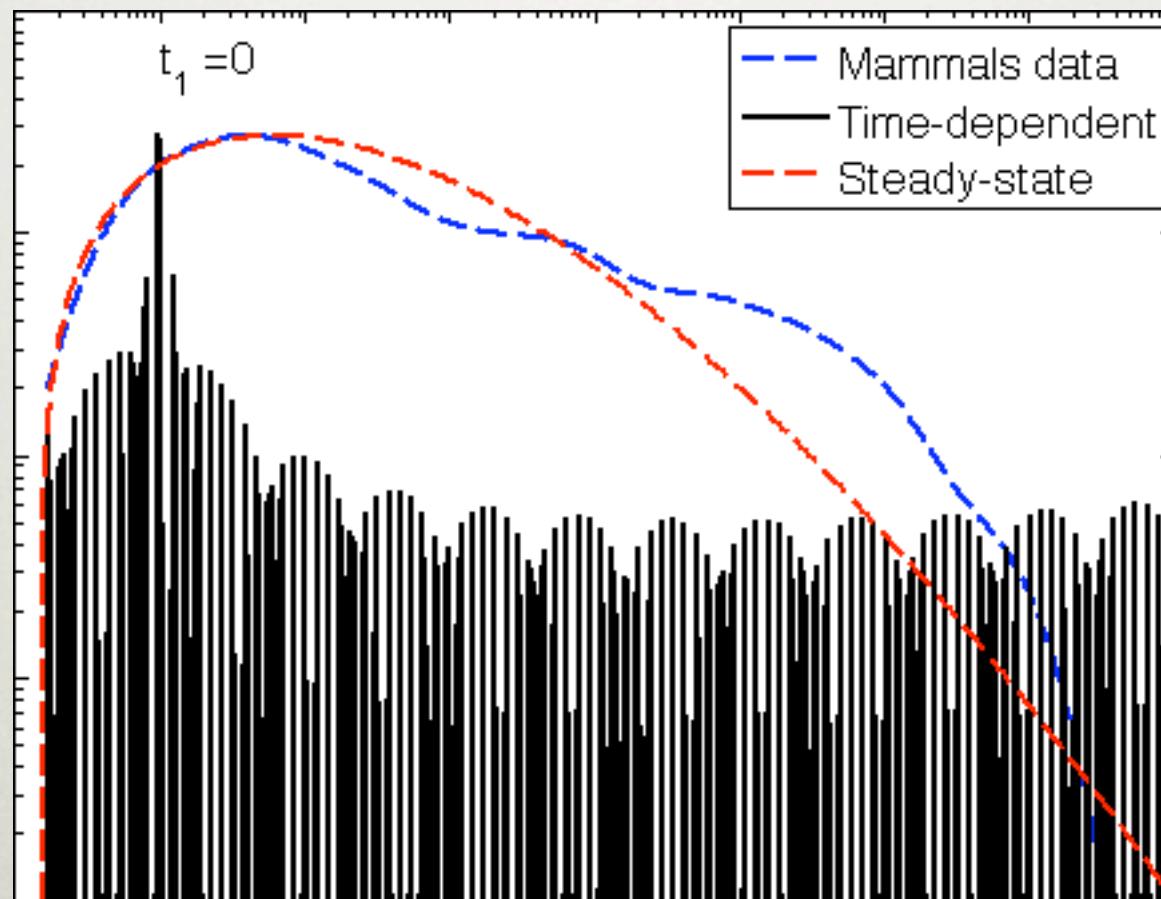
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simulate model dynamics

# SIZE DIVERSIFICATION

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simulate model dynamics



initially, 100,000 modes in eigen summation

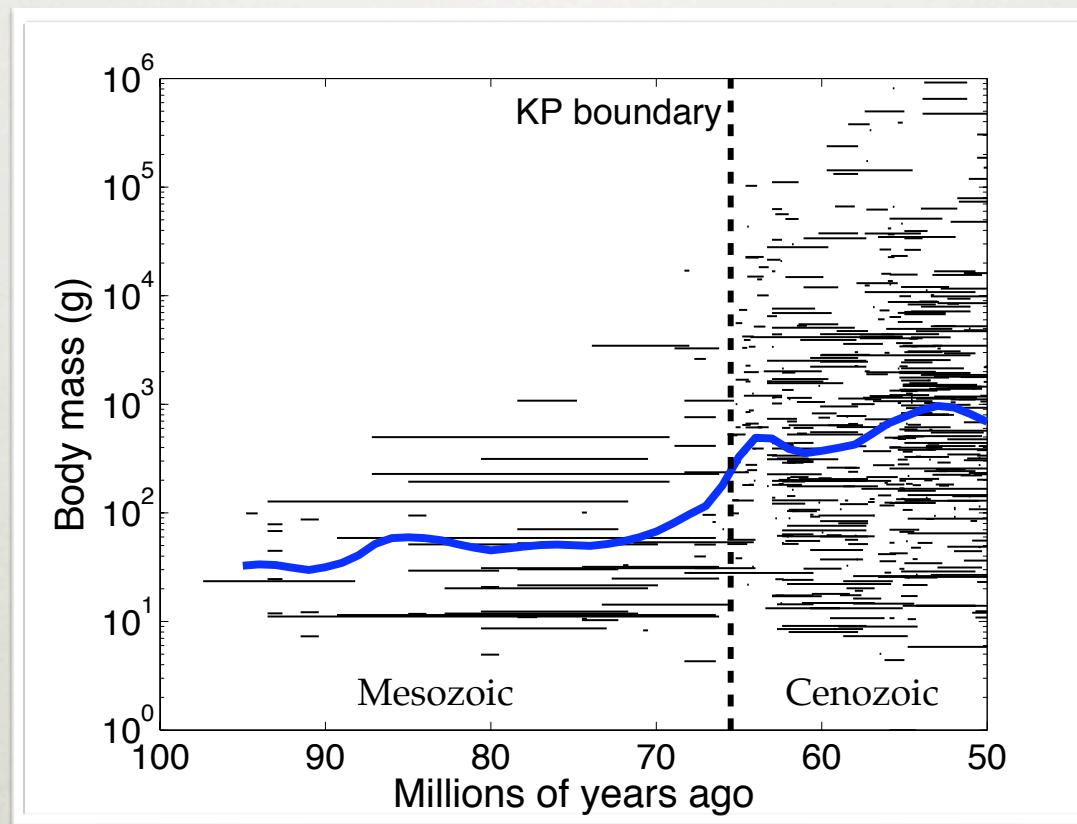
# SIZE DIVERSIFICATION

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but, do dynamics agree with fossil data?

# SIZE DIVERSIFICATION

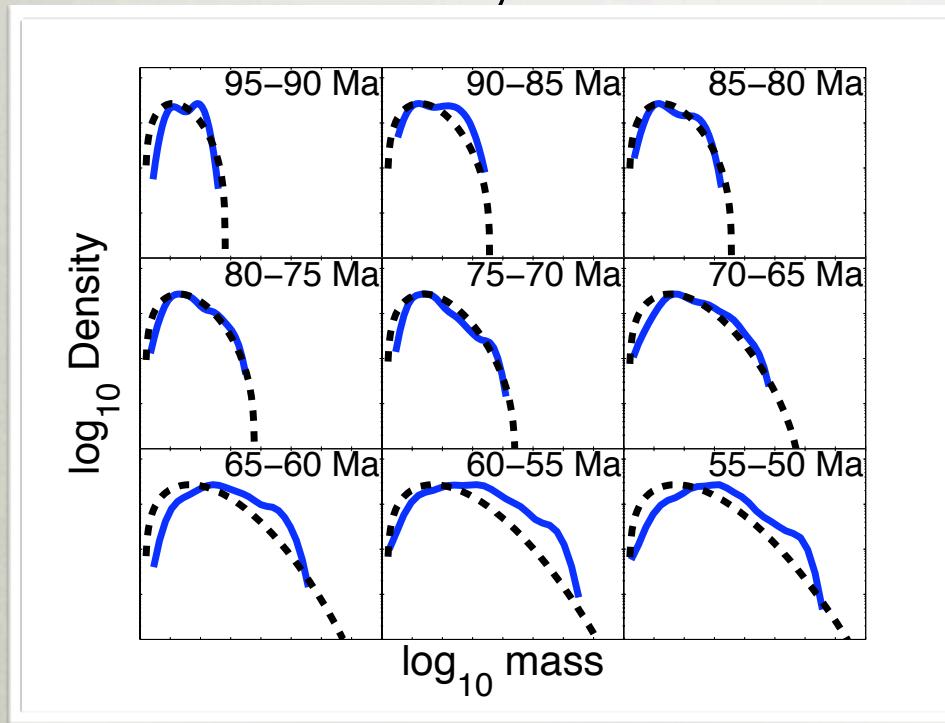
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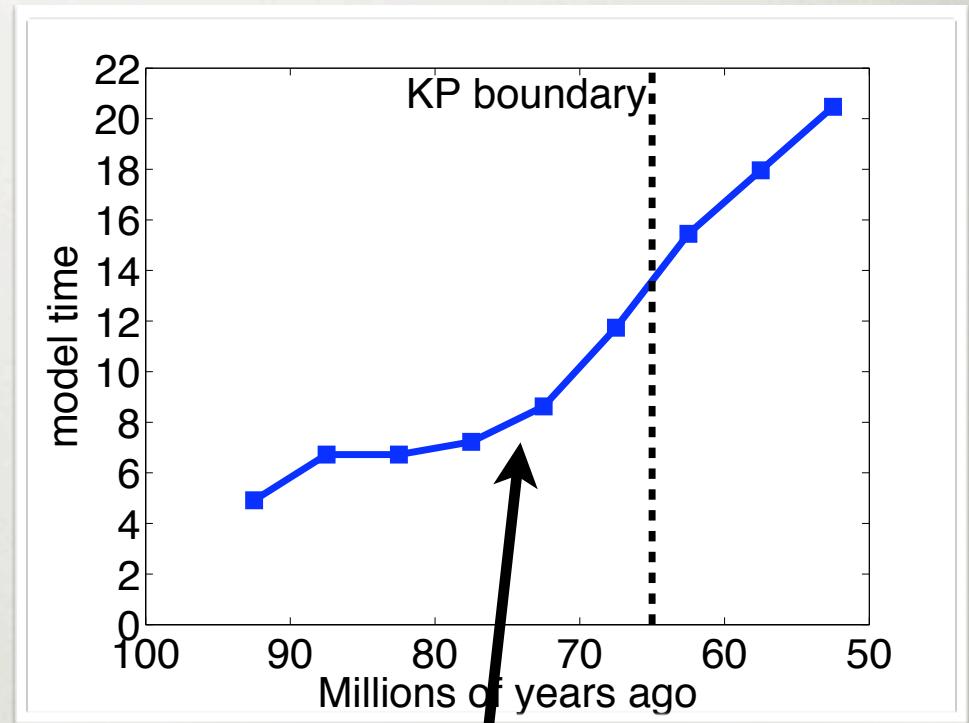
[North Am. terrestrial mammals]

# SIZE DIVERSIFICATION

model dynamics



model time



agrees with molecular-clocks  
for genetic diversification

# THURSDAY

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Southern Right whale (photo credit: Brian Skerry)