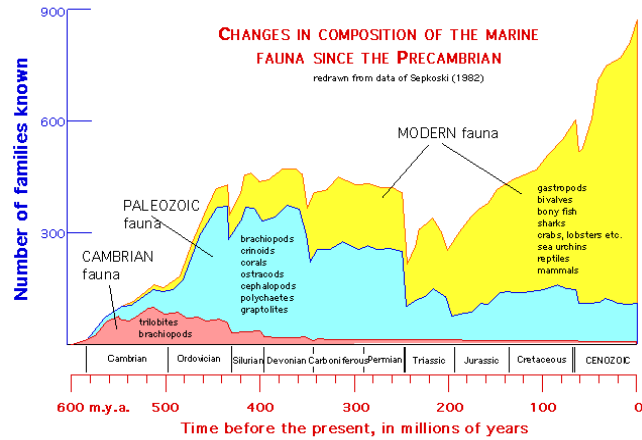


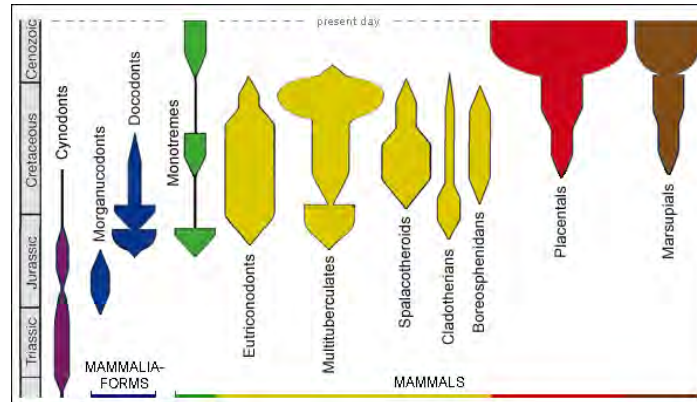
# Phenotypic Diversity in Macroevolution

# Taxonomic Diversity

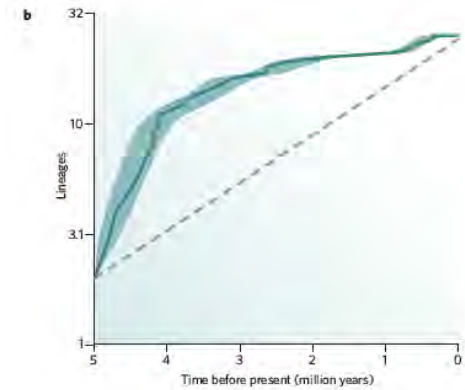
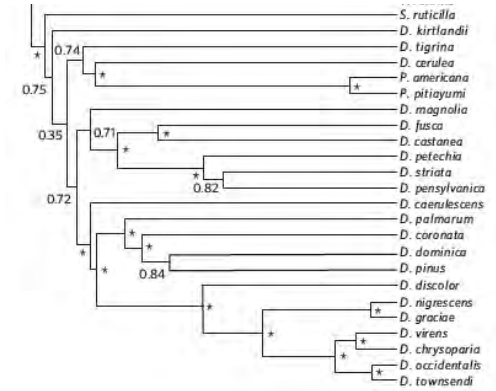
## Patterns of taxonomic diversity through time



after Sepkoski (1982)



Luo (2007)



Reznick and Ricklefs 2009

# Phenotypic Diversity

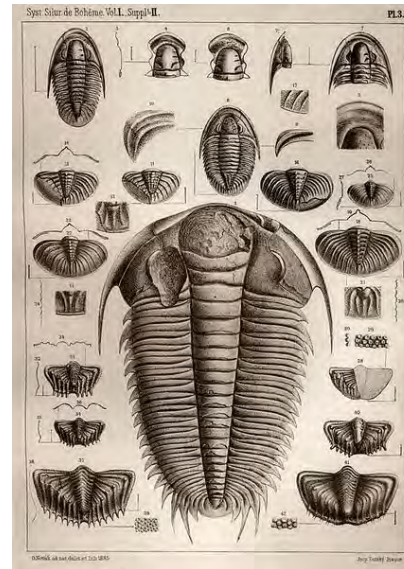
What about phenotypic diversity?



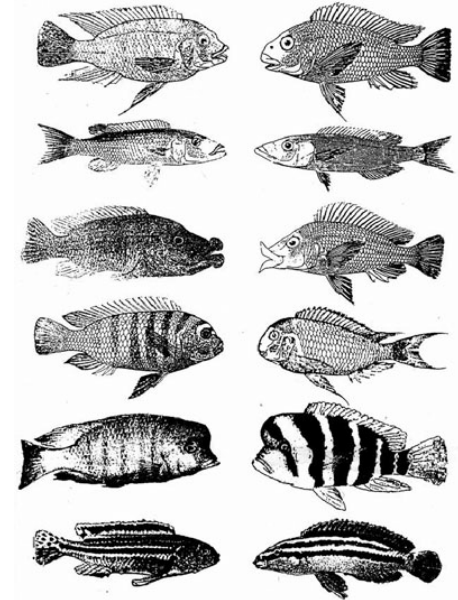
[www.ucmp.Berkeley.edu/taxa/inverts/Mollusca/gastropoda.php](http://www.ucmp.Berkeley.edu/taxa/inverts/Mollusca/gastropoda.php)



[dbs.umt.edu/research\\_labs/fishmanlab](http://dbs.umt.edu/research_labs/fishmanlab)



Wikimedia: from Barrande (1852)



Kocher et al. (1993)

-Understanding trends requires a quantification of phenotype

# Quantifying Phenotypic Diversity

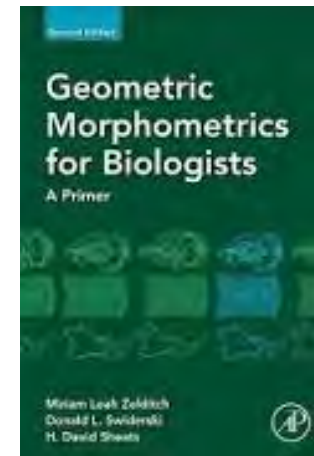
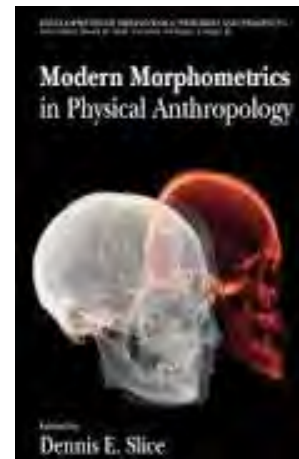
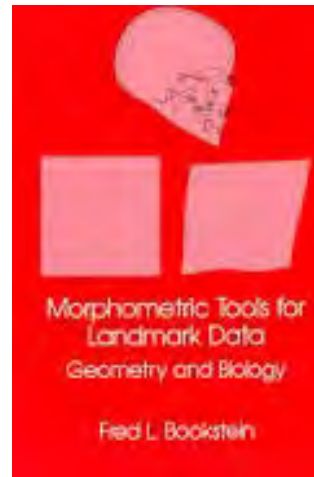
Challenge:

-How do we quantify morphology & phenotypic diversity?

Morphometrics (*morpho* = form; *metrics* = measure)

-The quantification of morphology

-The study of phenotypic variation and covariation

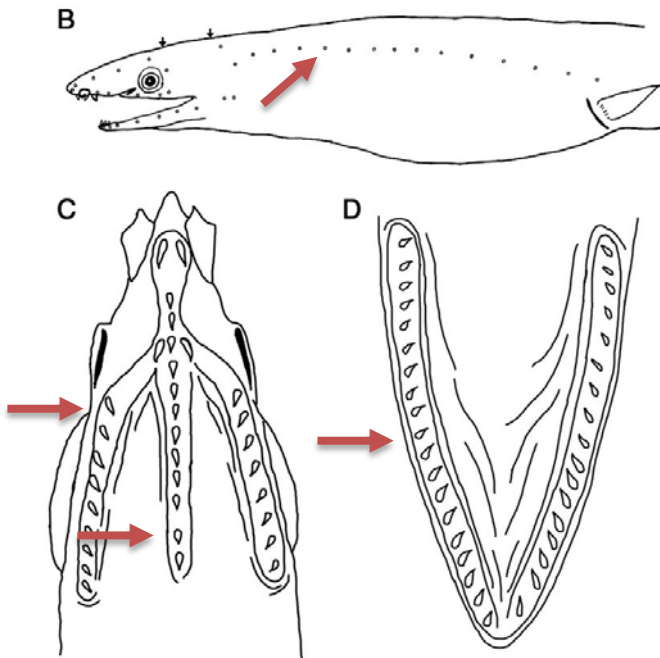


Different types of variables (and methods) have been used

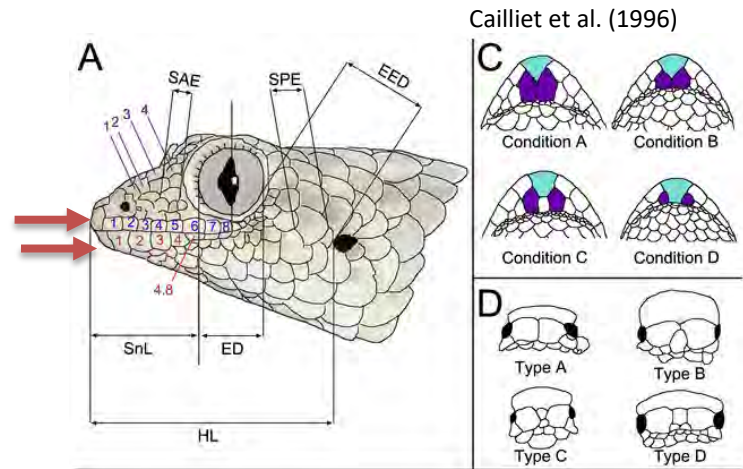
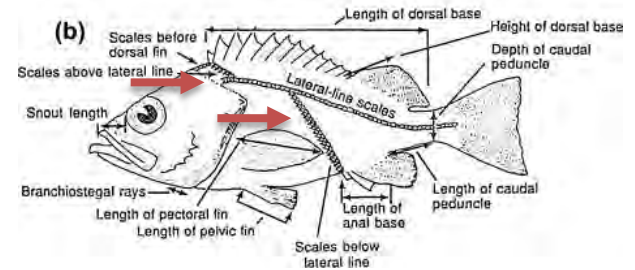
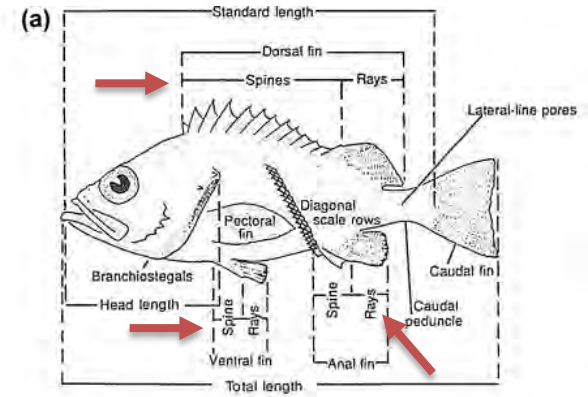
# Common Morphometric Data

## Meristic count data\*

-#teeth, pores, fin rays, scales, etc.



Ho et al. (2012)



Cailliet et al. (1996)

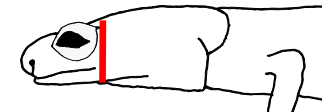
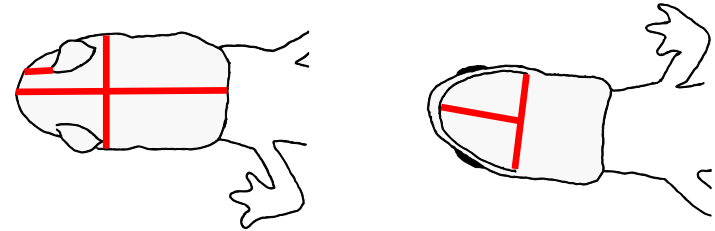
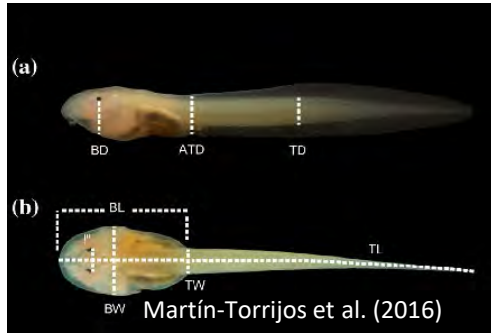
Scherz et al. (2017)

\*Frequently used for taxonomy and classification

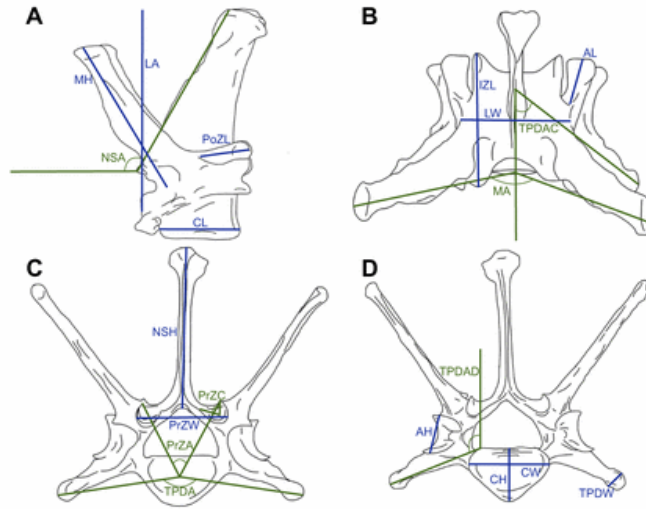
# Common Morphometric Data

## Linear measurements\*

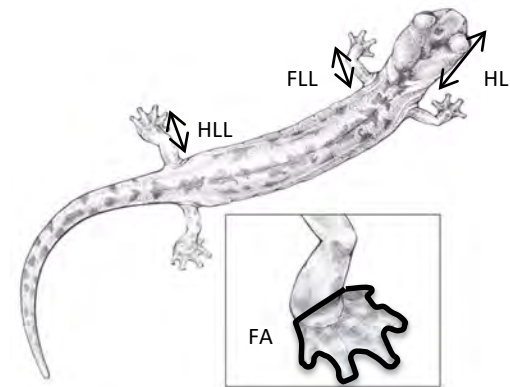
-Extents of, distances between, and angles among structures



Adams (2000)



Oliver et al. (2016)



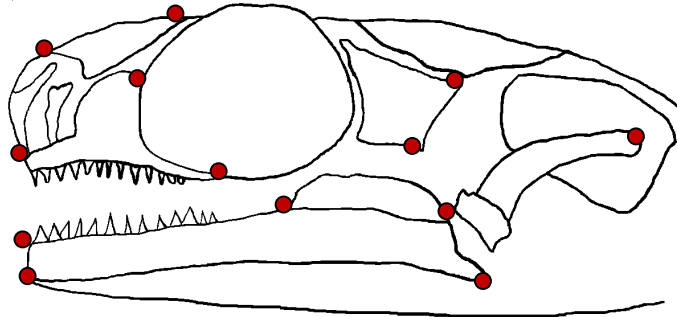
Adams et al. (2017)

\*Often called multivariate morphometrics

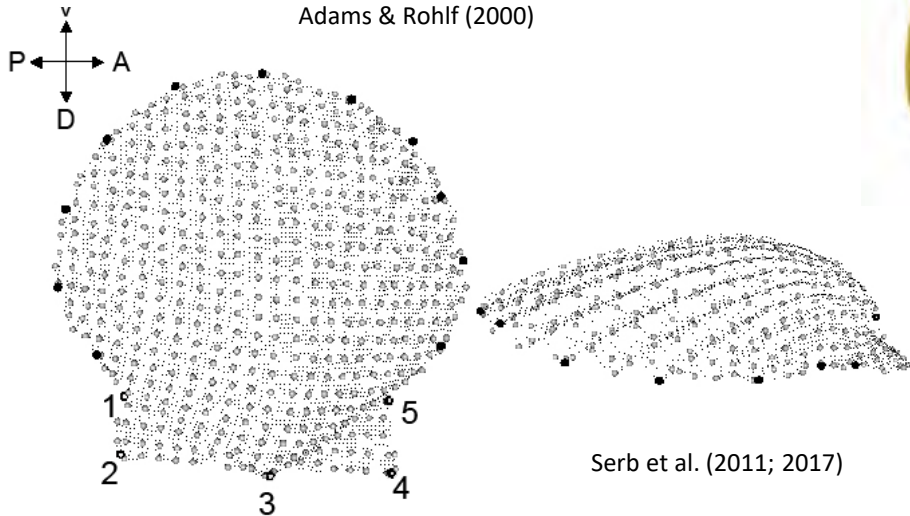
# Common Morphometric Data

## Geometric morphometrics\*

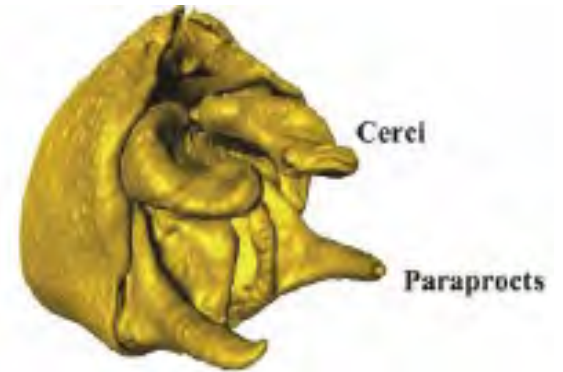
-Shape data from geometric variables



Adams & Rohlf (2000)



Serb et al. (2011; 2017)

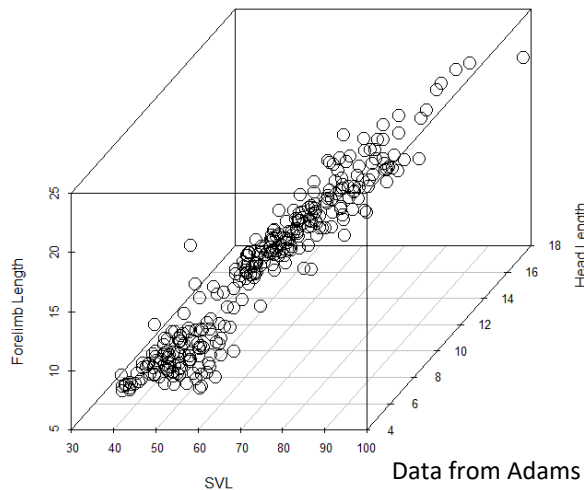
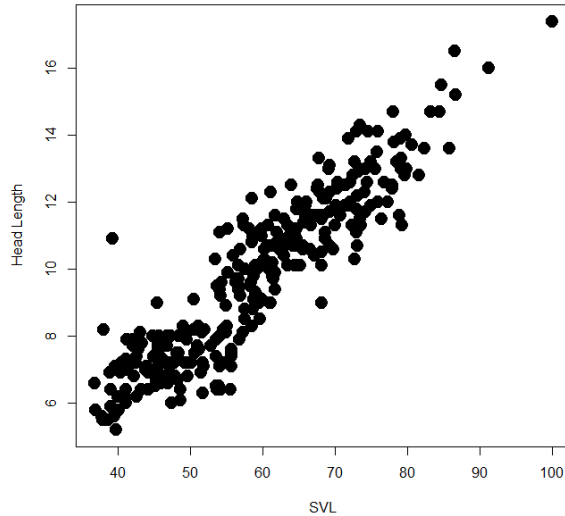


McPeck et al. (2008)

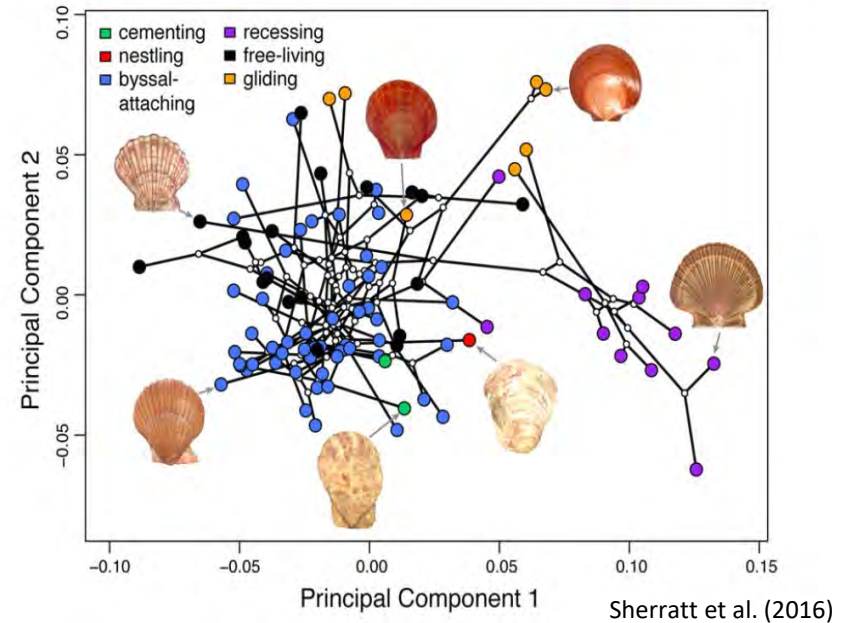
\*Highly multivariate, multi-dimensional data

What does one do with these variables?

-Morphospace: Dataspace summarizing morphological variables\*



Data from Adams (2000)

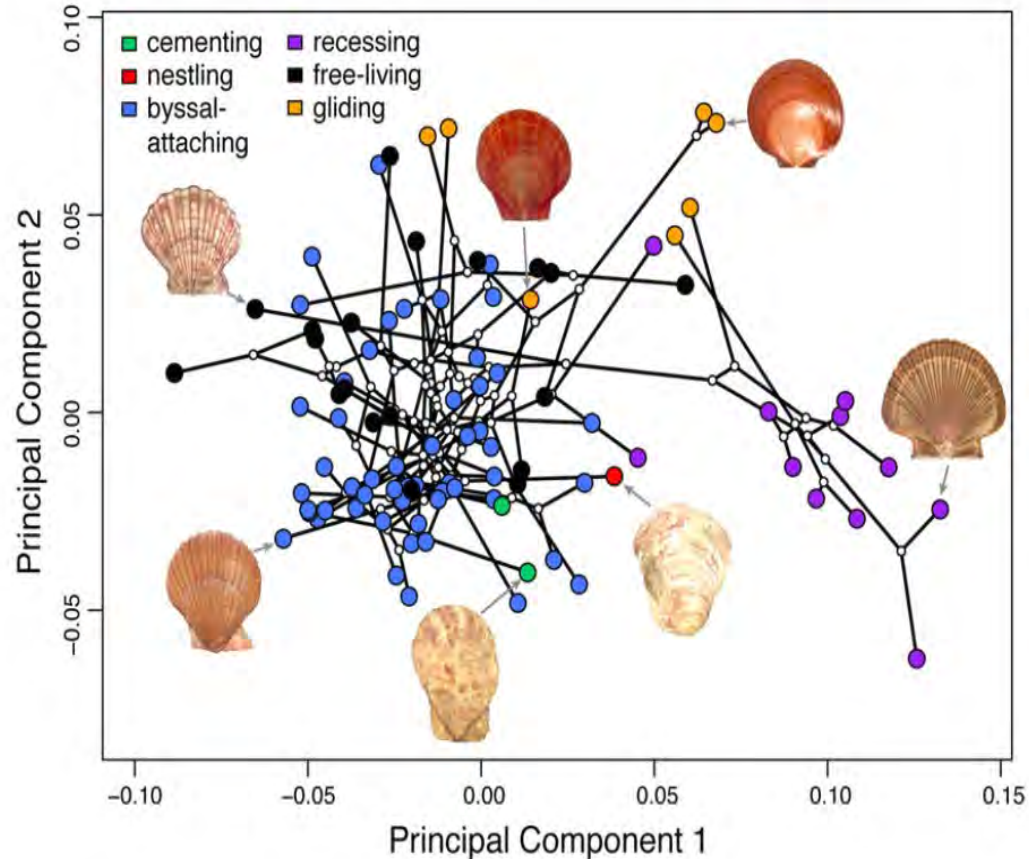


Sherratt et al. (2016)

\*PCA: principal components analysis often used to view space



## Morphospace with phylogeny superimposed

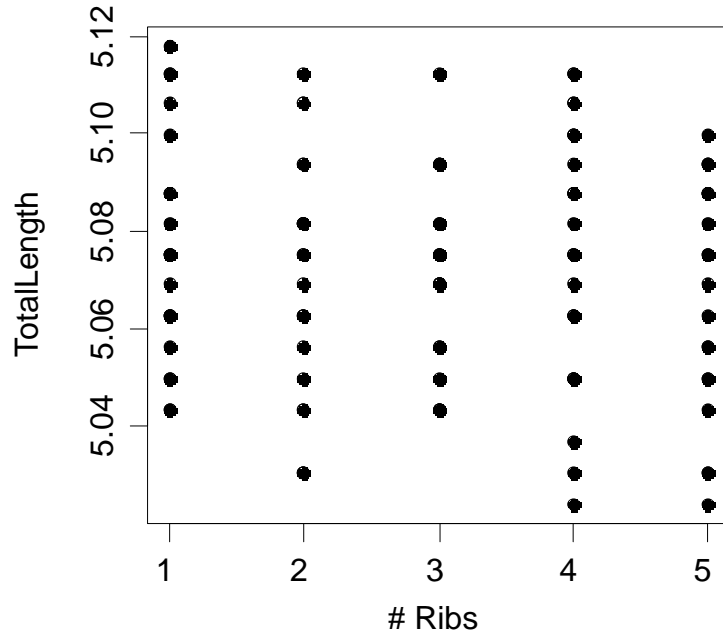


Sherratt et al. (2016)

-EXTREMELY useful for visual inspection of convergence, directional evolution, etc.

# Morphospaces: Careful in Application!

Axes of morphospace **MUST** be in commensurate units



-Technically, this is a morphospace  
(all variables are numeric)  
-but is total GIGO!!!

-Why? Axes are incommensurate units & scale

-Variances, covariances, disparity, distances, etc. have no meaning in this space

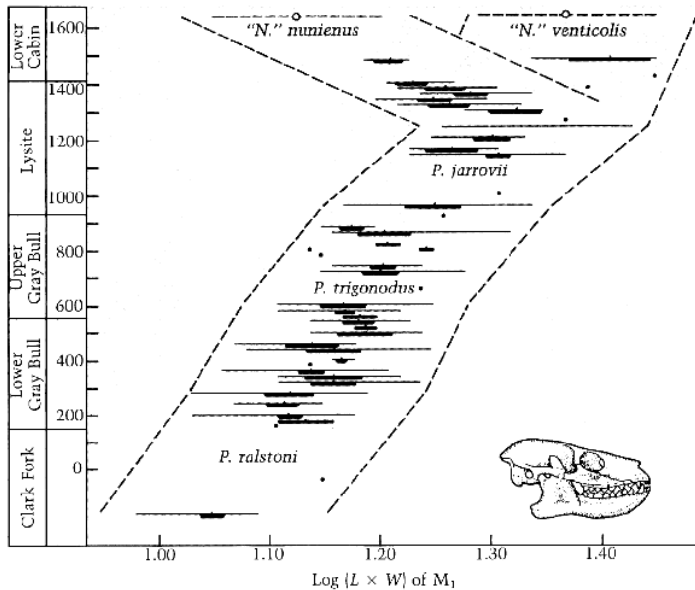
-Morphospaces should only be constructed with data in commensurate units!!!

# Some Phenotypic Trends

# Phenotypic Trends: Gradualism

## Phyletic gradualism

- $H_0$ : evolution is slow and gradual (ala Darwin's suggestion)
- Small changes accumulate over time
- Speciation from gradual accumulated divergence



Data from Gingerich (1976)  
Image: Gould & Eldredge (1977)

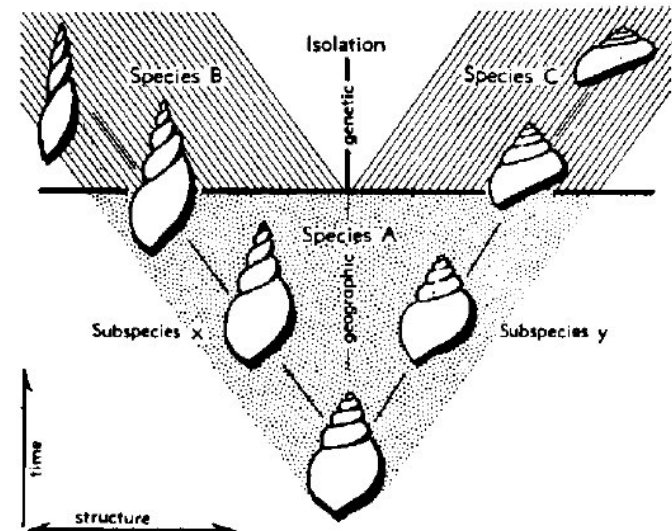


Figure 5-3:

A hypothetical case of geographic speciation viewed from the perspective of phyletic gradualism—slow and gradual transformation in two lineages.

From Moore, Lalicker, and Fischer, 1952; figure 1-15.

# Phyletic Gradualism: Anagenesis

Anagenesis: Phenotypic change within species over time

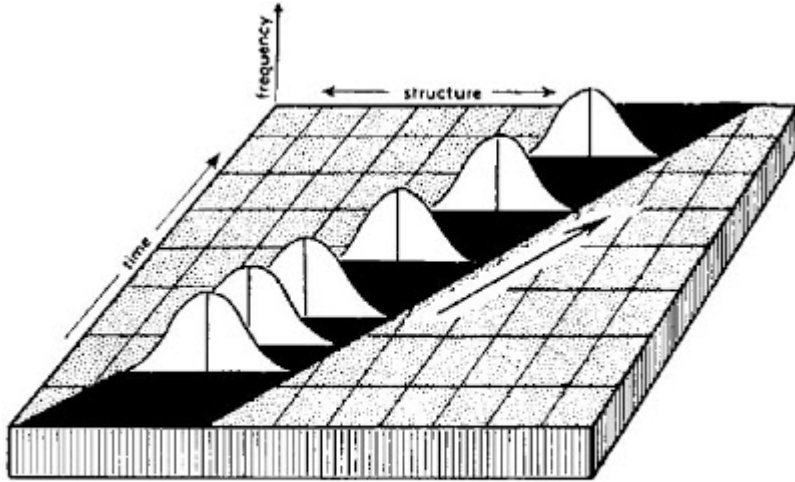
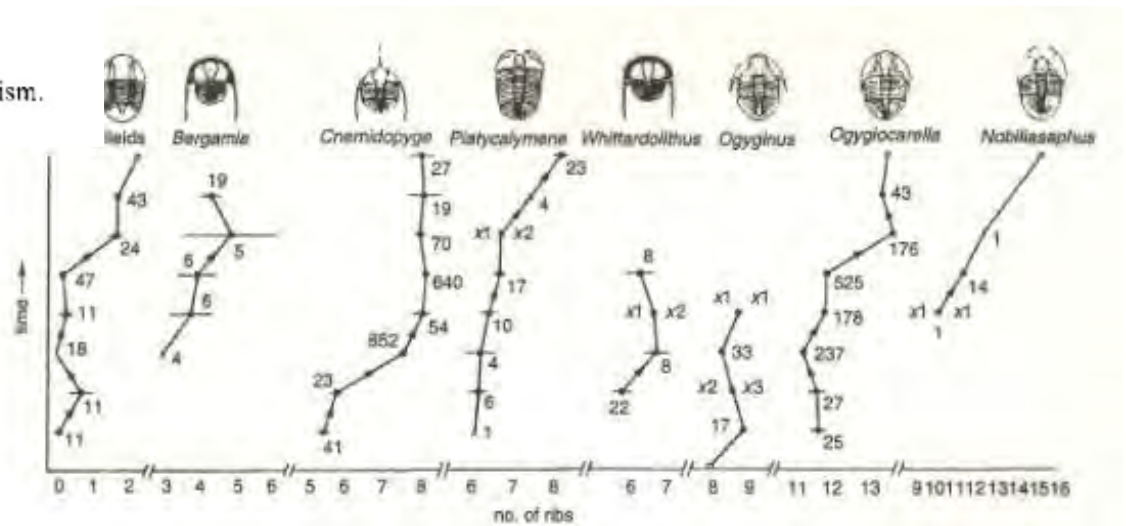
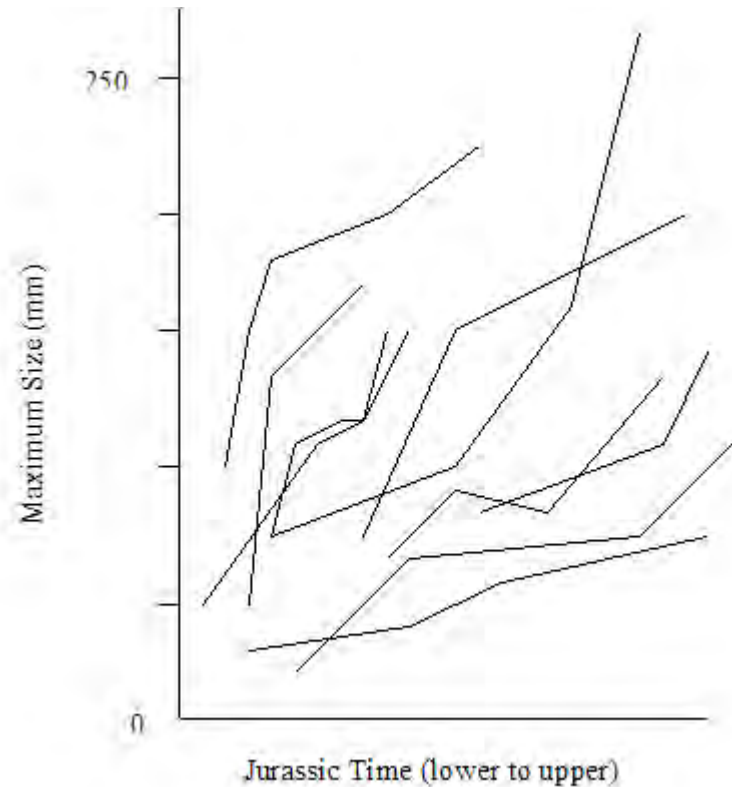
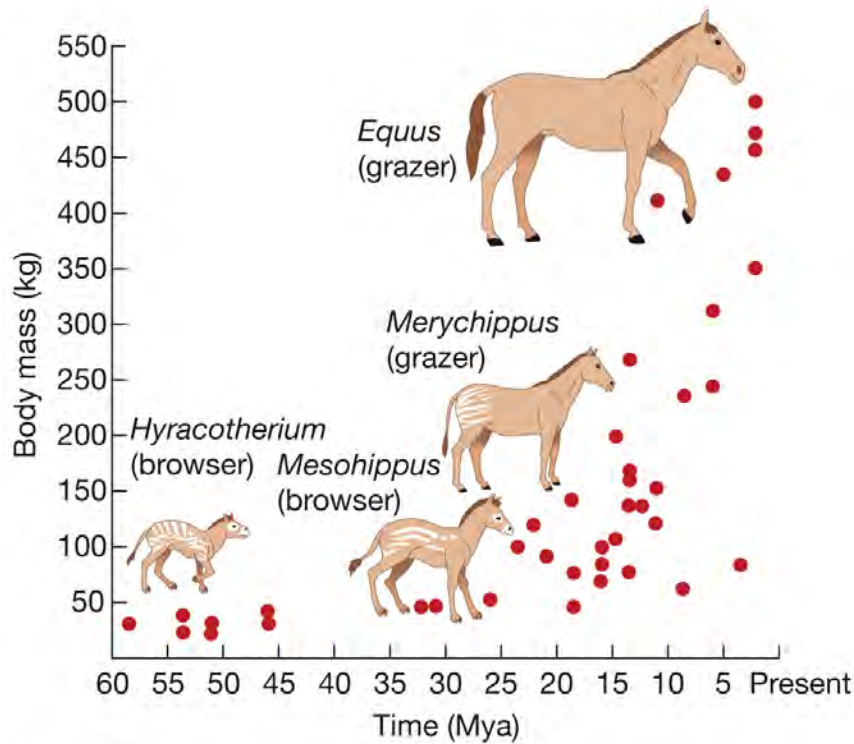


Figure 5-2:  
A standard textbook view of evolution *via* phyletic gradualism.  
From Moore, Lalicker, and Fischer, 1952; figure 1-14.

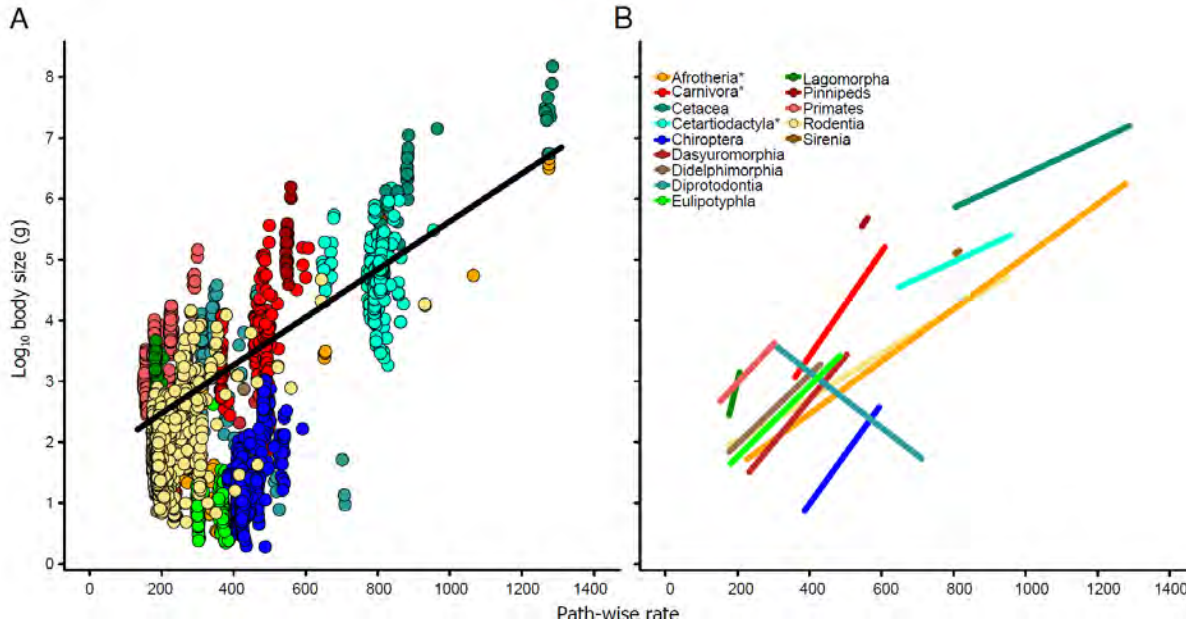
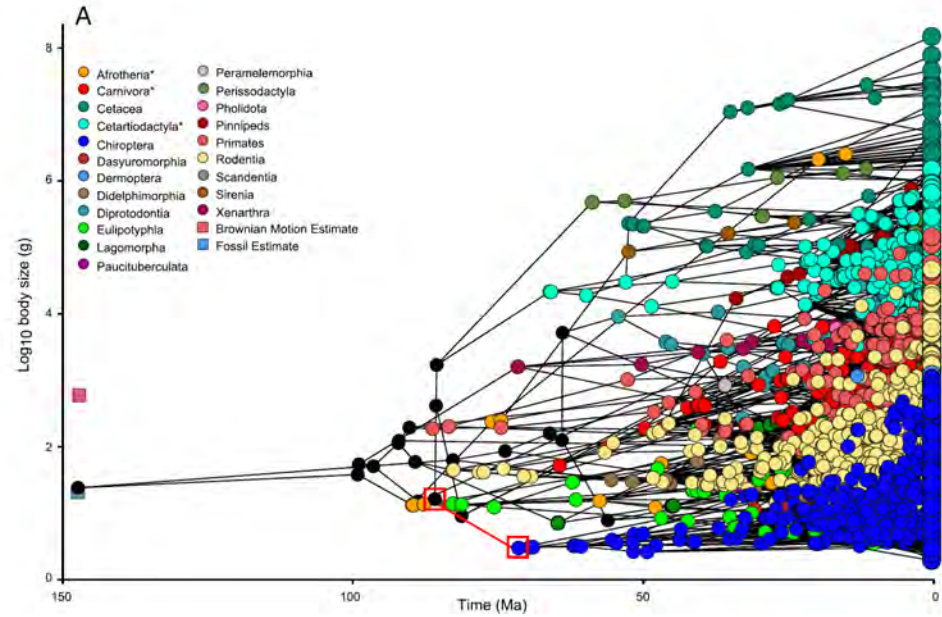


## Cope's rule

- Increase in body size in clade over time
- Frequently linked with gradualism

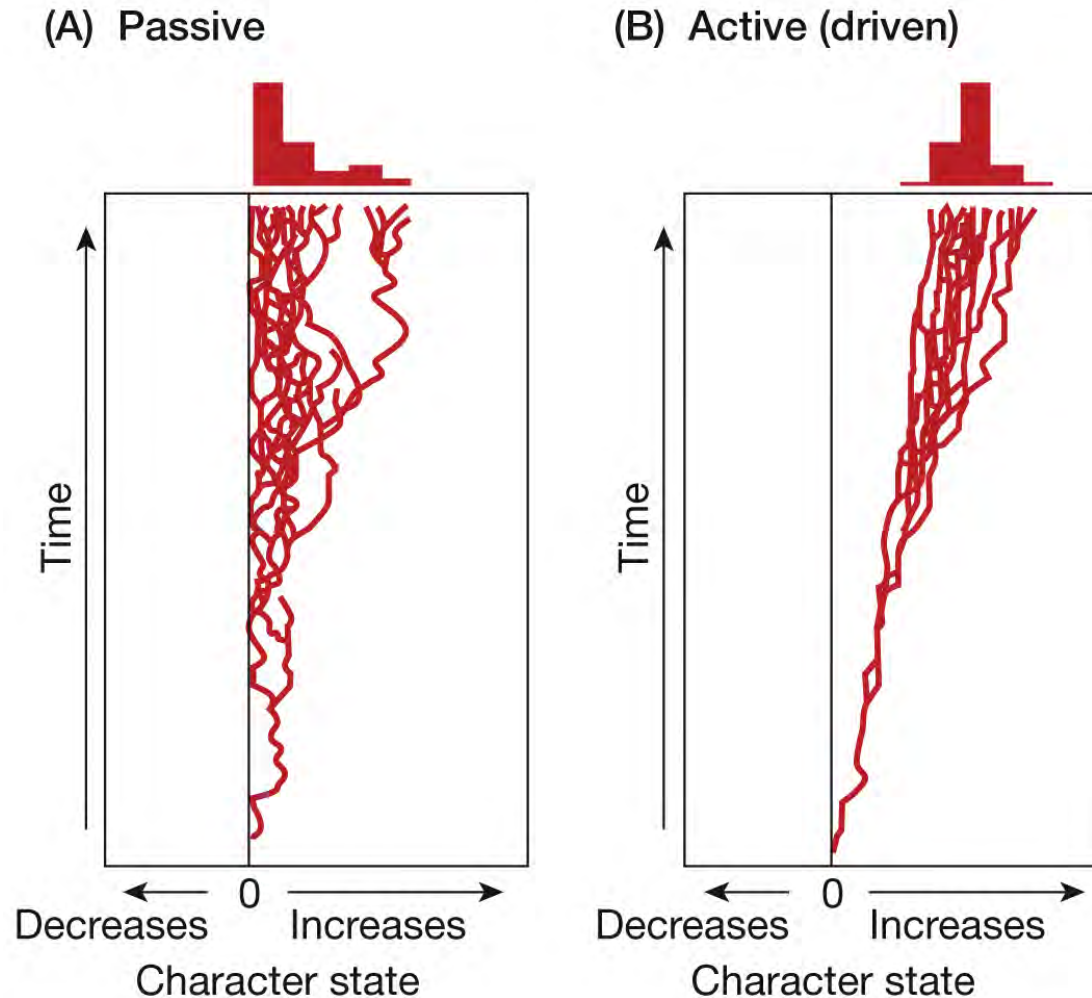


## A neontological example



Note that directional trends may take two forms: active and passive

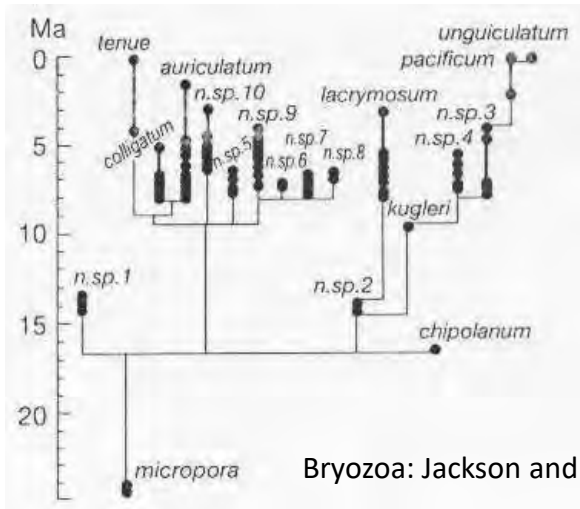
(the latter when trait value 'bump' into some limit over time)



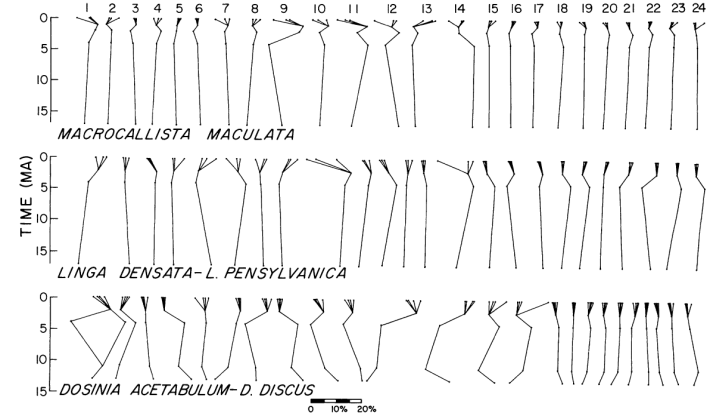
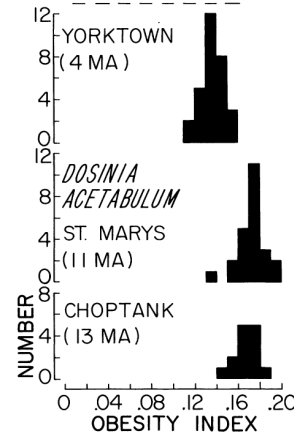


# Phenotypic Trends: Stasis

Stasis: Much of fossil record shows little change

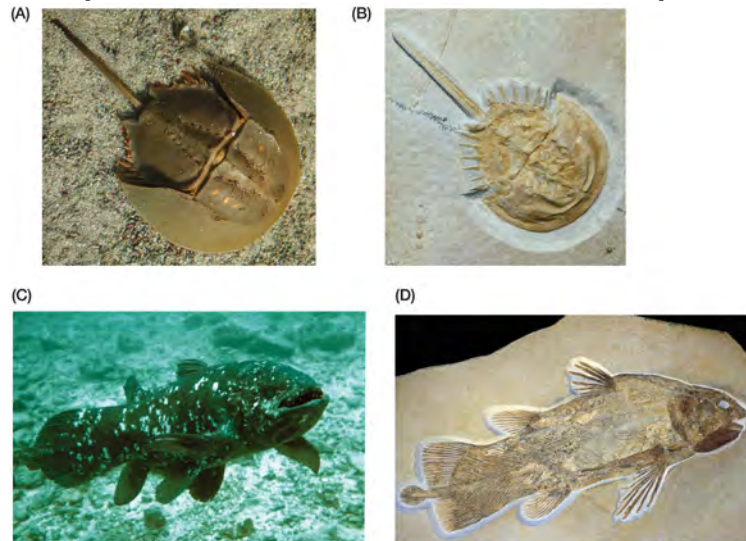


Bryozoa: Jackson and Cheetham (1994)



Bivalves: Stanley and Yang (1987)

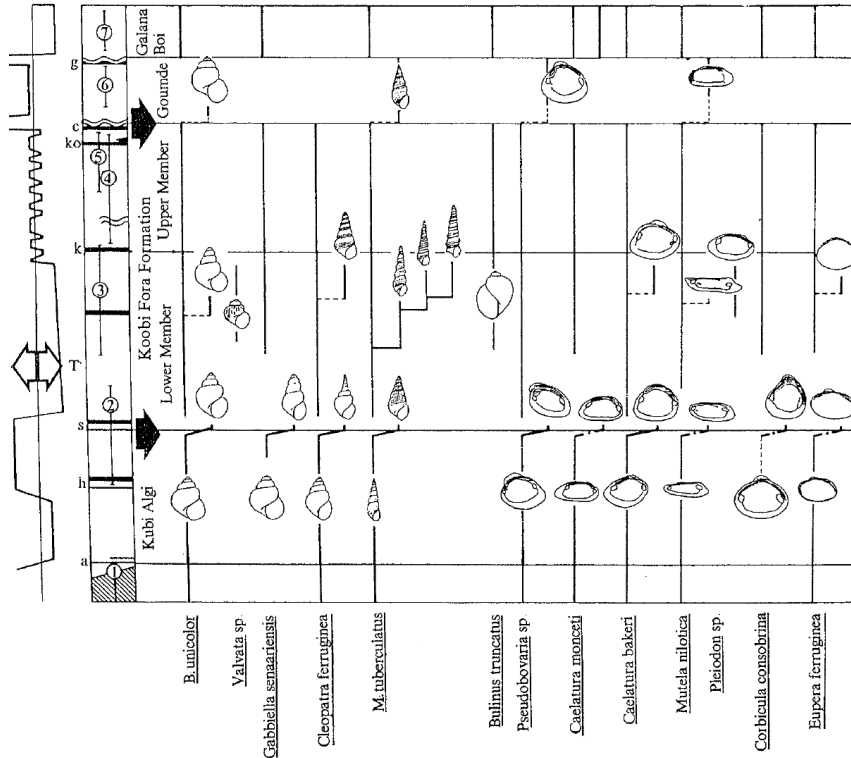
So called "Living fossils" provide another example



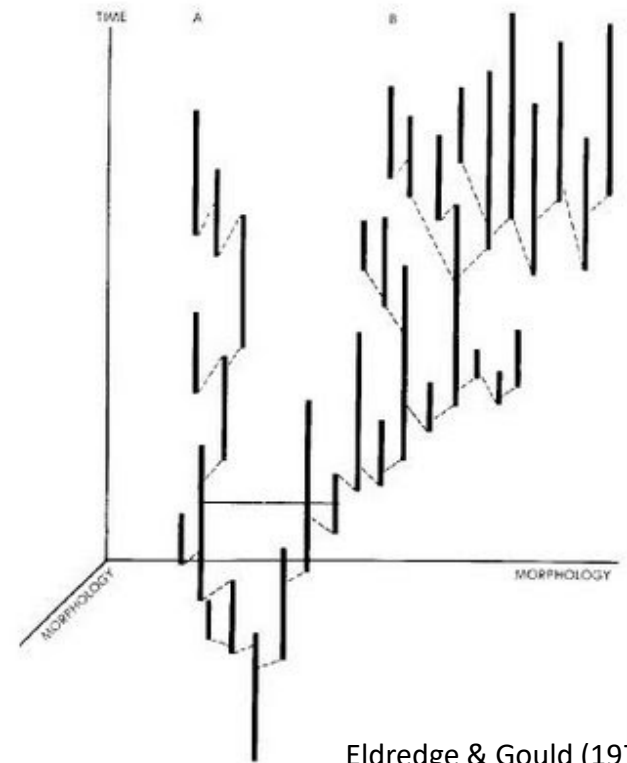
# Punctuated Equilibrium

## Punctuated Equilibrium: Stasis followed by rapid change

-Gould & Eldredge (1972) argued PE better explains many fossil trends



Williamson (1981)



Eldredge & Gould (1972)

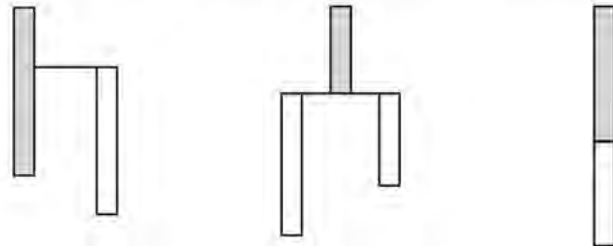
\*Note manner in which branching in phylogeny is depicted!

-Subtle implications: 1) change is punctual; 2) speciation is 'budding off' from ancestor, not 2 new descendants

# Speciation in the Fossil Record

As fossil species are defined phenotypically, linking trait change and lineage diversification results in distinct speciation modes

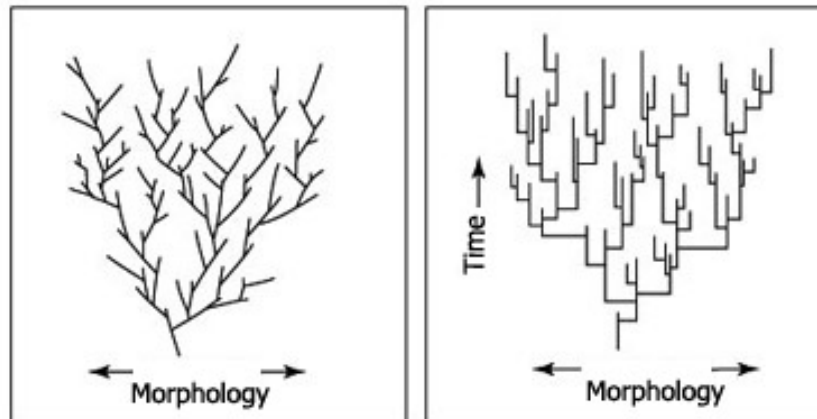
(i) asymmetric speciation (ii) symmetric speciation (iii) anagenetic speciation



Stadler et al. (2018)

after Foote (1996)

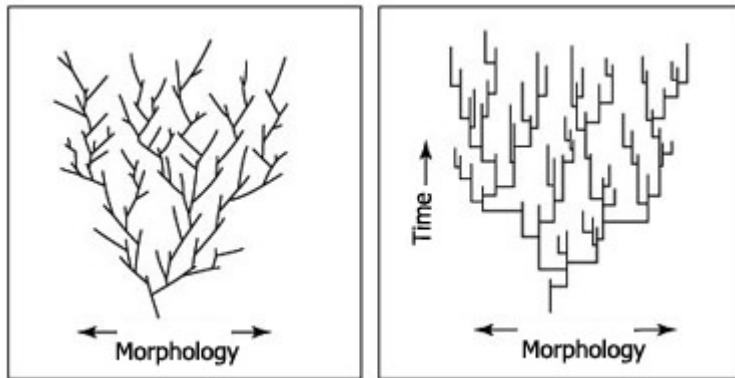
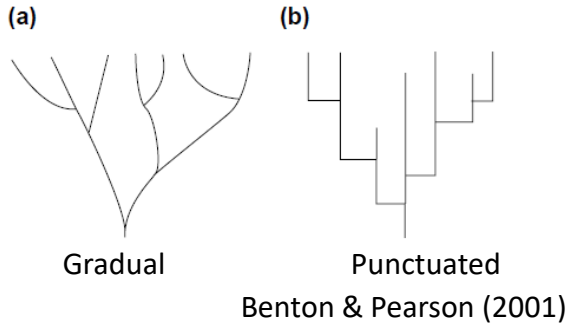
Neontologists & paleontologists often depict speciation differently  
(has implications for how we view phenotypic trends)



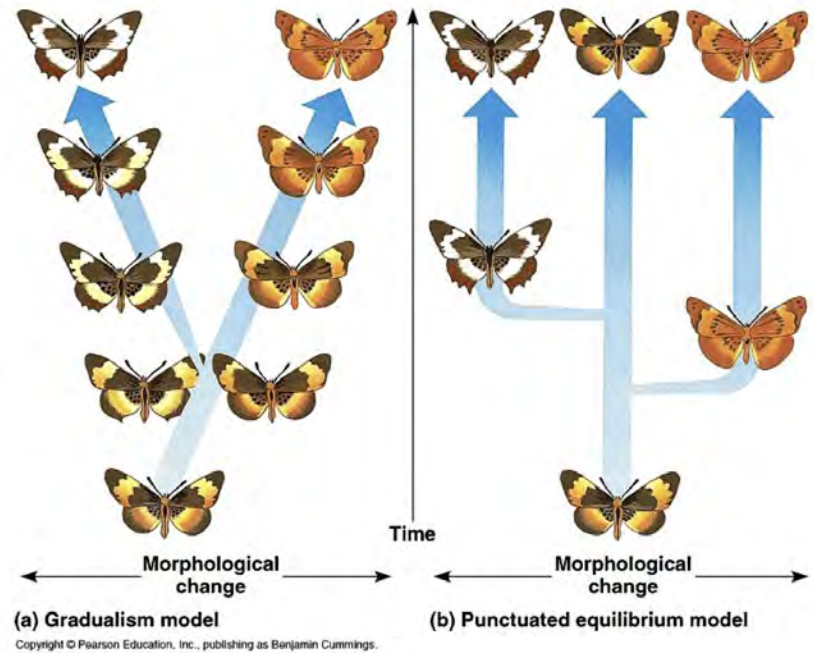
Freeman & Herron (1998)

# Distinguishing PE from Gradualism

Interpreting the fossil record can be challenging  
-Is the pattern punctuational or gradual?



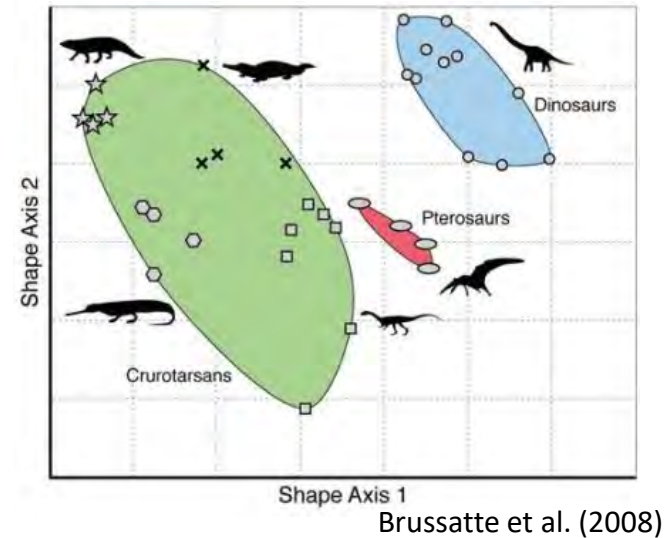
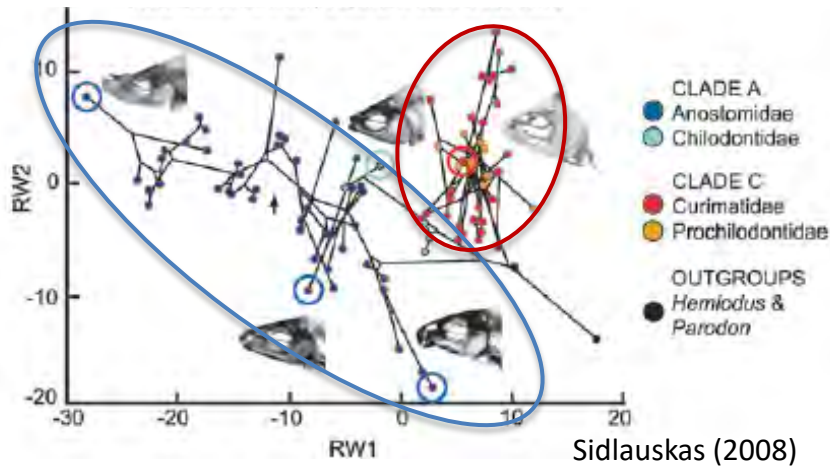
Freeman & Herron (1998)



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# Morphological Disparity

How much phenotypic diversity does a lineage display?



-Some lineages seem to occupy more of morphospace

Why? Hypotheses include:

-Constraints (competition, niche filling, biomechanical, etc.)

-Ecological release

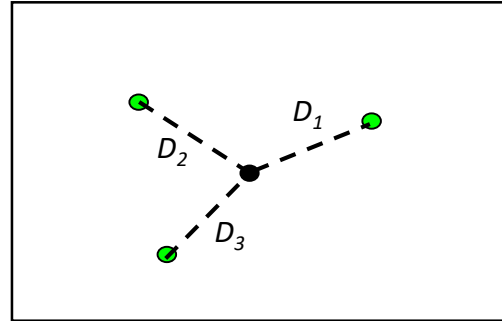
-Can this be quantified and compared?

# Quantifying Disparity

Morphological Disparity: a measure of phenotypic diversity\*

$$MD = \frac{\sum D_j^2}{N - 1}$$

$D_j$ : Distance from  $j^{\text{th}}$  object to centroid



Foote (1990; 1993)

MD is a measure of variance (for 1 trait it IS the variance)

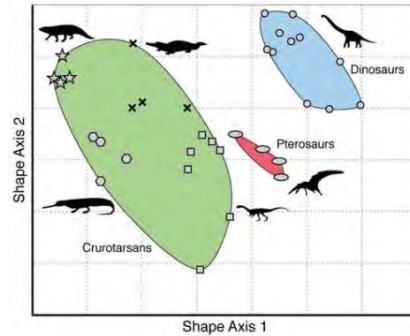
$$\text{Recall: } \sigma^2 = \frac{\sum (Y_j - \bar{Y})^2}{N - 1}$$

$$\sqrt{(Y_j - \bar{Y})^2} = D_j \text{ so } (Y_j - \bar{Y})^2 = D_j^2$$

\*One can obtain MD using pairwise distances among objects using SS → distance equivalency  
Gower, (1966); Anderson (2001); Adams (2014)

# Comparing Disparity

For multiple groups, which group displays greater MD?



Brussatte et al. (2008)

Compare MD statistically with permutation test

- 1) obtain  $MD_1, MD_2$ , etc.
- 2) calculate difference score:  $S = | MD_1 - MD_2 |$
- 3) Randomly assign taxa to groups, obtain MD and  $S_{perm}$
- 4) proportion  $S_{perm} > S_{obs}$  is level of significance

**NOTE: This MD test evaluates differences in *DISPERSION* (variance).  
Tests of *LOCATION* are performed using MANOVA!**

## Taxonomic and morphological disparity: Balstoidea and Trilobita

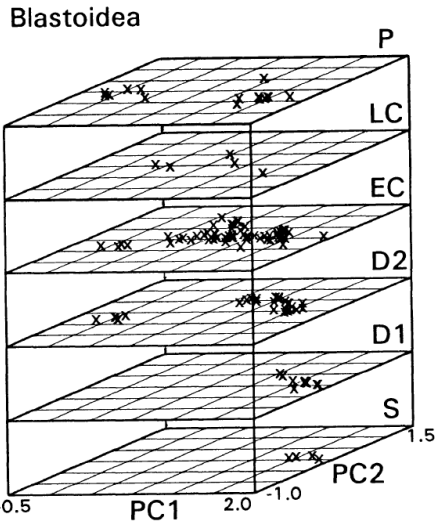


FIGURE 2. Temporal pattern of morphospace occupation in Blastoidea. Note overall increase in range of morphospace occupied. Early Carboniferous taxonomic diversity in Blastoidea.

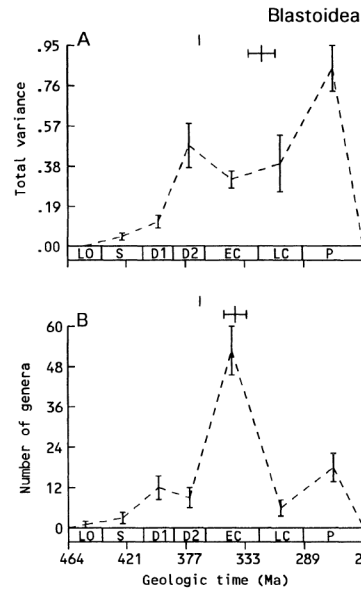


FIGURE 3. Comparison of morphological and taxonomic diversity in Blastoidea. Morphological diversity in this

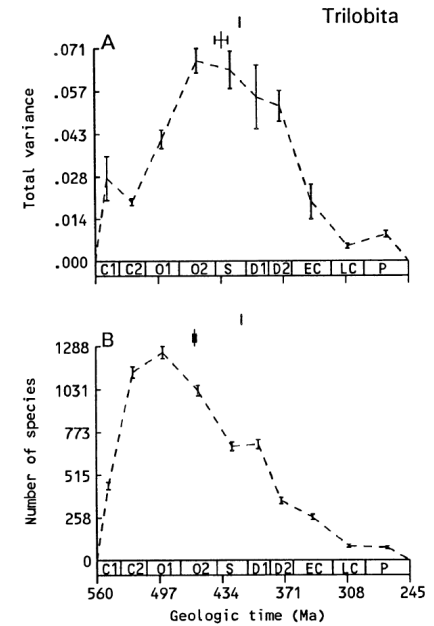
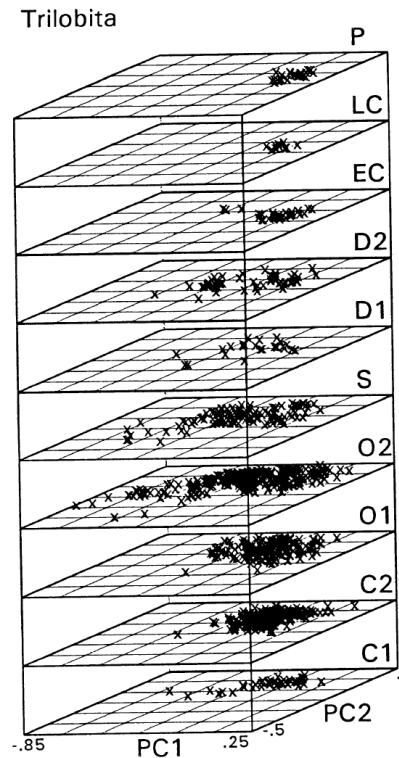


FIGURE 5. Comparison of morphological and taxonomic diversity in Trilobita. Morphological diversity is displaced forward in time relative to taxonomic diversity,

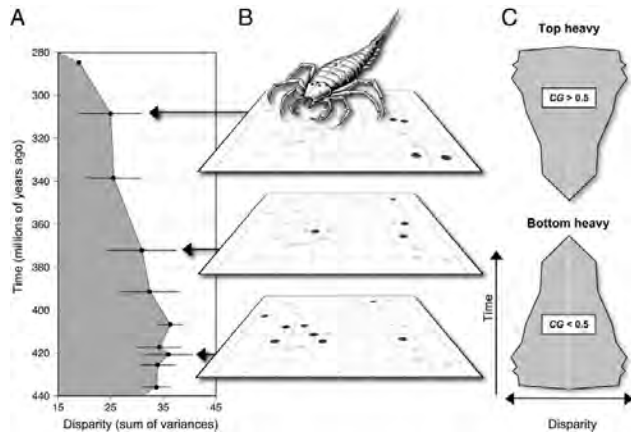


# Disparity Examples

Is there a common pattern of MD accumulation across time or taxa?

- $H_0$ :  $\uparrow$ MD from Cambrian  $\rightarrow$  recent, and early in lineage history

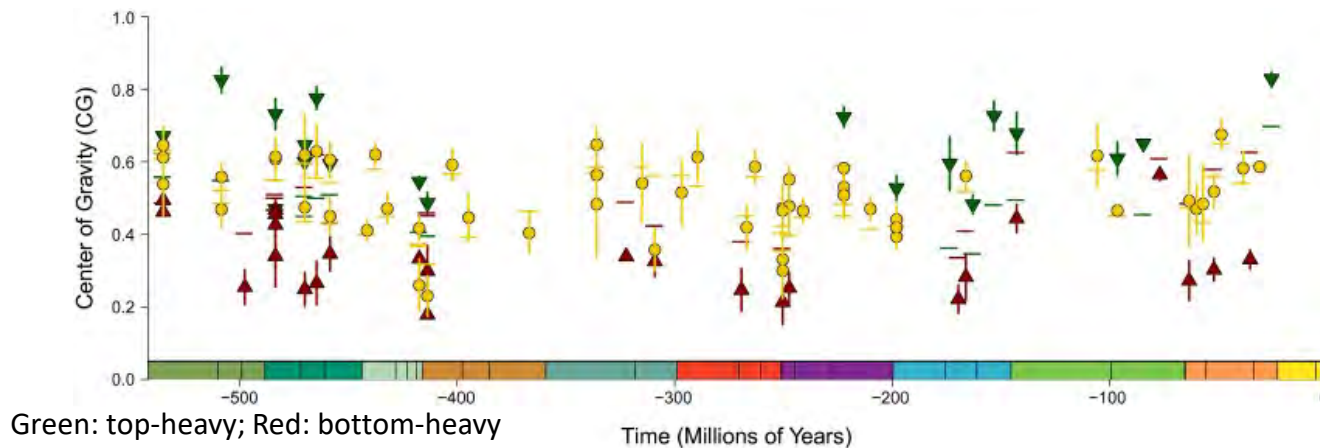
-Analysis of 98 metazoan datasets; most reach peak disparity early



No general trend Cambrian  $\rightarrow$  recent

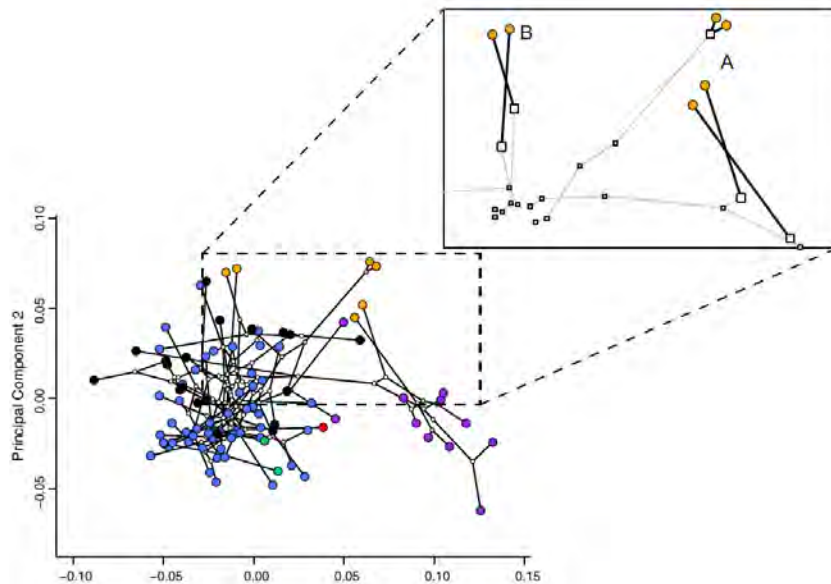
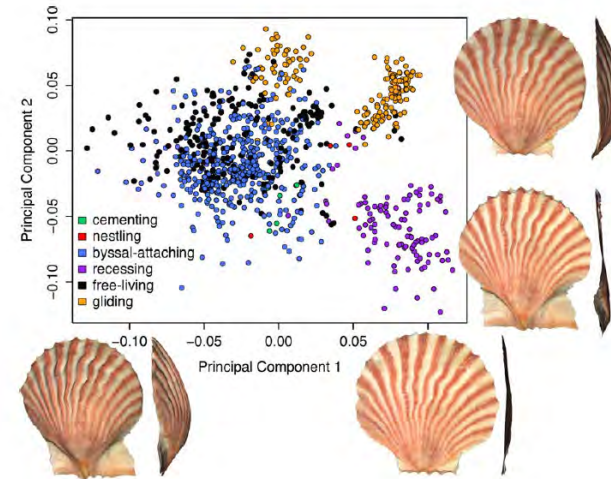
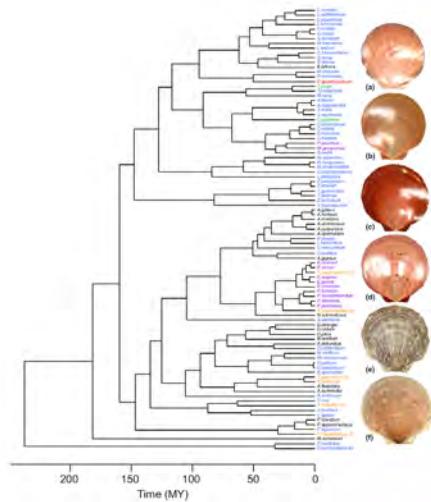
Most clades bottom-heavy

-latter consistent with EB-type radiation models



# Neontological Example

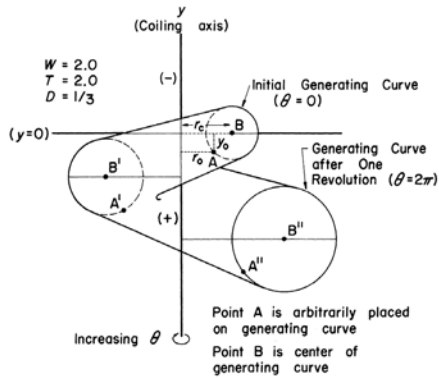
MD differs across scallop life habit eco-groups



-Gliders display less MD  
-Evidence of convergence

# Theoretical Morphospace

## Generate morphospace from mathematical rules



3 variables of coiling shell:

- Expansion whirl
- Distance from axis
- Translation

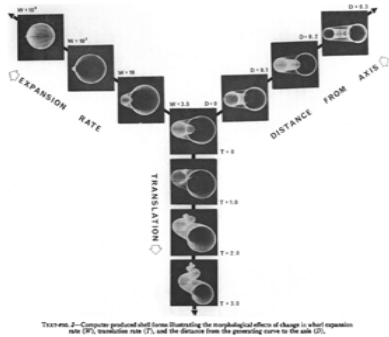
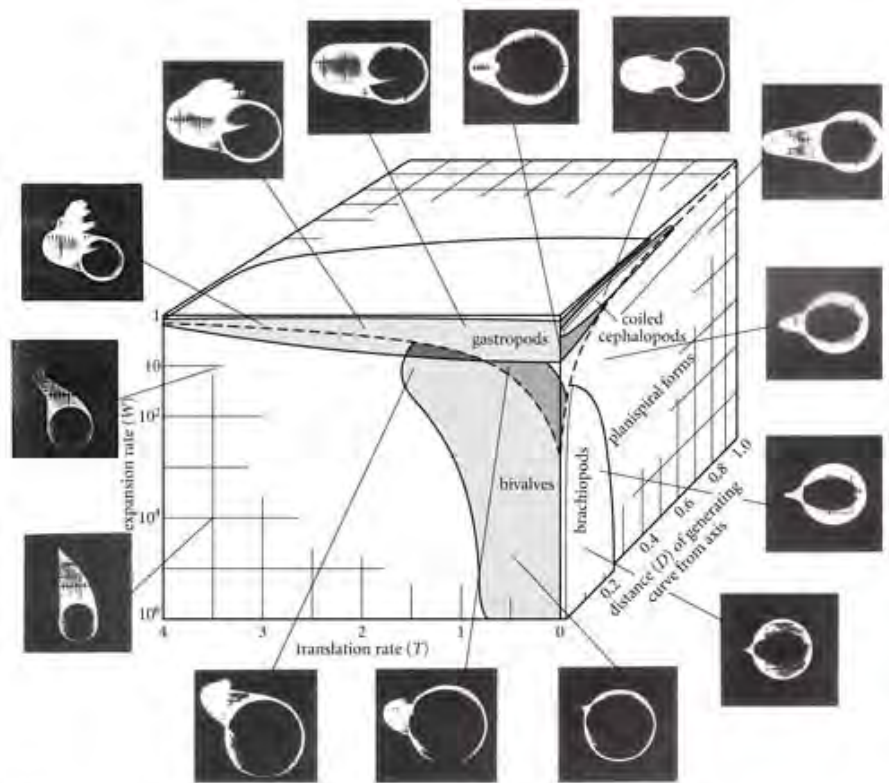


FIGURE 2.—Computer-generated shell forms illustrating the morphological effects of change in whorl expansion rate (W), translation rate (T), and the distance from the generating curve to the axis (D).

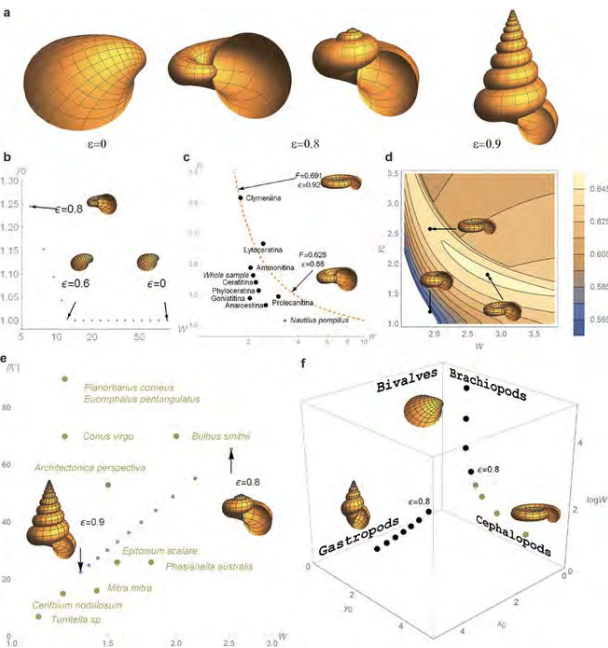
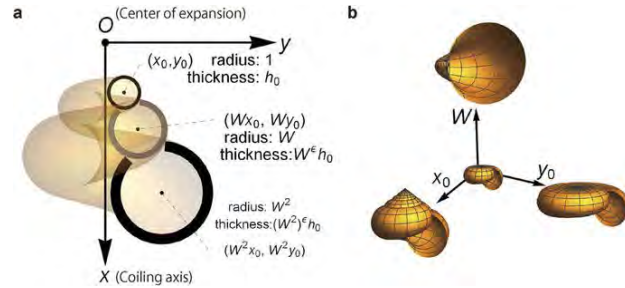


Resulting morphospace with 500 MYA of shell evolution superimposed

## -Helpful to understand structural limits to macroevolutionary change

(Why have certain morphologies not evolved?)

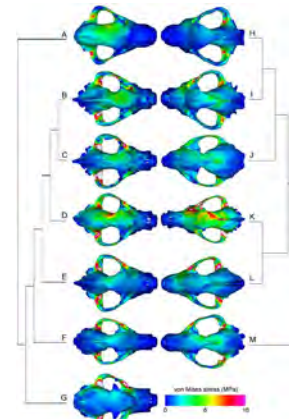
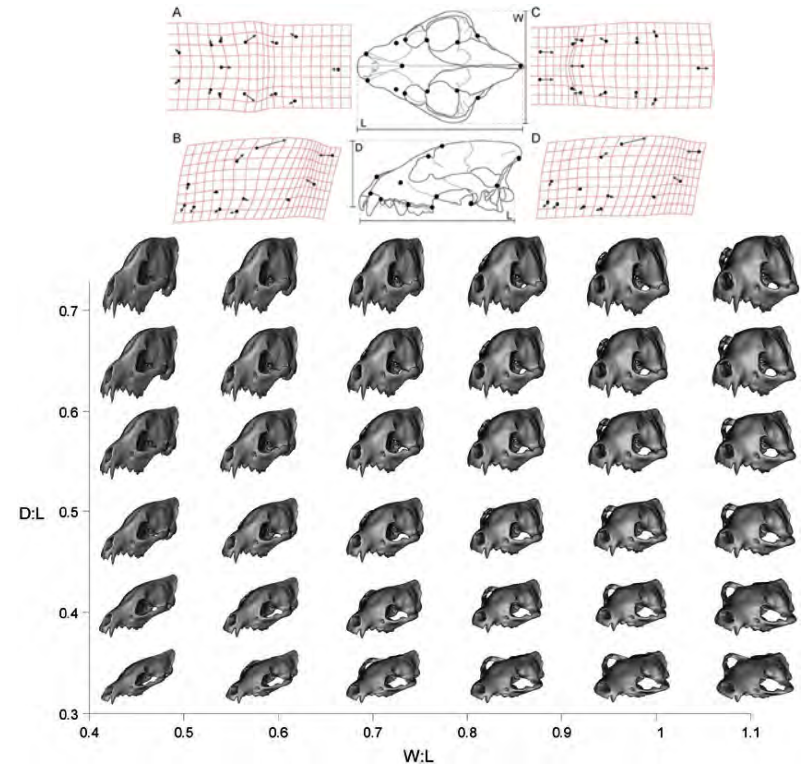
# Theoretical Morphospaces



Bivalves, gastropods, cephalopods

Coiling  $\rightarrow$  morphospace  $\rightarrow$  diversity

Okabe and Yoshimura (2017)



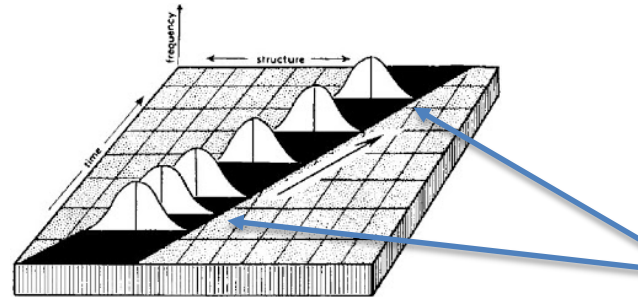
Hypercarnivores and Bite force

Shape  $\rightarrow$  morphospace  $\rightarrow$  function

Tseng (2014)

“How fast, as a matter of fact, do animals evolve in nature?” Simpson (1944)

## Rates of phenotypic evolution



Rate = Trait change / time

Figure 5-2:  
A standard textbook view of evolution *via* phyletic gradualism.  
From Moore, Lalicker, and Fischer, 1952; figure 1-14.

-Darwins:  $r_D = \frac{(\ln Y_1 - \ln Y_2)}{\Delta T}$  common for fossils;  $\Delta T$  typically in MYA

-Haldanes:  $r_H = \frac{(\ln Y_1 / \sigma_{Y_1}^2 - \ln Y_2 / \sigma_{Y_2}^2)}{T_1 - T_2}$  common for extant; t-generations

NOTE: these are lineage-specific (tree-based rates discussed later in semester)

# Evolutionary Rates

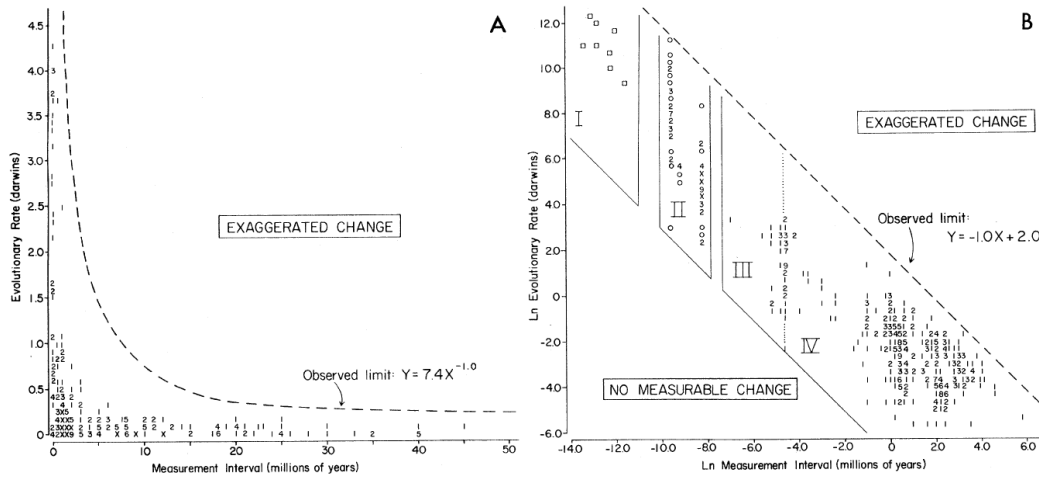
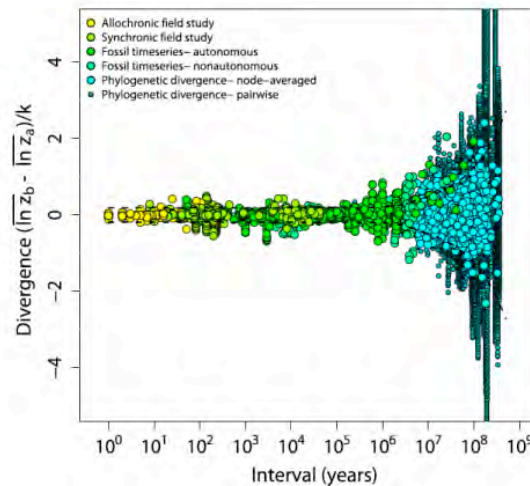


Fig. 1. Inverse relationship of evolutionary rates and interval of time over which rates were measured. (A) Central portion of distribution of 521

Paleontological studies  
Rates are slow  
Gingerich (1983)



Major change ~ 1 MYA  
Uyeda et al. (2011)

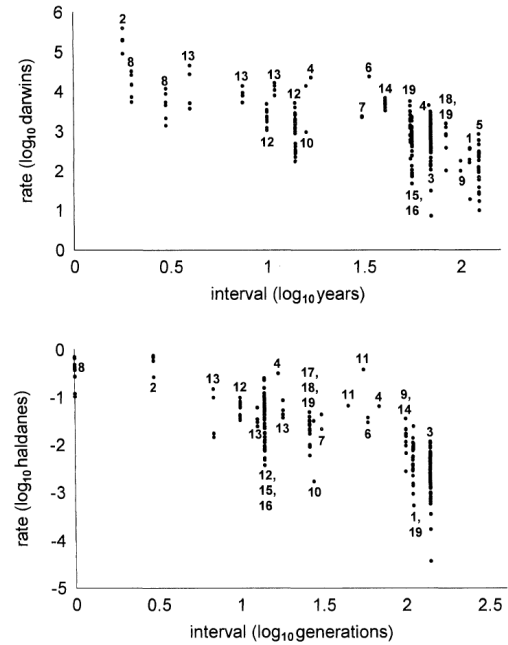
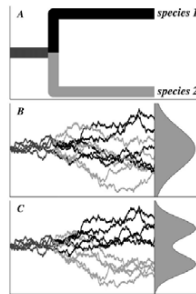
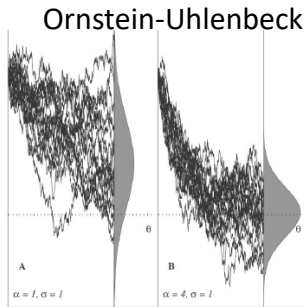
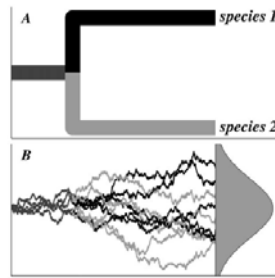
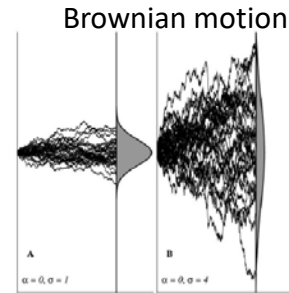


Fig. 4. Distribution of evolutionary rates in darwins (top panel) and haldanes (bottom panel) for the studies summarized in Table

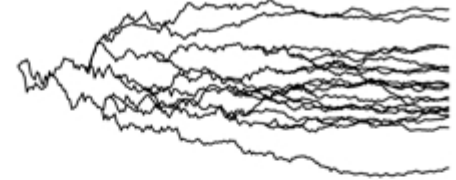
Neontological studies  
Rates are fast  
Hendry & Kinneson (1999)

# Tempo and Mode

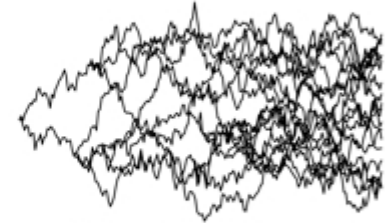
The mode of evolution: the manner in which disparity accumulates



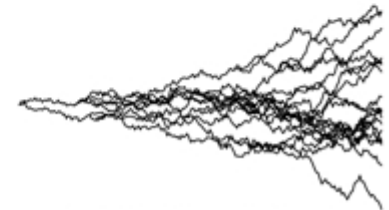
Early Burst



"OU"



Late Burst



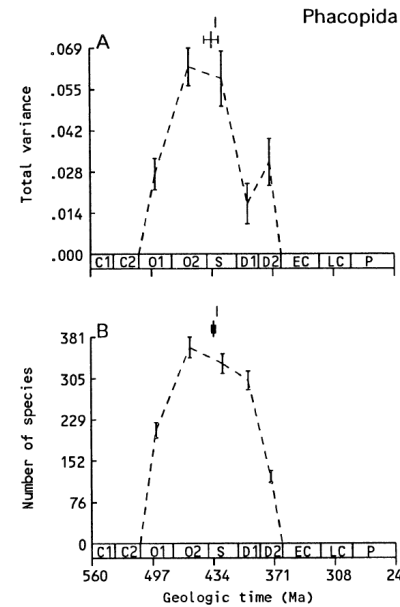
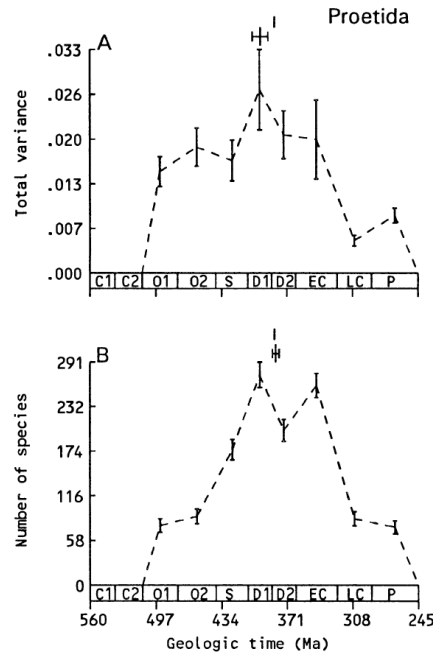
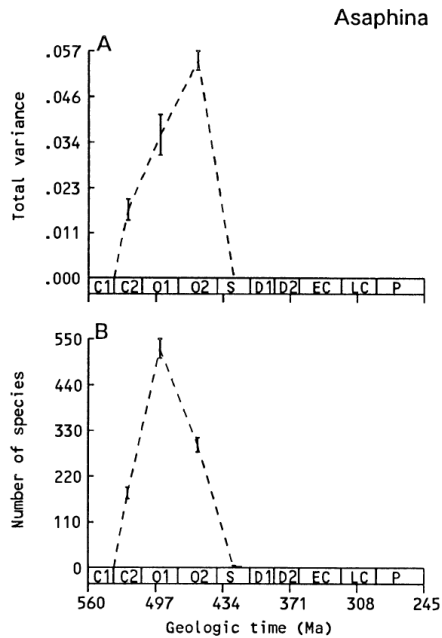
Butler and King (2004)

T. Ingram: [www.anoleannals.org](http://www.anoleannals.org)

-We will discuss this later in the semester

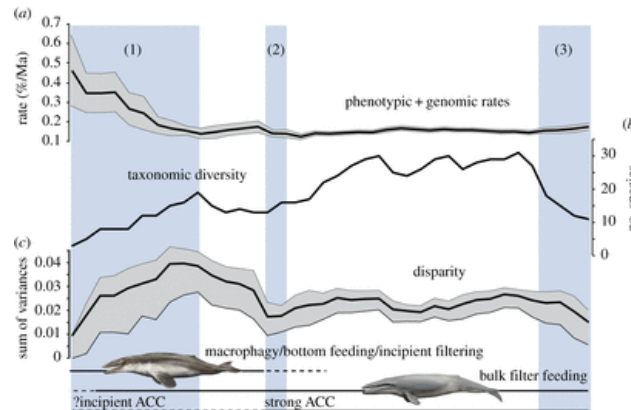
# Combining Diversity and Disparity

## When are diversity and disparity associated?



Diversity & disparity  
Correlated through time

Foote (2015)

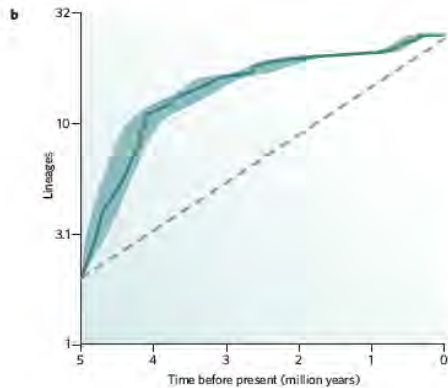
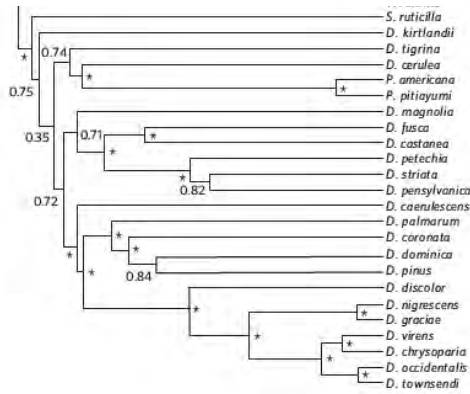


Diversity & disparity  
Decoupled through time

Marx & Fordyce (2015)

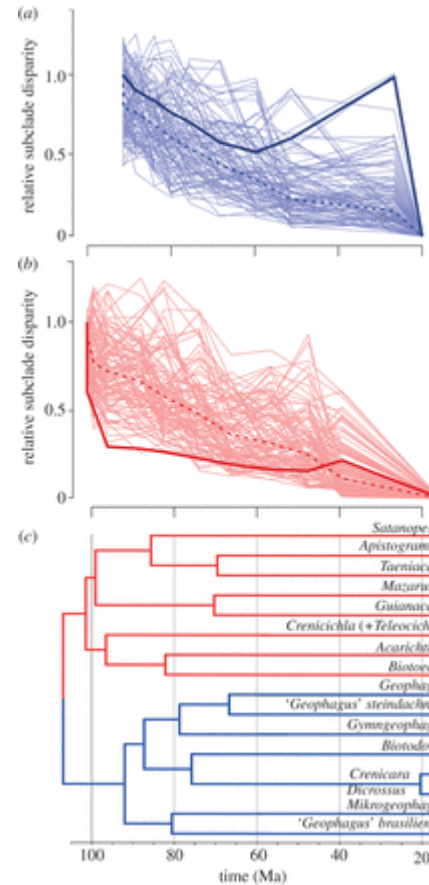


How do taxonomic & phenotypic diversity accumulate over time?



Reznick and Ricklefs (2009)

LTT plot



Arbour and López-Fernández (2013)

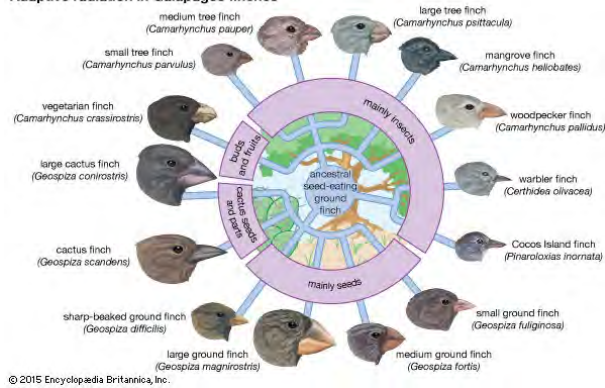
DTT plot

\*NOTE: convention for DTT is opposite LTT, because measured as MD BETWEEN sub-clades which must decrease over time (began with Harmon et al. 2003)

# Adaptive Radiations

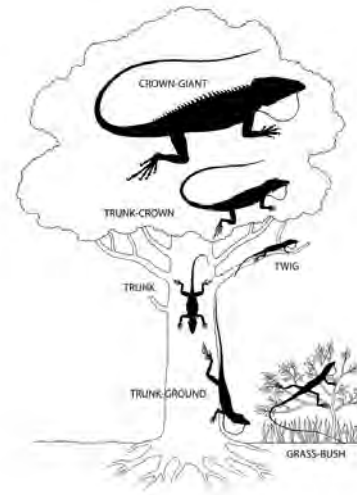
Phenotypic and taxonomic diversification; exploiting new niches

Adaptive radiation in Galapagos finches

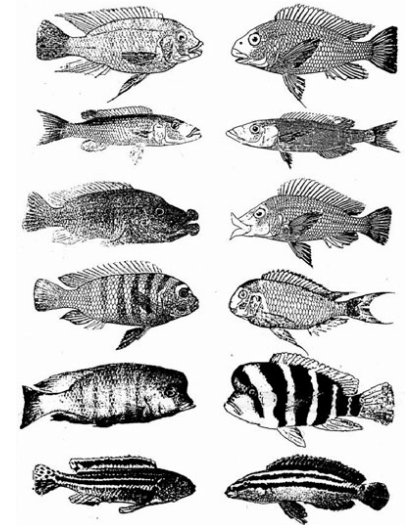


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Encyclopaedia Britannica, Inc (2010)



Losos (2009)



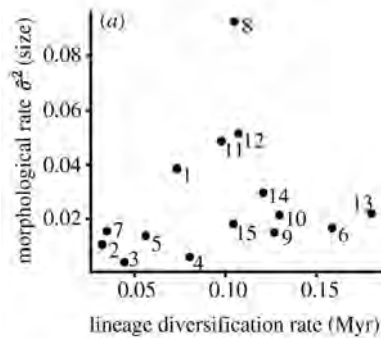
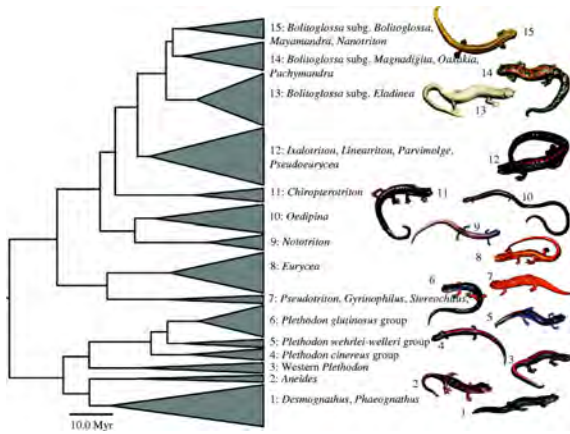
Kocher et al. (1993)

Predictions from AR hypothesis:

- Disparity follows early-burst (EB)
- Diversity: LTT plot shows EB
- Diversity and disparity rates expected to be coupled

# Adaptive Radiations

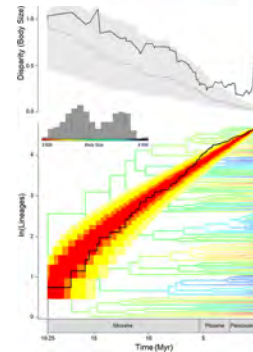
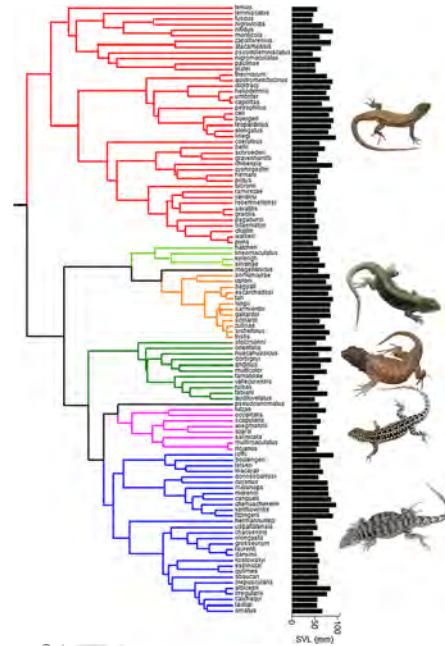
Sometimes, it's just a radiation...



Plethodontids

-Rates not correlated

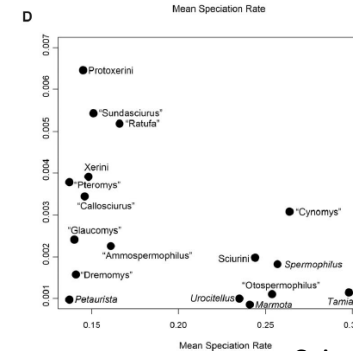
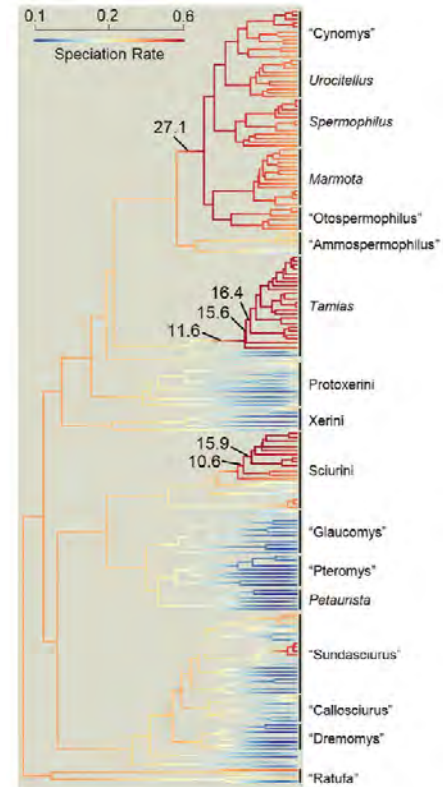
Adams et al. (2009)



S. Amer. *Liolaemus* lizards

-LTT & DTT: not EB

Pincheira-Donoso et al. (2015)



Sciuridae rodents

-Rates not correlated

Zelditch et al. (2015)

