

# Evolution of Body Size in Bears



**How to Build and Use a Phylogeny**

# Lesson II

## Using a Phylogeny

## Objectives for Lesson II:

### 1. Overview of concepts

#### 1. Simple ancestral state reconstruction on the bear phylogeny

- class activity

#### 1. Ancestral state reconstruction of polyploidy in birches.

- In pairs

Students will use the following example:

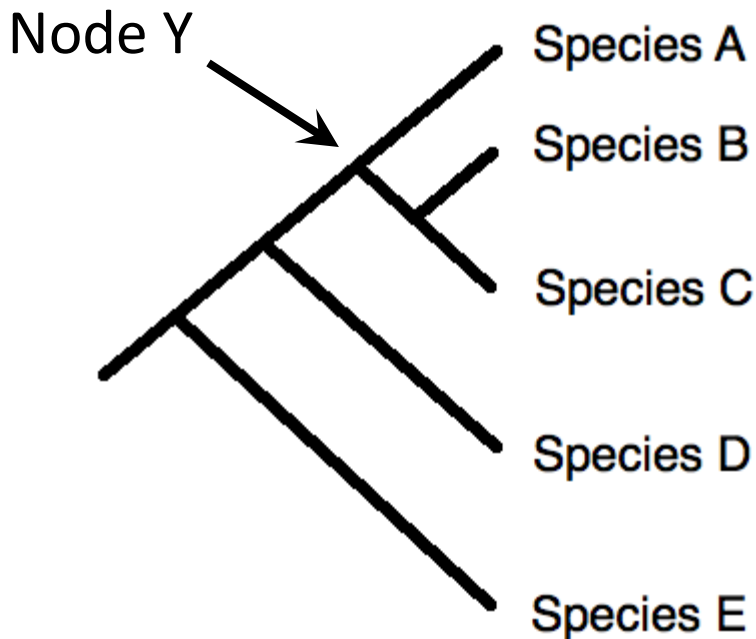
- Examining frequency of polyploidy in *Betula* (birches), based on Mayrose *et al.* (2011) *Science*

# Character Evolution

- Character evolution is the study of how a trait like color has evolved across a phylogenetic tree
- Character states impact an organism's ecology - where it lives, what it eats, how often it reproduces etc...
- Understanding the evolution of a trait through time reveals the origin of current biodiversity.

# Terms Review:

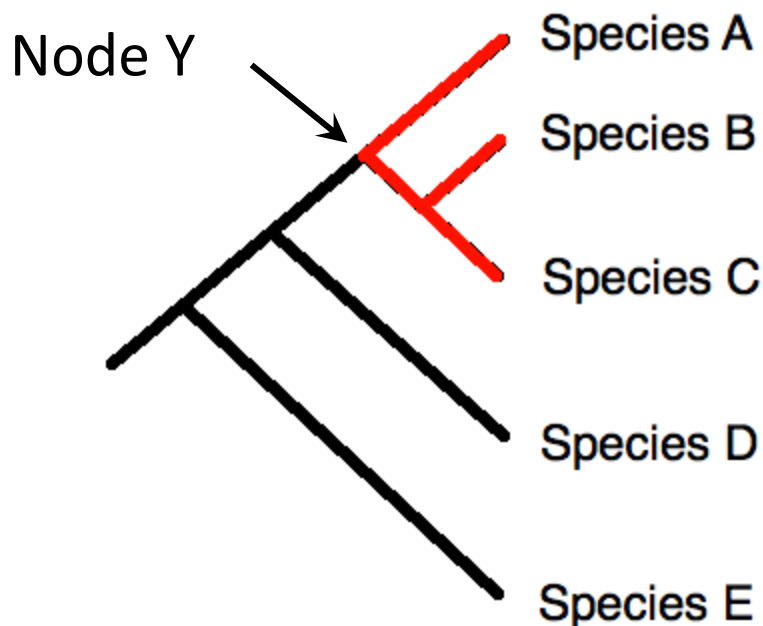
- A *phylogeny* describes the evolutionary relationships of a set of organisms. It is often based on morphological traits or, more commonly, DNA sequences.



- Species B and C are *sister* taxa
- Species E is an *outgroup* to ABCD
- Node Y is the *ancestor* to A, B & C

# Terms cont...

- *Ancestral state reconstruction* is a method to estimate the character state of an ancestor based on phylogenetic relationships and character states of present-day organisms

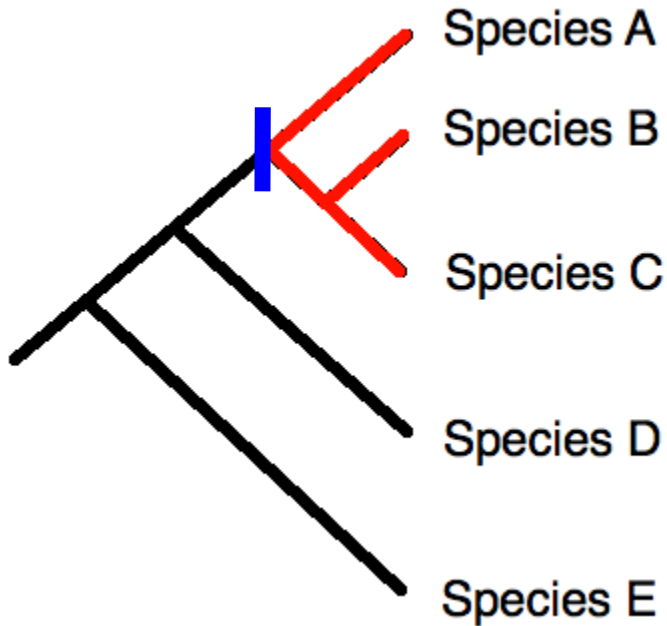


- In this example, the *character* is color & the *character state* is red or black
- Species A, B & C share the same character state (red). In this reconstruction, the character state red evolved in the ancestor at Node Y

# Ancestral state reconstruction

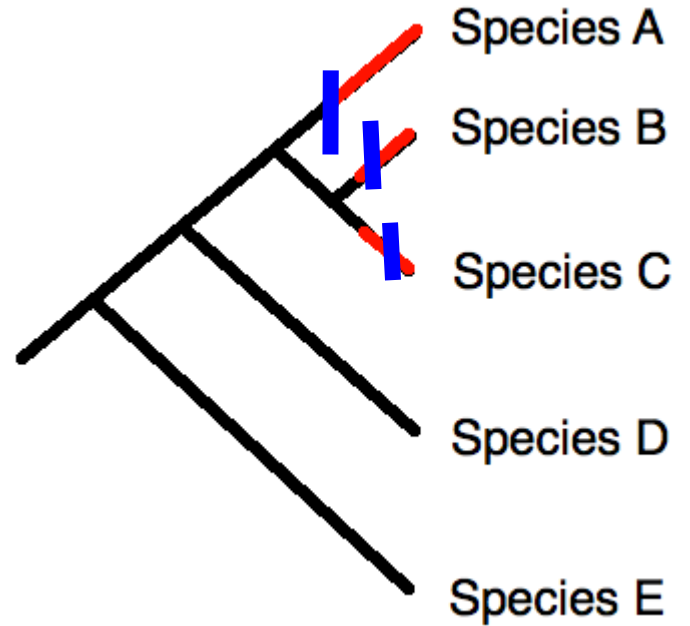
- There are multiple methods to reconstruct ancestral states.
- *Parsimony* is one of the simplest methods.
- Parsimony attempts to **minimize** the number of character state changes across the phylogeny

█ - Character state change



- One character state change

Most parsimonious



- Three character state changes



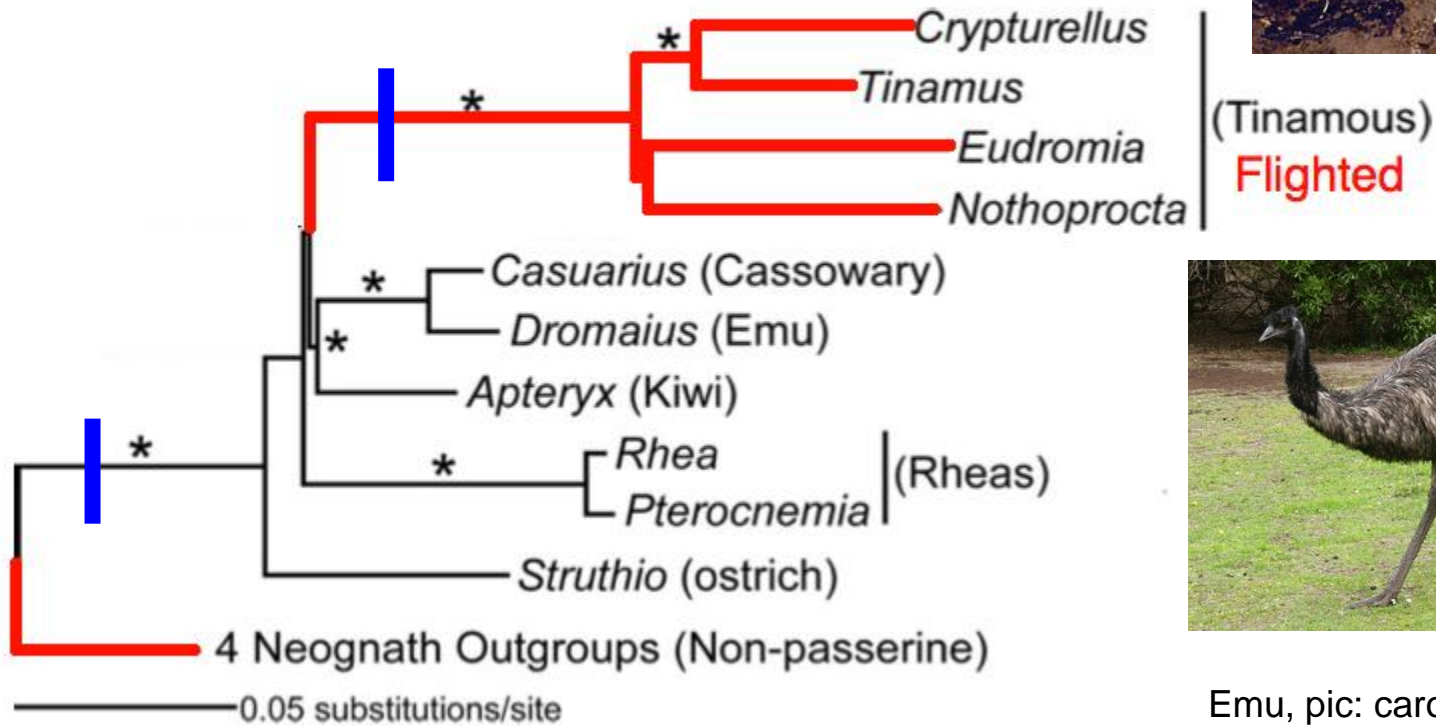
# Ancestral state reconstruction

- All methods of ancestral state reconstruction have assumptions
- *Parsimony* assumes that characters have evolved in a way that involves the fewest possible changes.
- There are examples when parsimony reconstruction is NOT the most likely

Great tinamou, pic: Patrick Coin (1992)



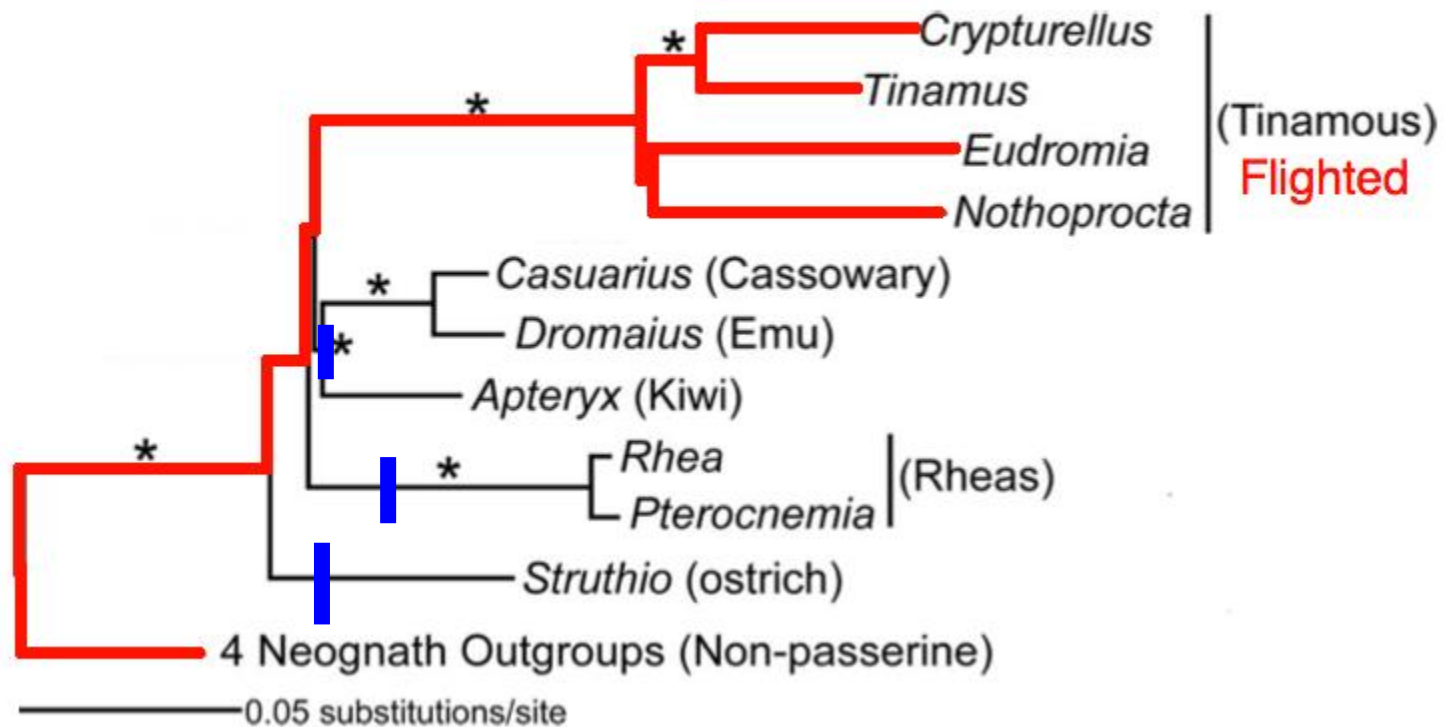
# Ratite Birds



Emu, pic: carolinaemu.com

Modified from Harshman et al (2008) *PNAS* 105:13462-13467

# Non-parsimonious reconstruction - most likely



# Ancestral state reconstruction

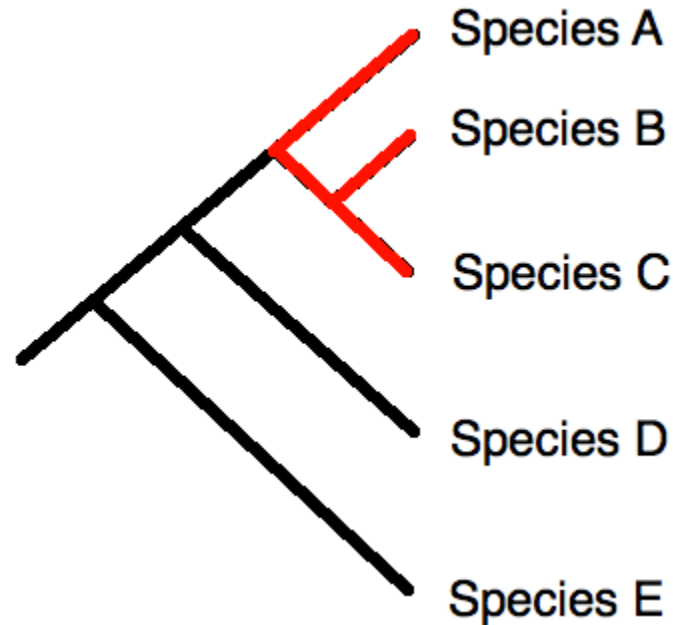
- The data used to generate a phylogeny and the character mapped onto the tree must be *independent*.
- You cannot use ancestral state reconstruction for a character you used to build the tree (circular).

# Uses of ancestral state reconstruction

- Indicate character state of ancestor
- Does a trait change often or rarely across the tree? Is a trait *labile* (changes frequently/easily) or *conserved* (changes infrequently)?
- Is the state change associated with an increase or decrease in diversification rates?

# Examples...

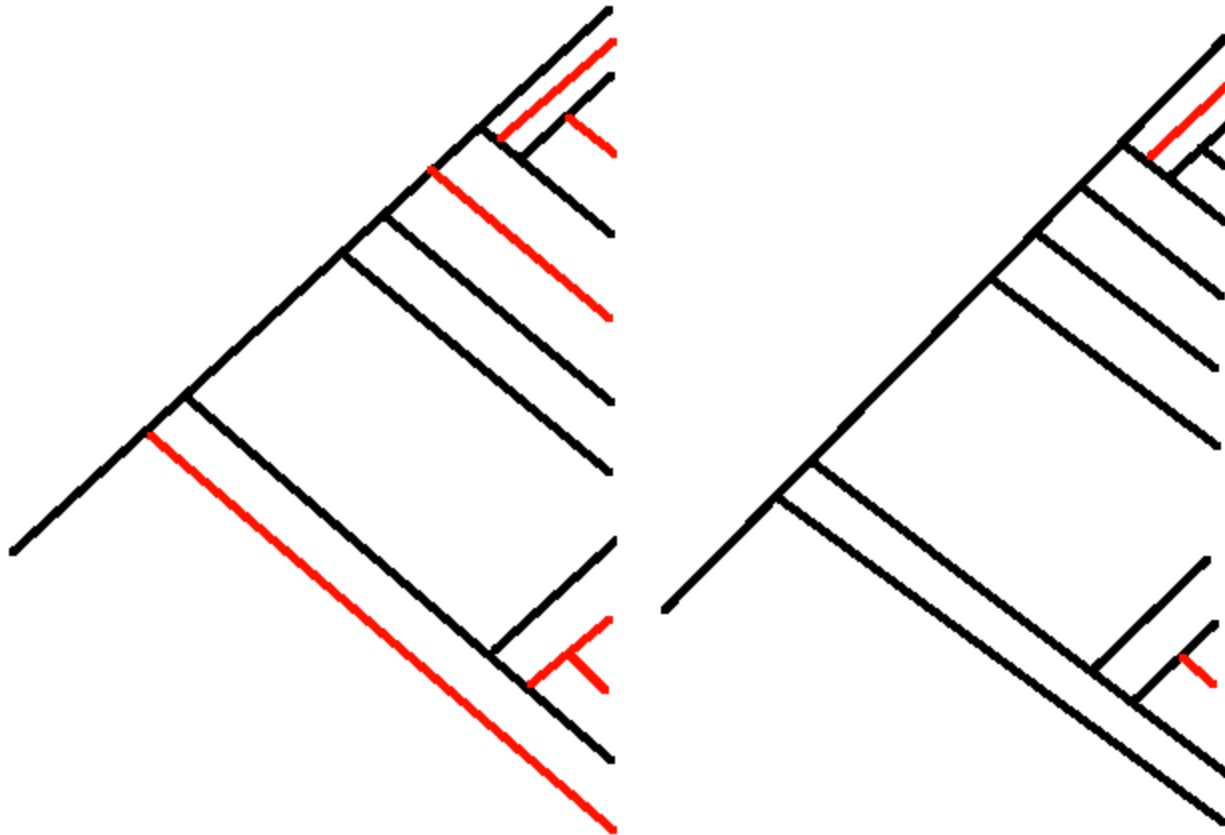
What is the character state of the ancestor?



Eg. When & how did migration originate in Wood-warblers?  
(Winger et al. 2012. *Proc.Roy.Soc.B.* **279**: 610-618)

# Examples...

Is a character labile or conservative?

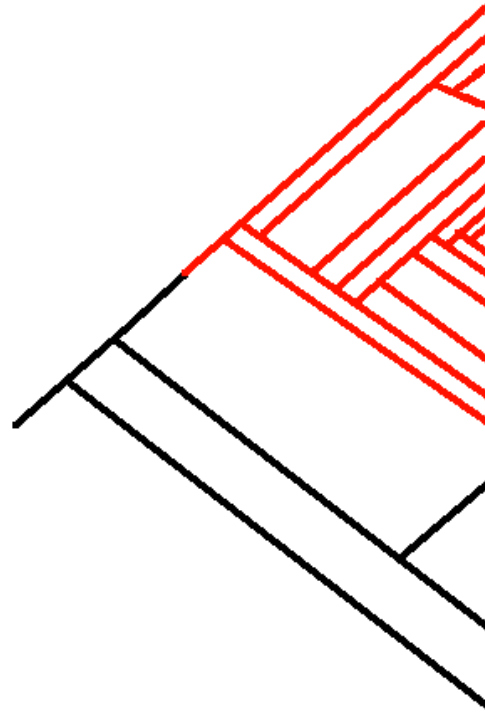


Eg. Are behavioral traits more or less conserved than morphological and physiological characters?

(Blomberg et al. 2003. *Sys.Bio.* **57**: 717-745 )

# Examples...

Is a character state change associated with a change in diversification rate?



Eg. Are key innovations in angiosperms associated with an increased rate of diversification?

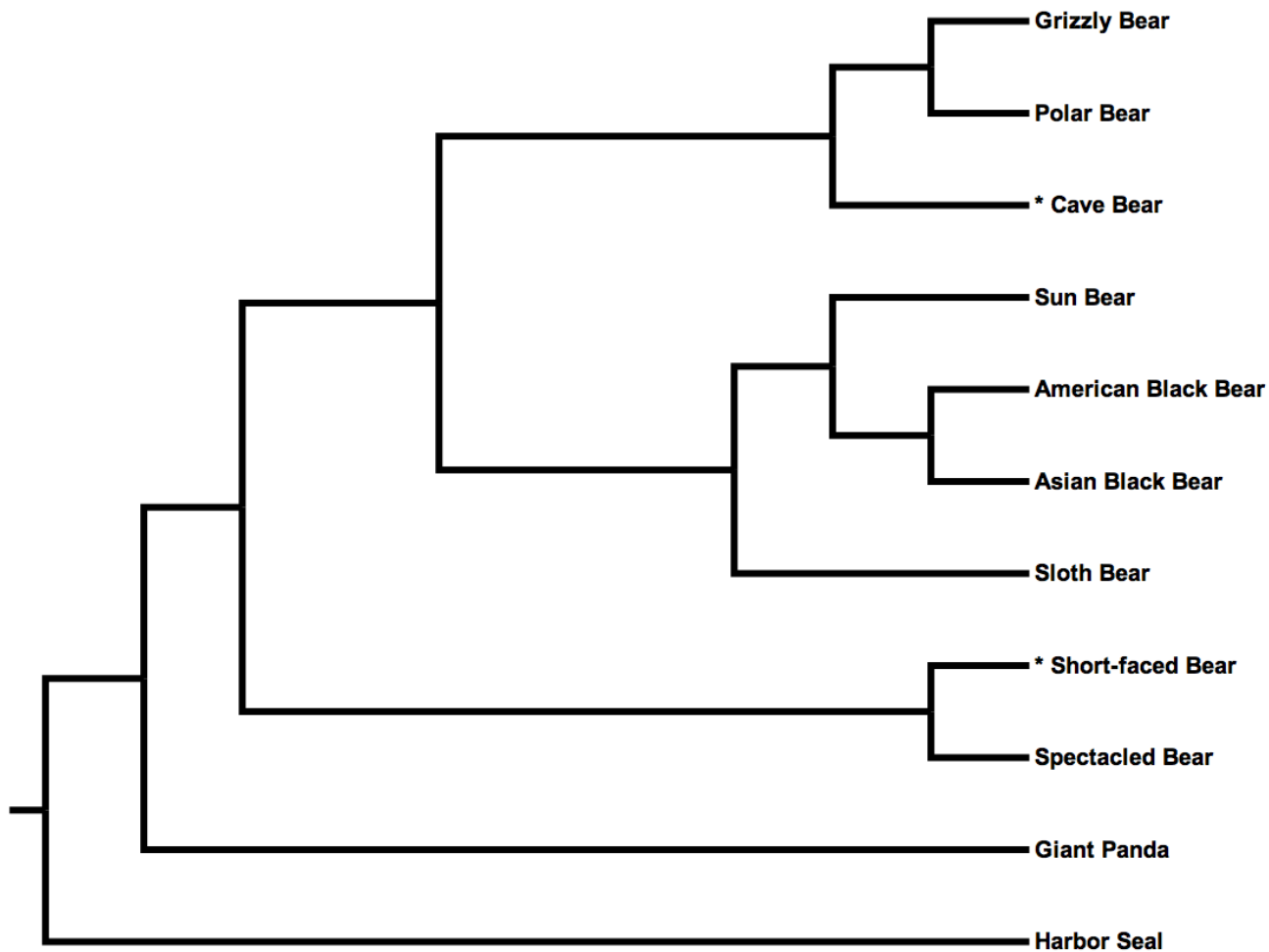
(Sanderson & Donoghue. 1994. *Science*. **264**: 1590-1593)



# Biology of Bears

- 8 living species (and many extinct species)
- Widespread (mostly Northern Hemisphere and parts of Southern Hemisphere)
- Carnivores
- Closest relatives are dogs (canids) and seals






1.1

\* = extinct species

## Body Mass Database:



**Felisa A. Smith**  
Department of Biology, University of New Mexico

Home | Research | Members | Publications | **Datasets** | Join Us | Popular Writing | PIBBs | In the News

### 1. MOM v4.1.

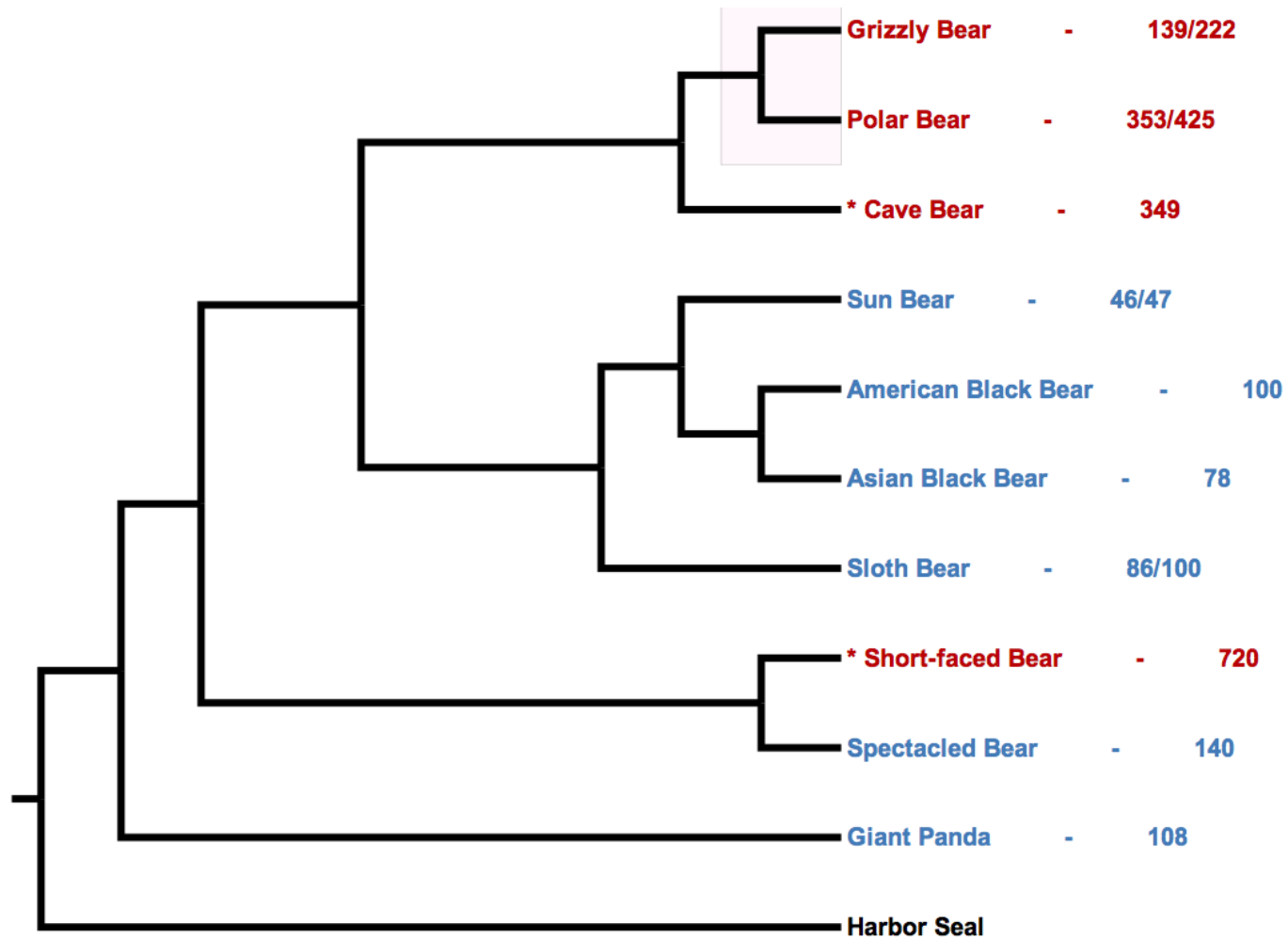
Global database of late Quaternary mammal body mass arranged by continent.

<http://biology.unm.edu/fasmith/Datasets/>

- Body mass is a *continuous* trait. For simplicity, we are going to lump bears into two groups: **large** and **small**.

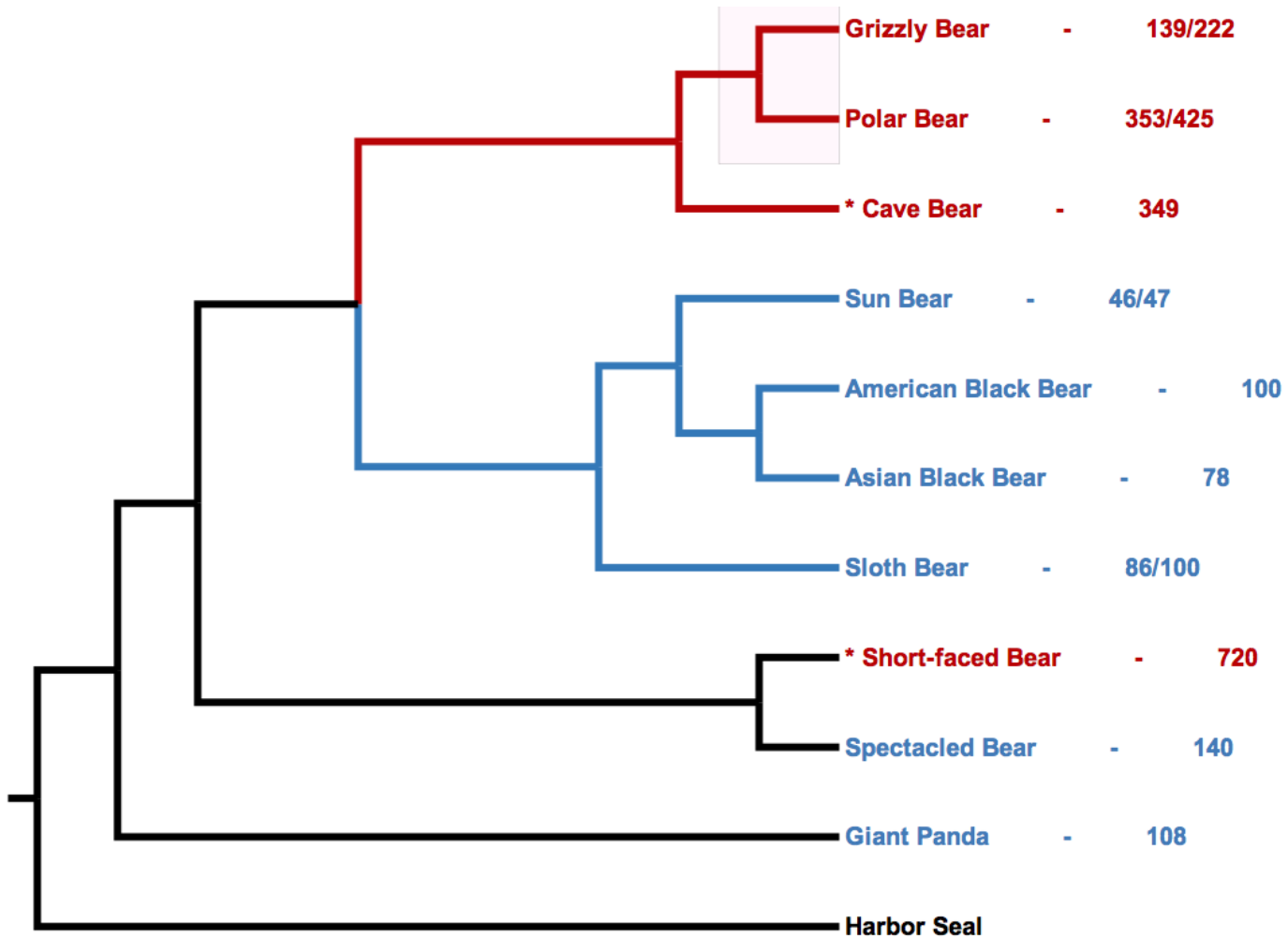
- **large** bears > 200 lbs
- **small** bears < 200 lbs

- We will then conduct a simple ancestral state reconstruction of body mass in bears.

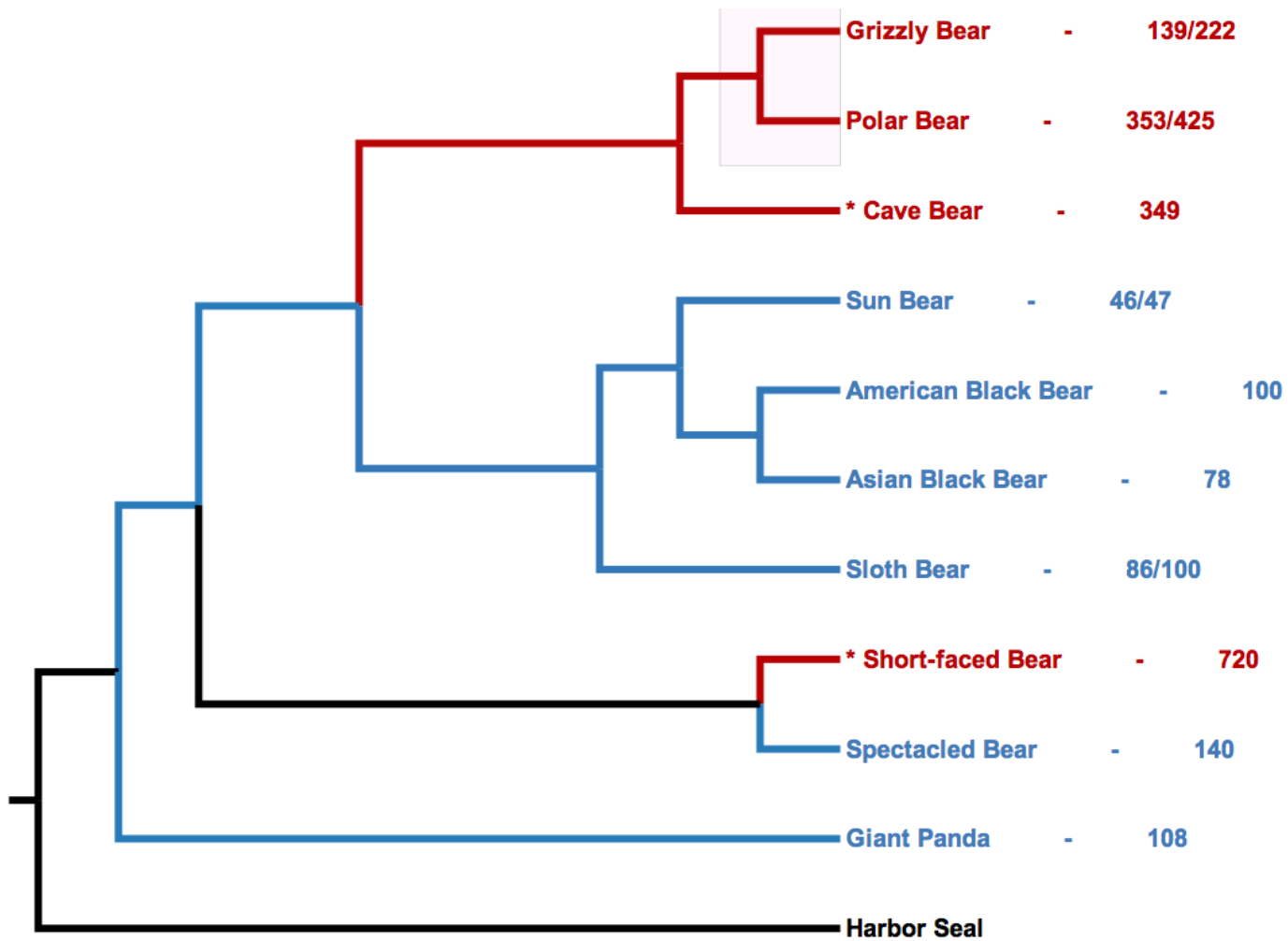


1.1

*\*\* Average masses (male + female) are shown. Masses for extinct species are estimates based on the fossil record. See MOM metadata for more information.*



1.1



- Based on our parsimony approach, the ancestor of all bears appears to have been a **small**-bodied bear (that is, <200kg).
- Now form a hypothesis for why some bears have evolved larger body sizes.
- Any ideas??



Evolution of large body size could be due to:

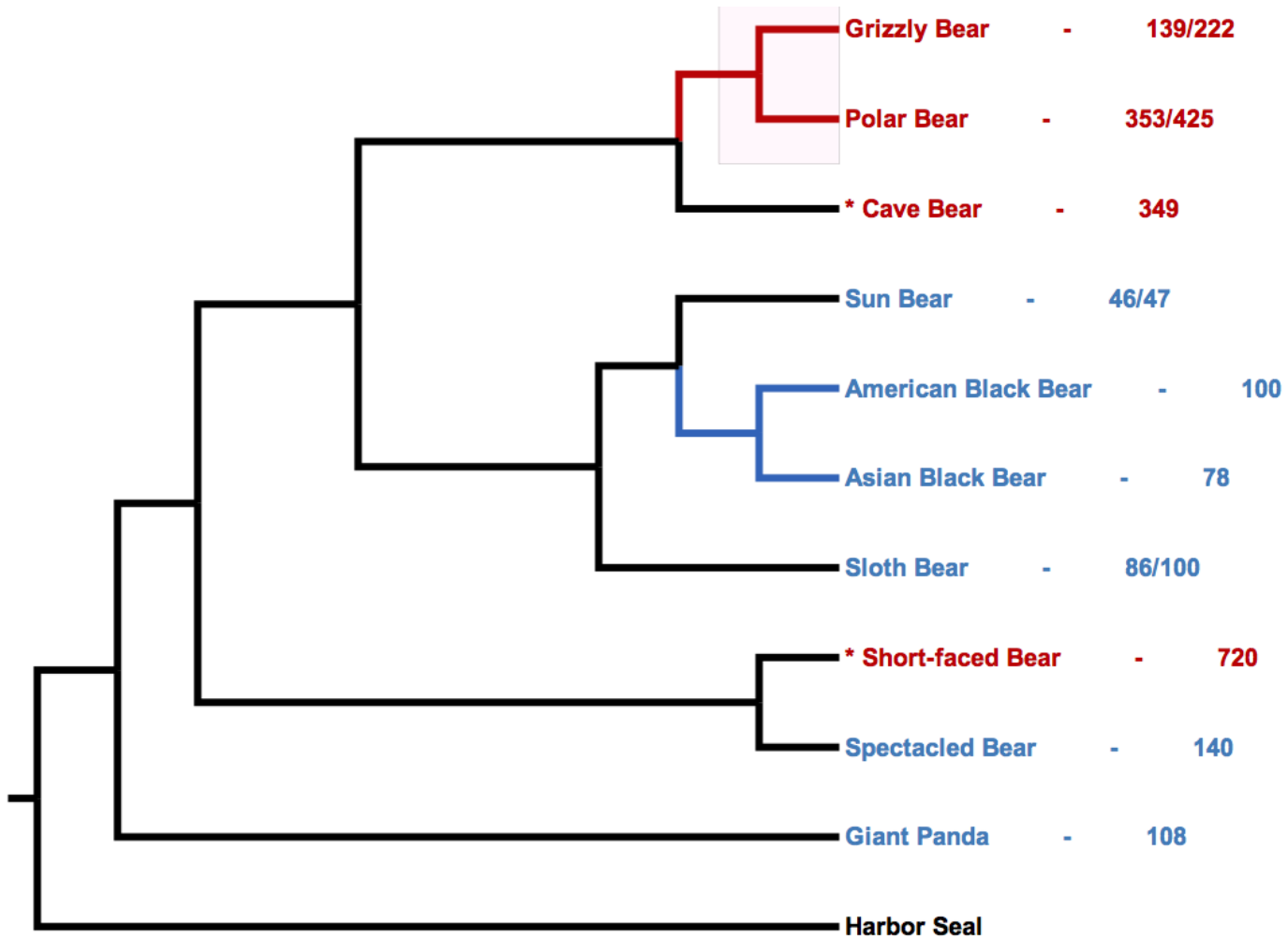
- adaptation to high-latitude environments
- adaptation to ice-age climates  
(larger size aids in retention of body heat)

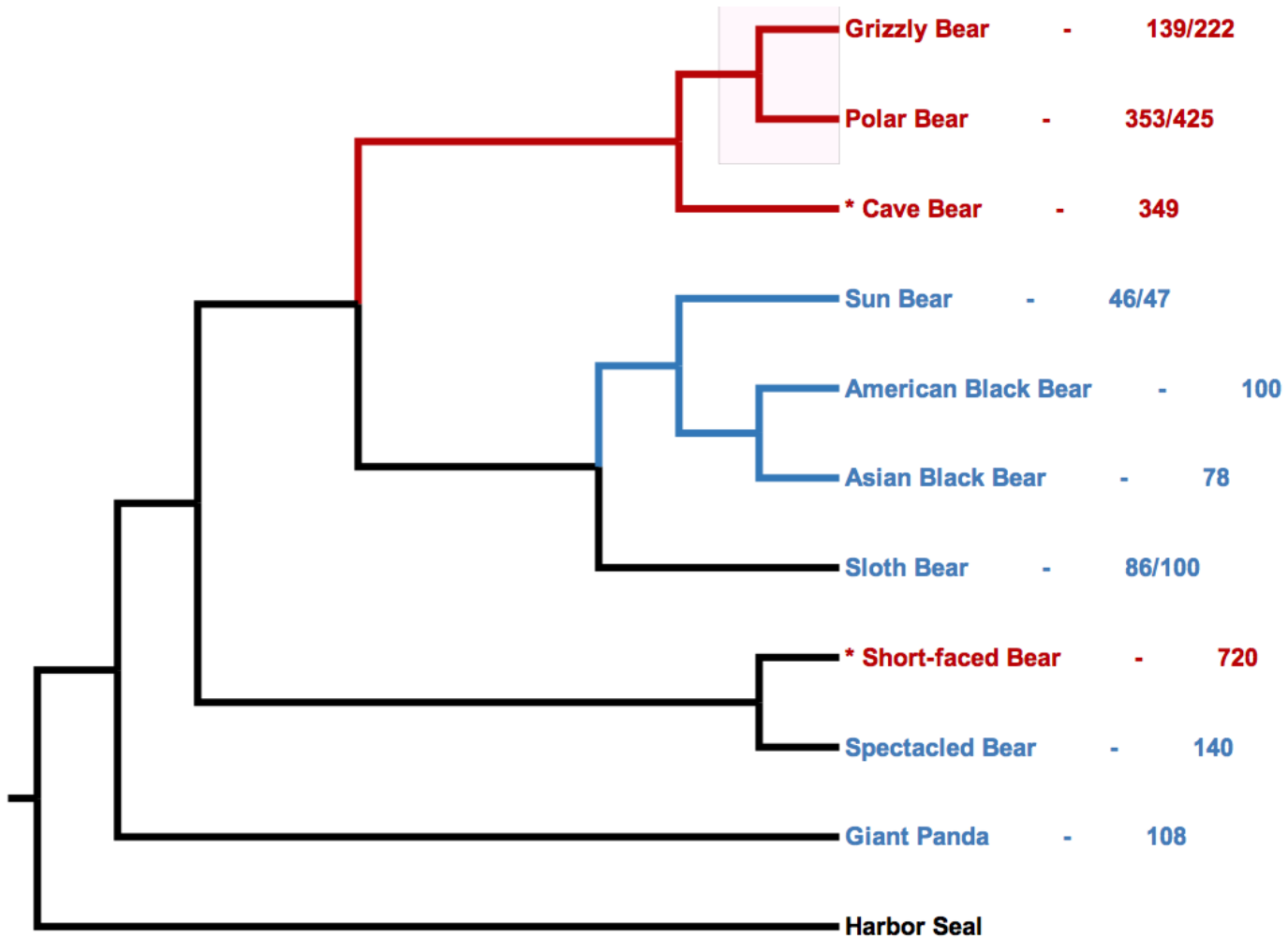
OR

- adaptation to larger prey sizes

The End







1.1

