


species = groups of actually or potentially interbreeding natural populations

population = individuals of the same species living in some defined area.

Evolution = (in a biological context) a change in the genetic content of a population with time. If the time period for the change is short, the genetic change is small and a new species does not result (microevolution). However, if the time period for the change is long, the genetic change is large and a new species will result (macroevolution).




EXACTLY HOW *WINGS* EVOLVED IS SOMETHING OF A MYSTERY.

THE PROBLEM IS THAT *PARTLY* EVOLVED WINGS ARE USELESS FOR FLYING!



SINCE TRYING TO FLY WOULD BE FATAL, IT'S HARD TO SEE HOW A HALF-WINGED ANIMAL COULD LIVE TO REPRODUCE. AND YET...

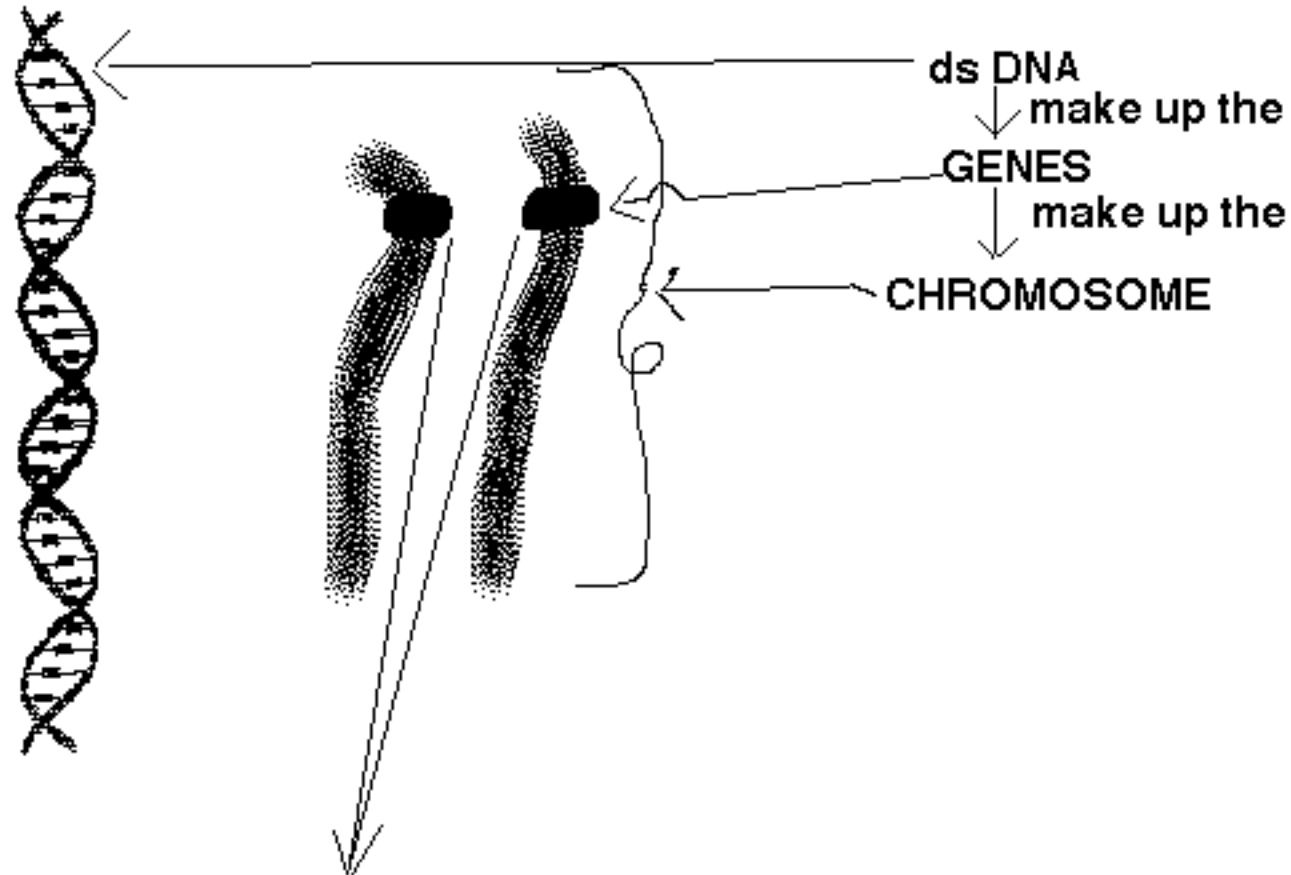


THIS IS BATTY!

THE ANSWER SEEMS TO BE THAT HALF A WING HAS SOME *OTHER* FUNCTION, LIKE SNAGGING BUGS, AND ONLY LATER IS USED FOR FLIGHT.



WOW! LOOK HOW BERNADETTE IS USING HER BUG-SNAGGERS!



ALLELES: Two different forms of the gene

For many hereditary traits, genes exist in two or more different forms called alleles. On each pair of chromosomes, there is one allele for a particular gene on each. Ex. A, B, O blood groups. In humans there are 3 alleles: A, B, and O. ↩

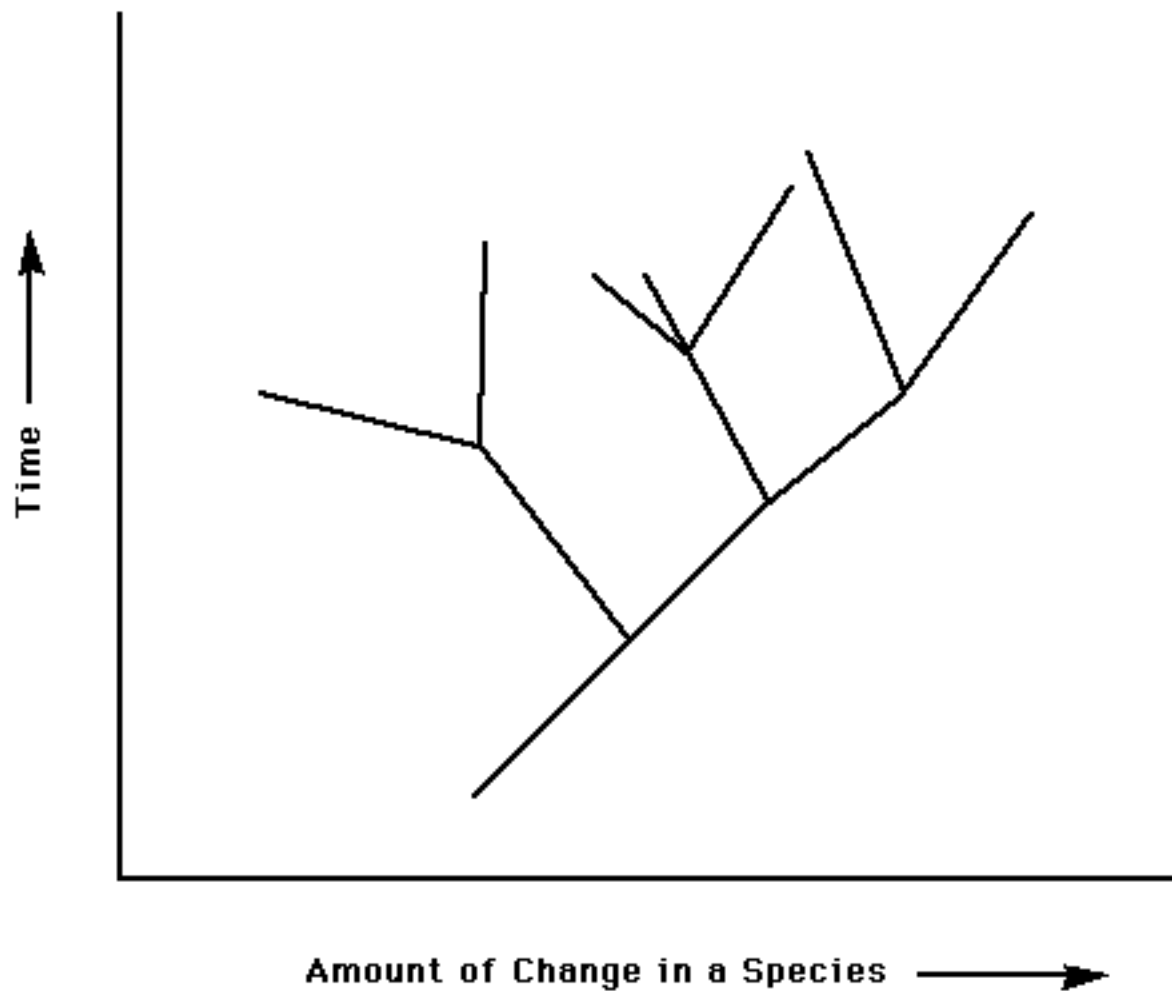
1. Factors that produce new alleles

Mutation

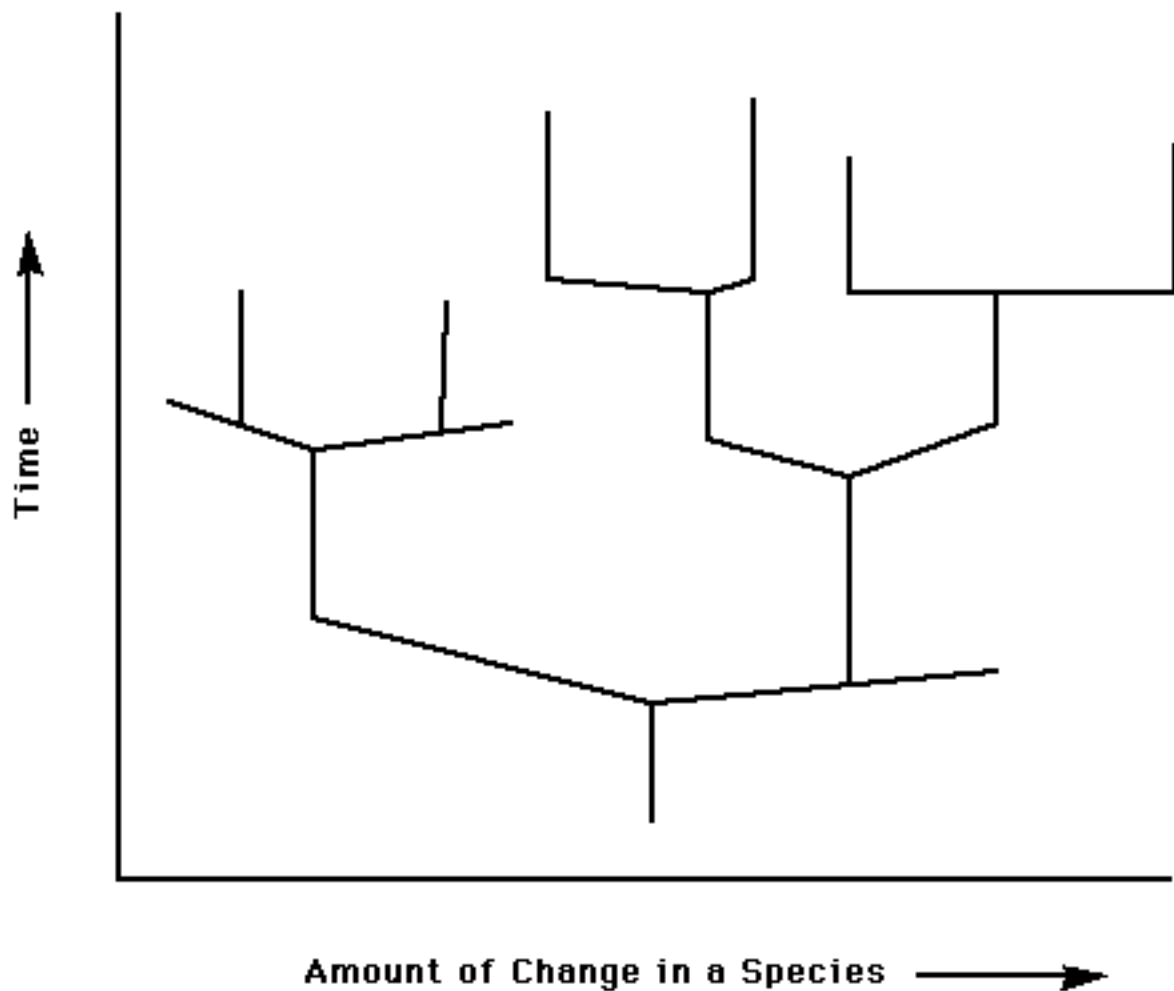
2. Factors which just reshuffle already existing alleles

- a) recombination – alleles from each parent**
- b) genetic drift – Separation of small populations**
- c) migrations – into or out of a population**
- d) selection pressures**
 - 1. Natural selection**
 - 2. Artificial selection**

Gradualism



Punctuated Equilibria



There are two approaches to tracing the family tree of any organism:

1. Examine the fossil record – Paleontology

Archaeopteryx Fossil

Archaeopteryx Picture

2. Examine the genes of living organisms – Molecular genetics

Archaeopteryx

Camel Evolution



ARCHAEOPTERYX

150 MILLION OLD
FOSSIL

Molecular Genetics



The sequence of monomers in DNA, RNA or a protein can be used to estimate the degree of "relatedness" between organisms. As an example, consider the partial proteins below in which each letter represents an amino acid.

Which pair is the most closely related ?

SYKTSDAKGS

1

FAKPVSICAE

2

VAKHVSICAE

3

VAKLYQIHAE

4

SYKTSDAKGS

1

FAKPVSICAE

2

VAKHVSICAE

3

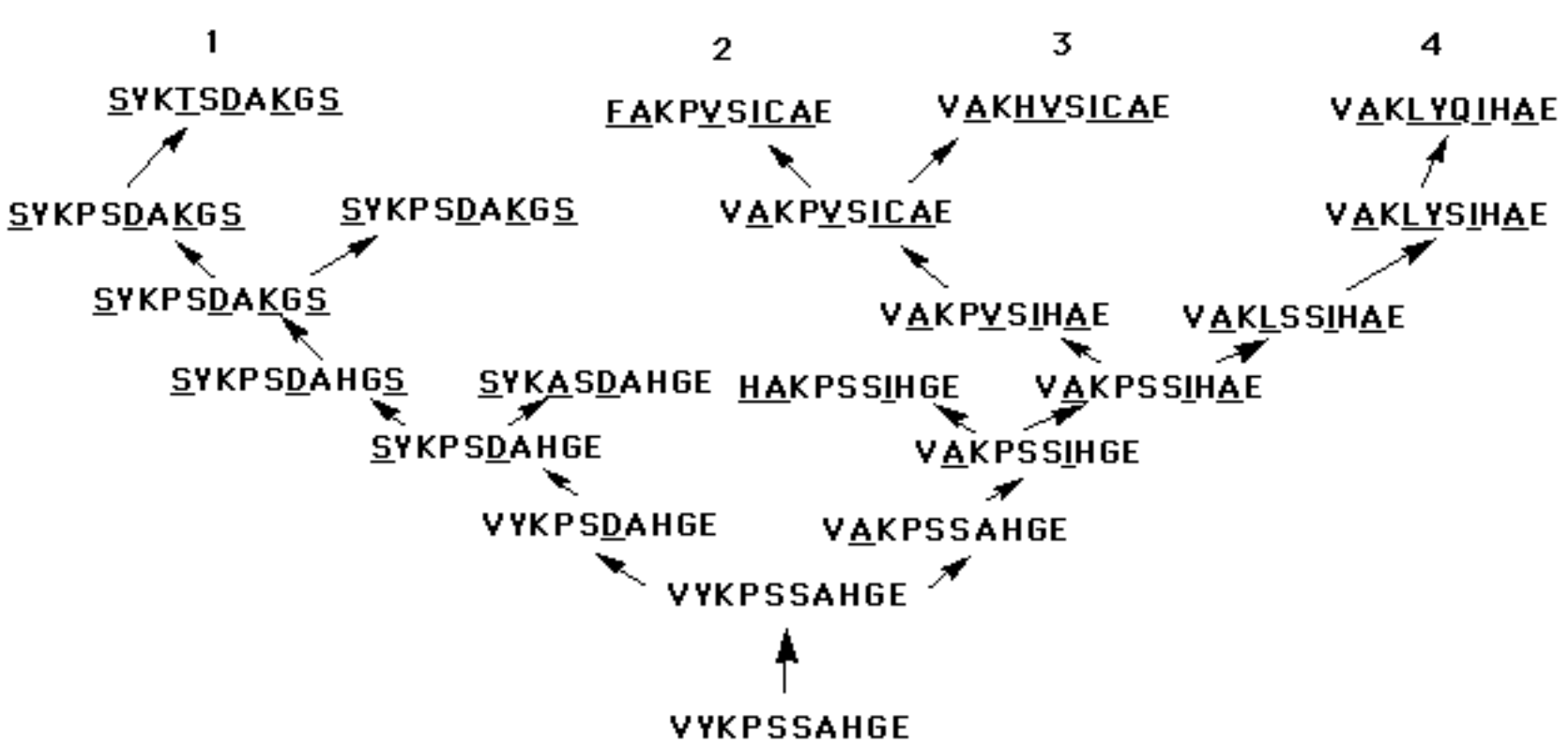
VAKLYQIHAE

4

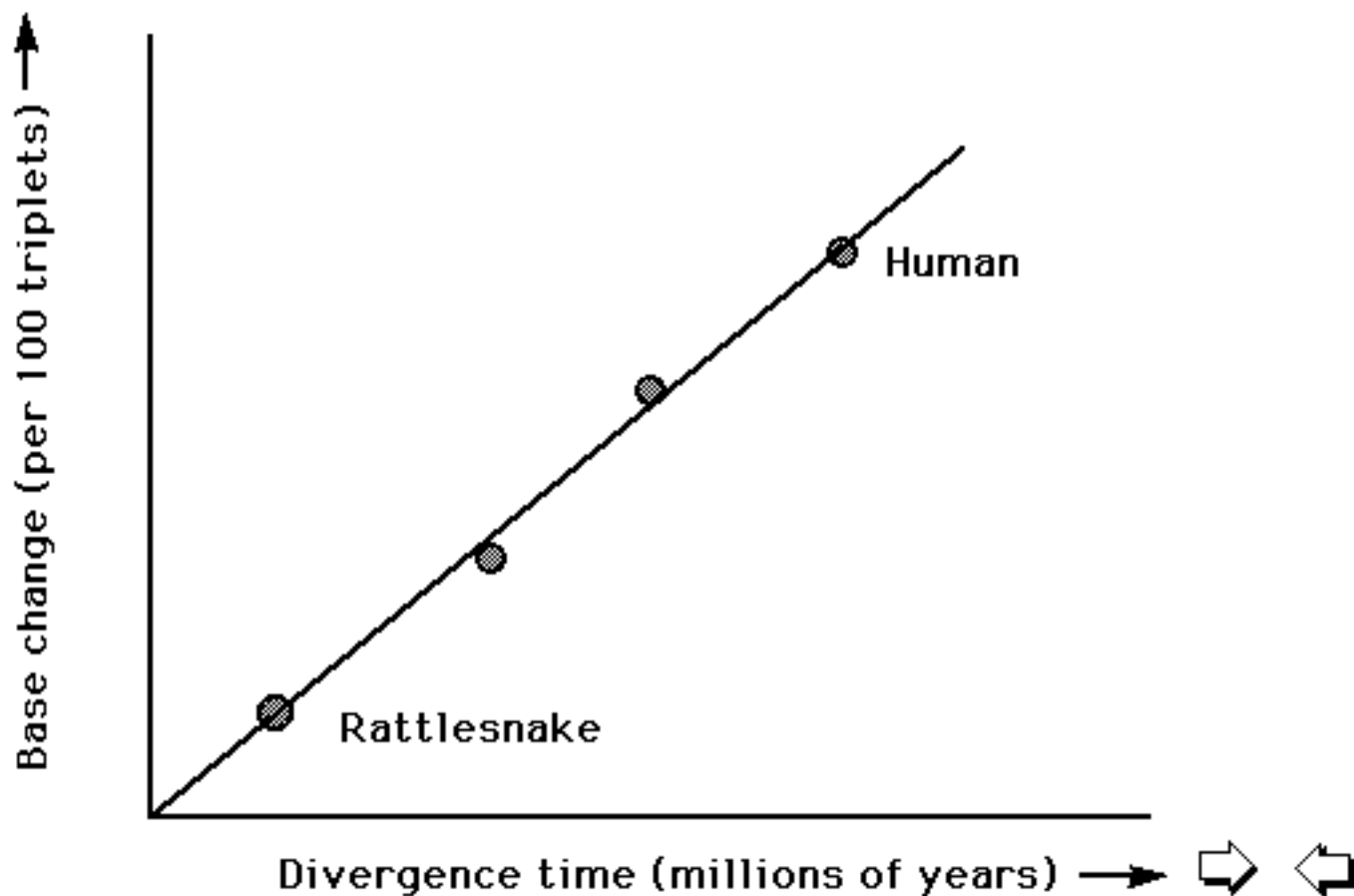
Compare their sequences:

1	SYKTSDAKGS	}	9 different
2	FAKPVSICAE		
3	VAKHVSICAE	}	2 different
4	VAKLYQIHAE		
1	SYKTSDAKGS	}	4 different
		}	9 different

You would obviously conclude that 2 and 3 were the closest relatives and that 4 was closer to 2 and 3 than 1.



If a constant rate of mutation is assumed, the Divergence time may be calculated

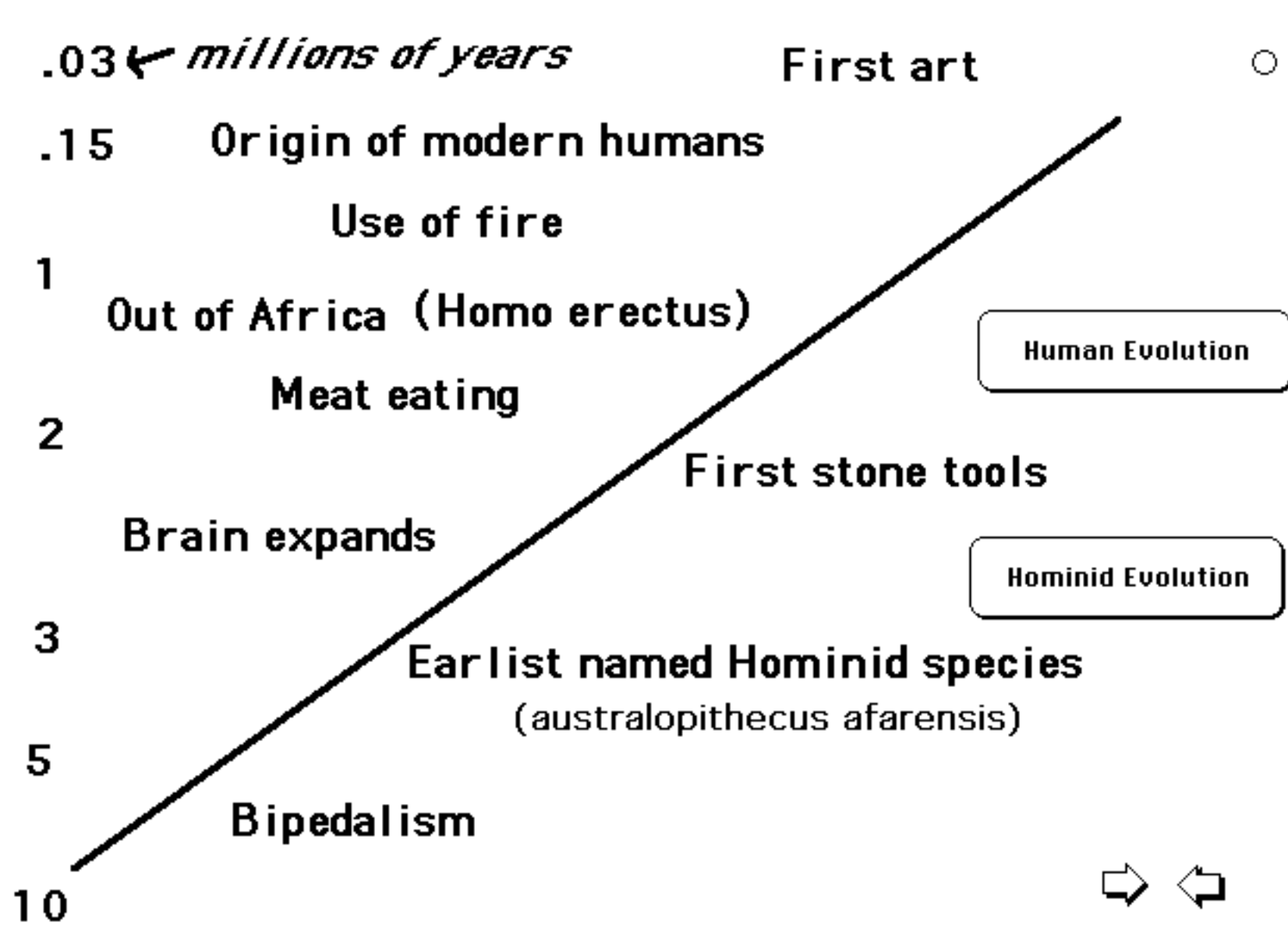


There are two approaches to tracing the family tree of humans:

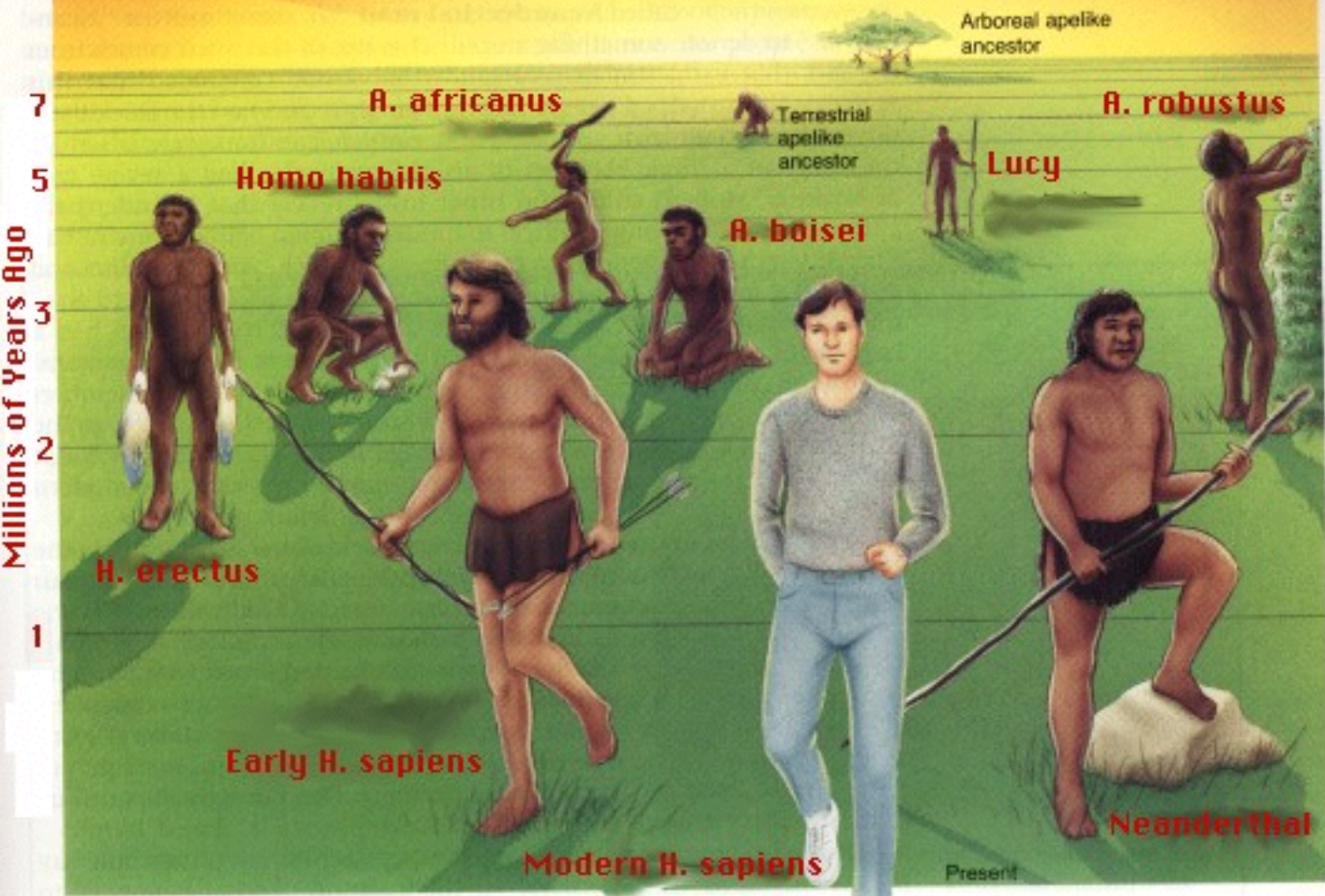
- 1. Examine the fossil record – Paleontology**
- 2. Examine the genes of living organisms – Molecular genetics**

Genetic studies have been taken to indicate that modern man (Homo Sapiens) arose in Africa 200,000 years ago

Fossil evidence argues that various human groups arose where they are found today.



TIMELINE OF HOMINID EVOLUTION



Mass Extinctions

The geological record show a number of instances in which the number of different species dropped dramatically These are mass extinctions brought about by some change in the environment:

Global climate change

Glaciation (Ice ages)

Continental Drift

Global warming (rise in sea levels)

Extraterrestrial impacts

Where is evolution going now?

Should we try to control the gene pool?

Cloning

Making a genetic copy of an organism