Genes and culture in language evolution
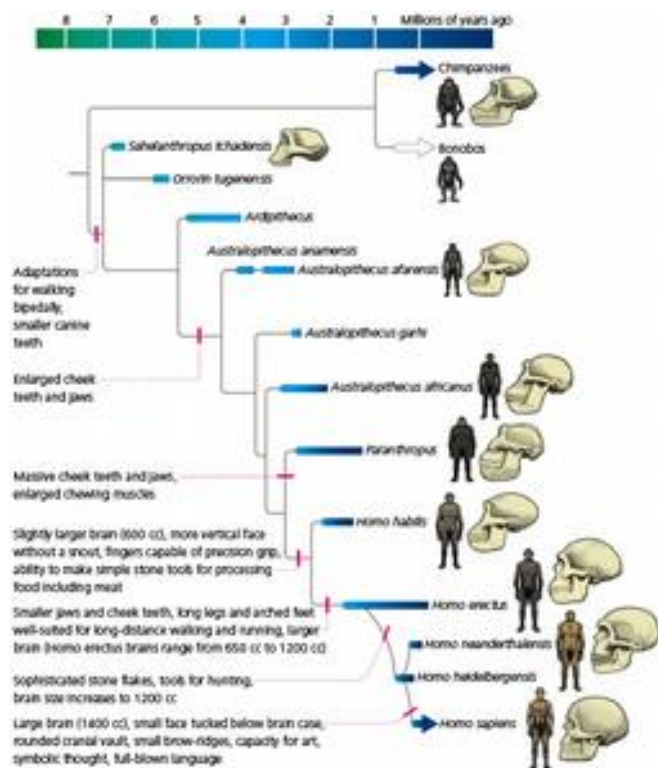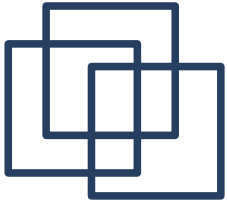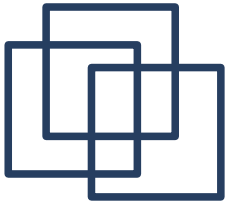
Dan Dediu

Language and Genetics
Max Planck Institute for Psycholinguistics
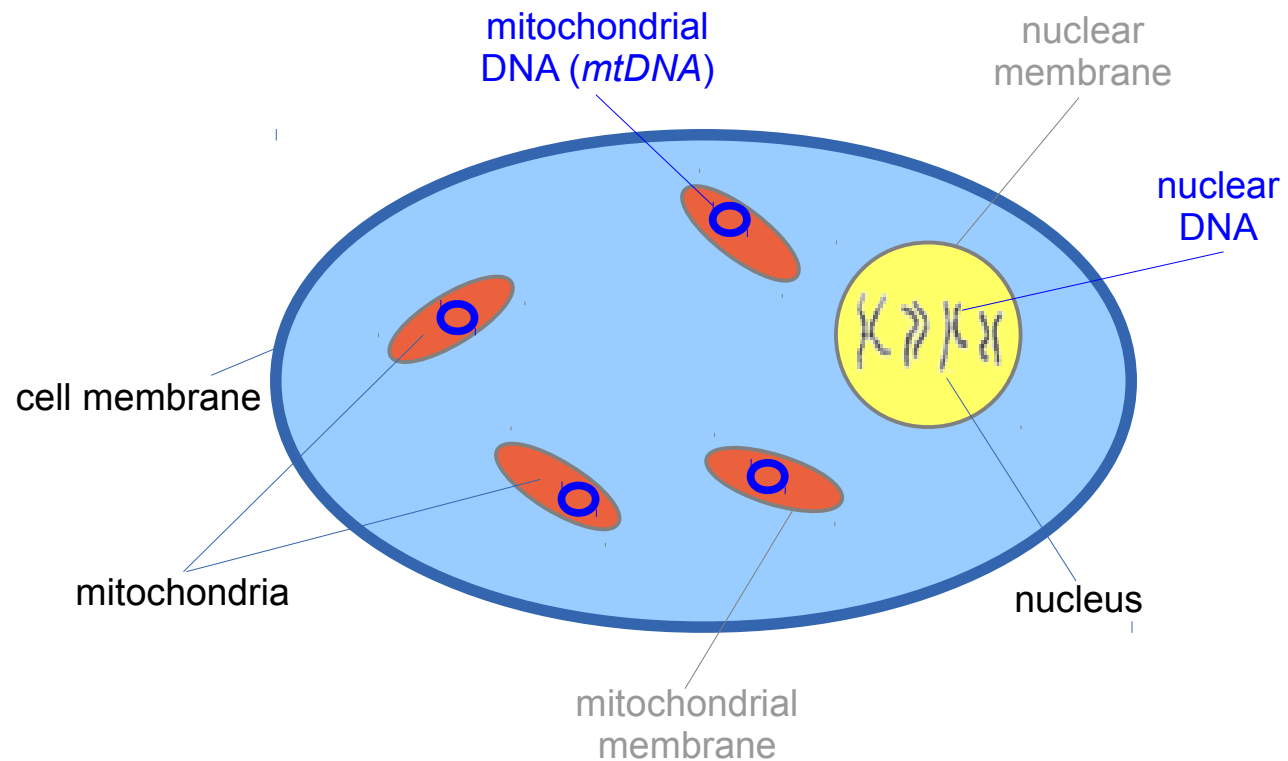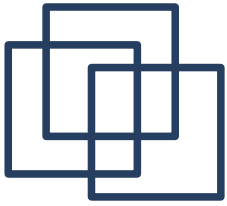Nijmegen
The Netherlands

Overview

- Population and evolutionary genetics

- Human evolution

- Human genetic and linguistic diversity

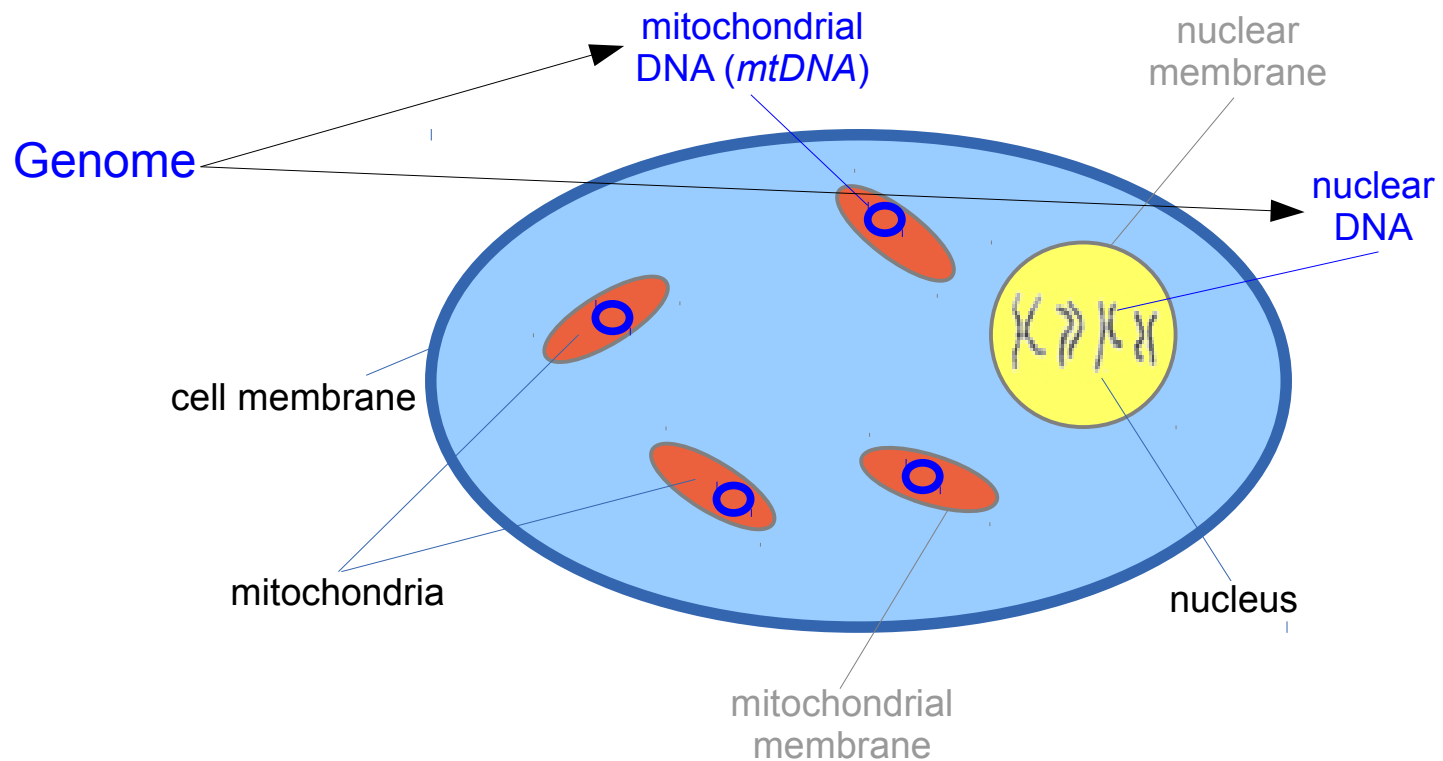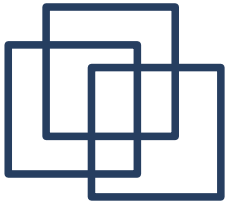- Genetic biasing, gene-culture co-evolution, niche construction

Population and evolutionary genetics

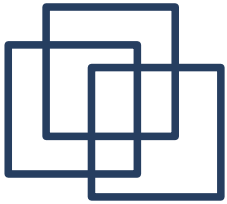# Population and evolutionary genetics

Population and evolutionary genetics

- Some genes (energy production)
- Own genetic code
- "D-loop" → *hypervariable regions* (HVR-I & HVR-II)

Population and evolutionary genetics

- Some genes (energy production)
- Own genetic code
- "D-loop" → *hypervariable regions* (HVR-I & HVR-II)

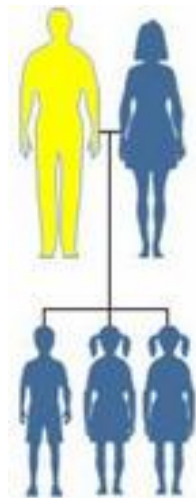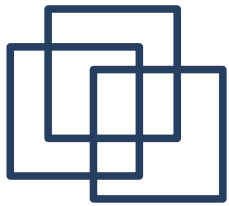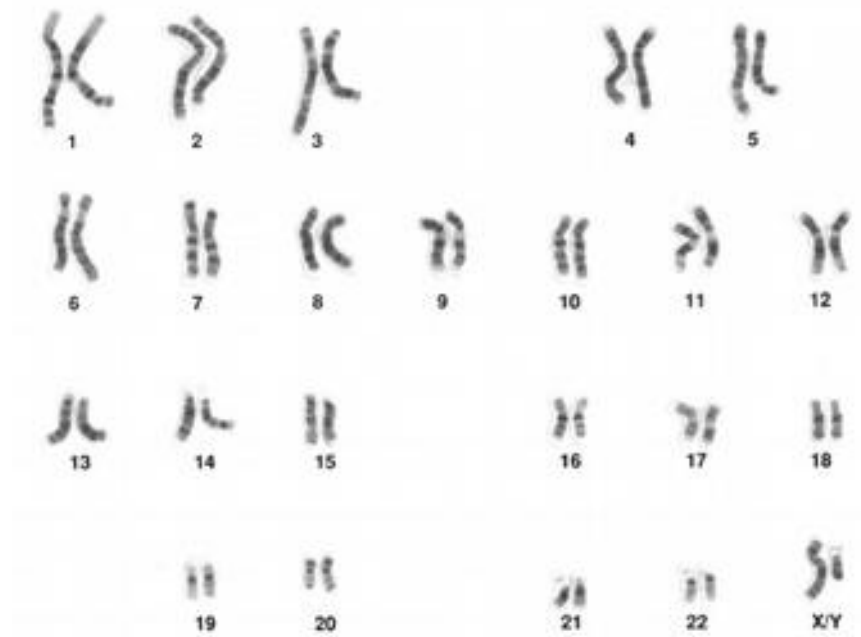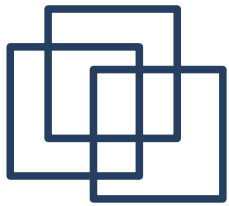**Maternal inheritance**

Population and evolutionary genetics

- **Chromosomes**: single linear molecules of DNA
- **Karyotype** = all chromosomes in the nucleus
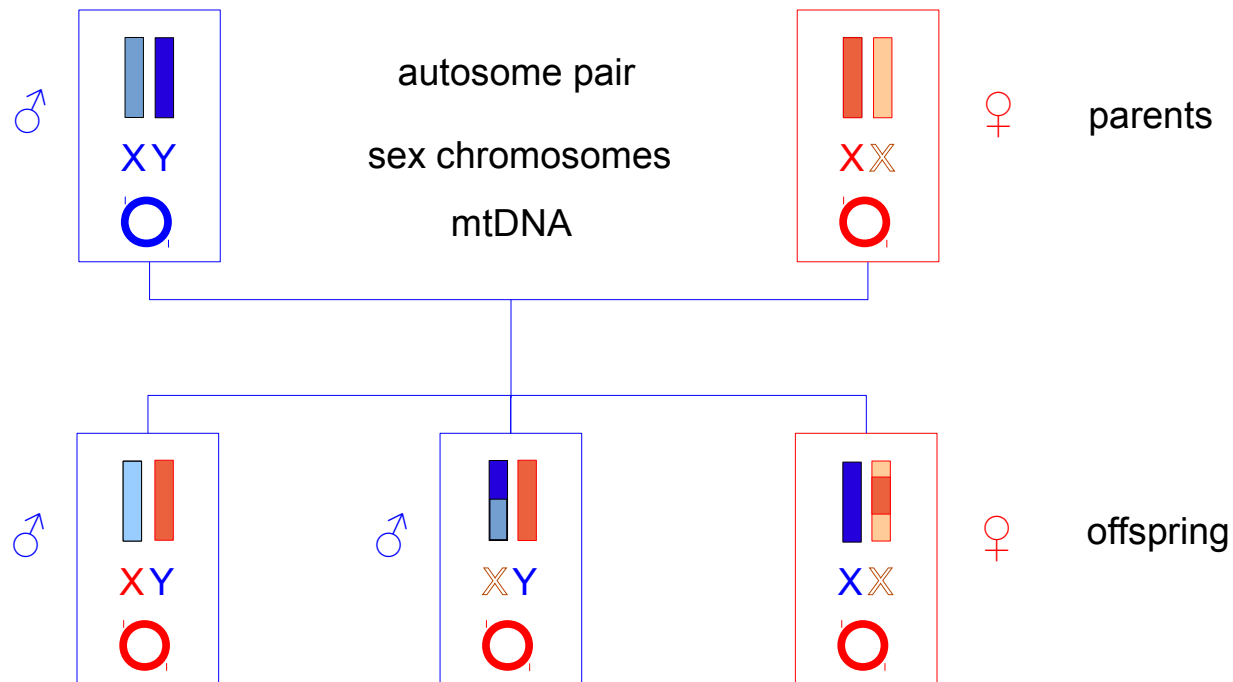- Humans: **23 pairs** of chromosomes (diploid)

- 22 pairs **autosomes**
- 1 pair **sex chromosomes**
  - **XX**=♀ **XY**=♂
- **Inheritance**:
  - X & autosomes: both parents
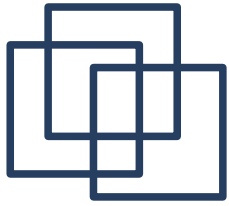 - Y: paternal inheritance

Population and evolutionary genetics
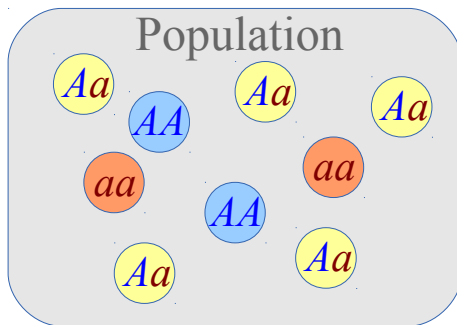
- the **inheritance** of the nuclear genome
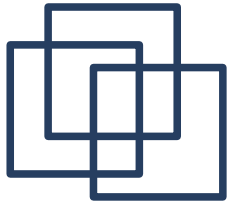- **recombination**, **independent assortment**

- Bi-allelic autosomal locus $\rightarrow A$ & a

Population and evolutionary genetics
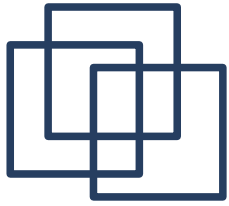
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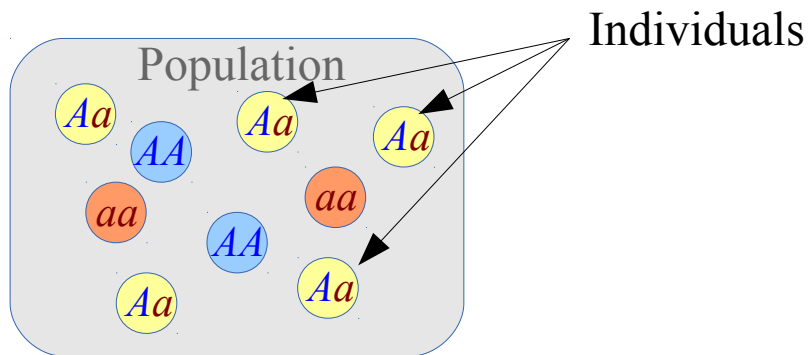
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Population and evolutionary genetics

- Bi-allelic autosomal locus $\rightarrow A$ & a

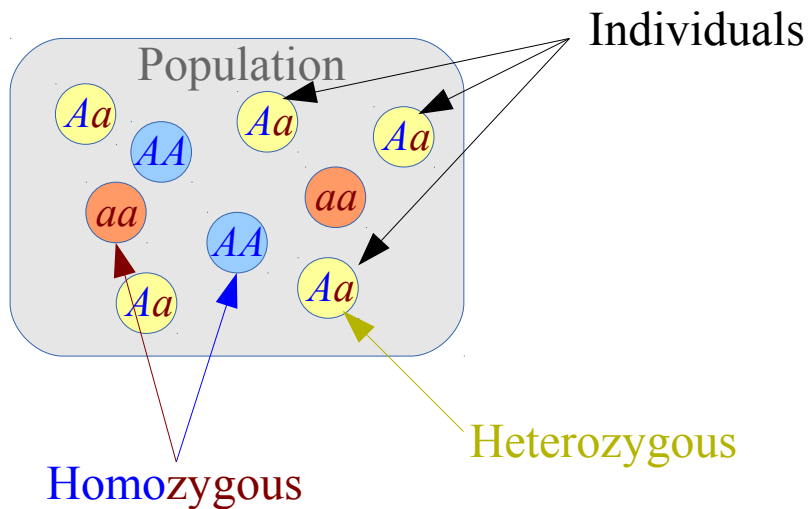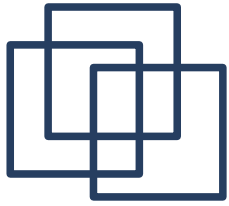- Allele frequencies p_A & p_a

- Genotype frequencies p_{AA}, p_{Aa} & p_{aa}

Population and evolutionary genetics

- Bi-allelic autosomal locus $\rightarrow A$ & a

- Allele frequencies p_A & p_a

- Genotype frequencies p_{AA}, p_{Aa} & p_{aa}

t

p_A, p_a; p_{AA}, p_{Aa} & p_{aa}

Time

Population and evolutionary genetics

- Bi-allelic autosomal locus $\rightarrow$ A & a

- Allele frequencies p_A & p_a

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p_A, p_a; p_{AA}, p_{Aa} & p_{aa}

$t+1$

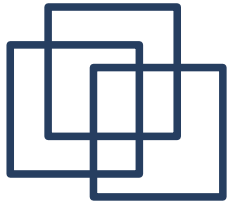p'_A, p'_a; p'_{AA}, p'_{Aa} & p'_{aa}
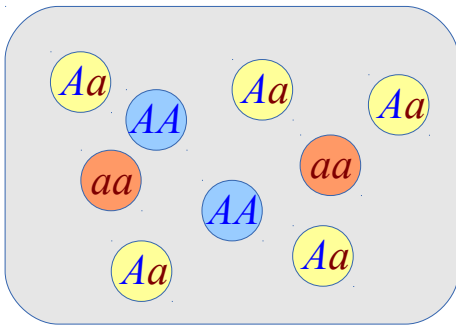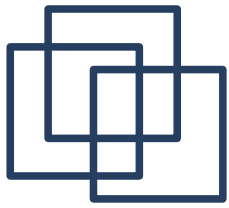
Time

Population and evolutionary genetics

- Bi-allelic autosomal locus $\rightarrow A$ & a

- Allele frequencies p_A & p_a

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t

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

conditions & factors

$t+1$

$p'_A, p'_a; p'_{AA}, p'_{Aa}$ & p'_{aa}

Time

Population and evolutionary genetics

- Bi-allelic autosomal locus $\rightarrow A$ & a

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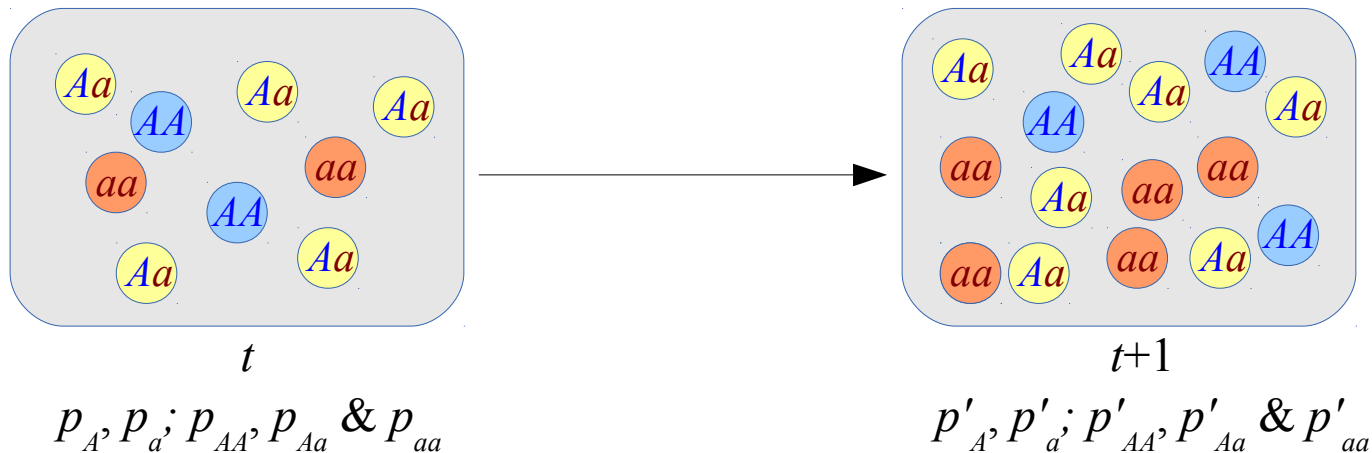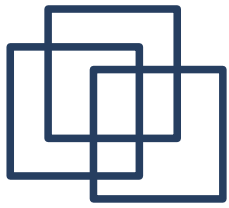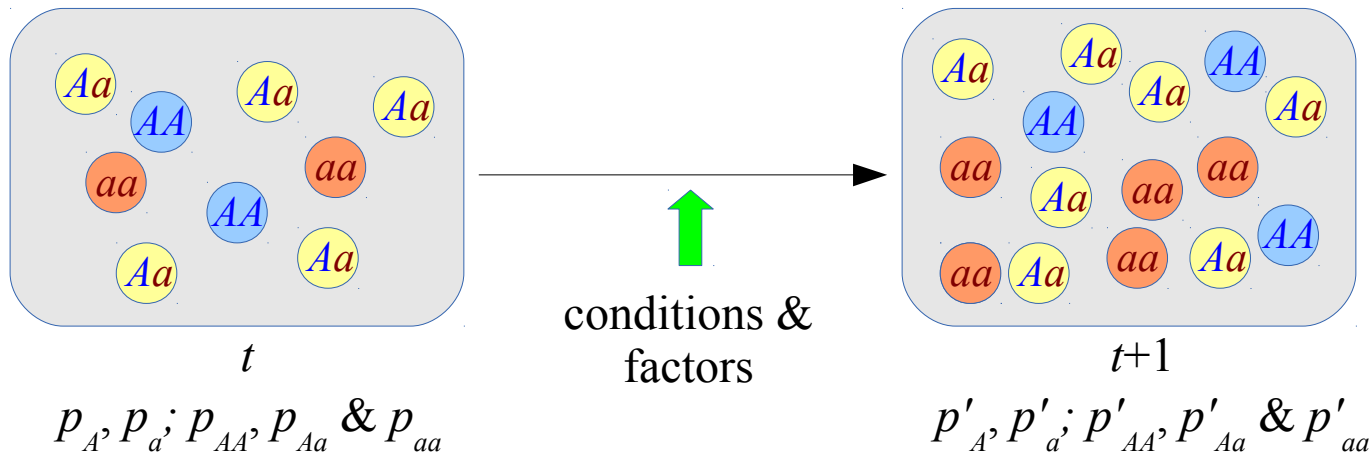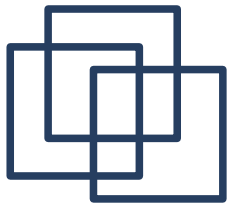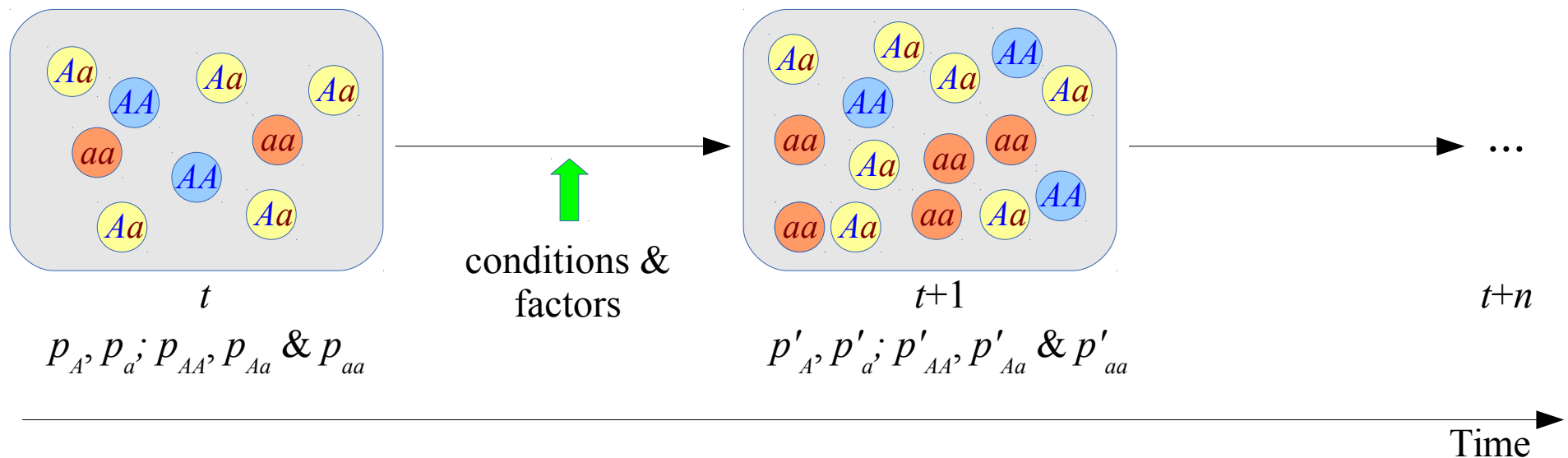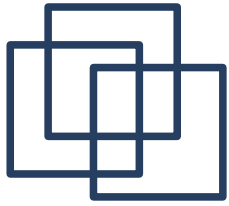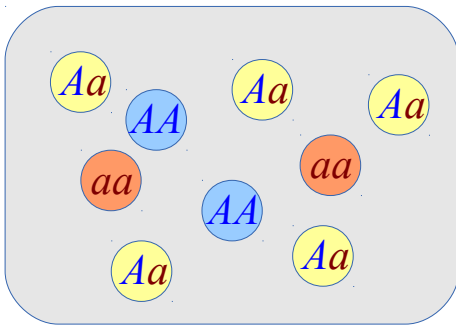- Genotype frequencies p_{AA}, p_{Aa} & p_{aa}

t

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

conditions & factors

$t+1$

$p'_A, p'_a; p'_{AA}, p'_{Aa}$ & p'_{aa}

$t+n$

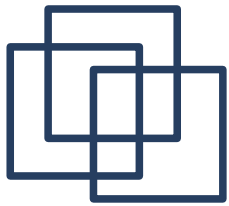...

Time

Population and evolutionary genetics

- **Hardy-Weinberg Equilibrium** (HWE):

  if nothing interesting happens → no change

t

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

Time

Population and evolutionary genetics

- Hardy-Weinberg Equilibrium (HWE):

  if nothing interesting happens → no change

t

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

- infinite population
- no mutation
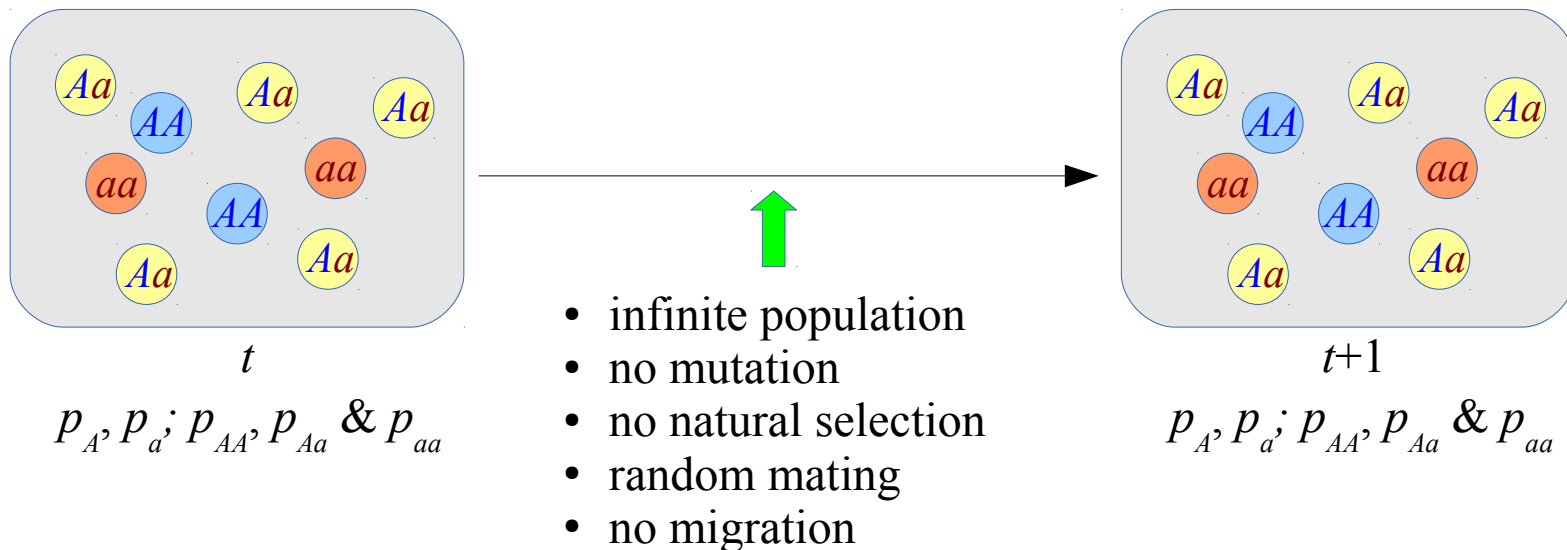- no natural selection
- random mating
- no migration

$t+1$

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

Time

Population and evolutionary genetics

- Hardy-Weinberg Equilibrium (HWE):

  if nothing interesting happens → no change
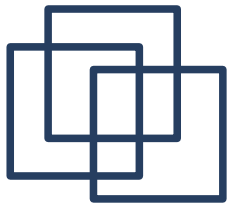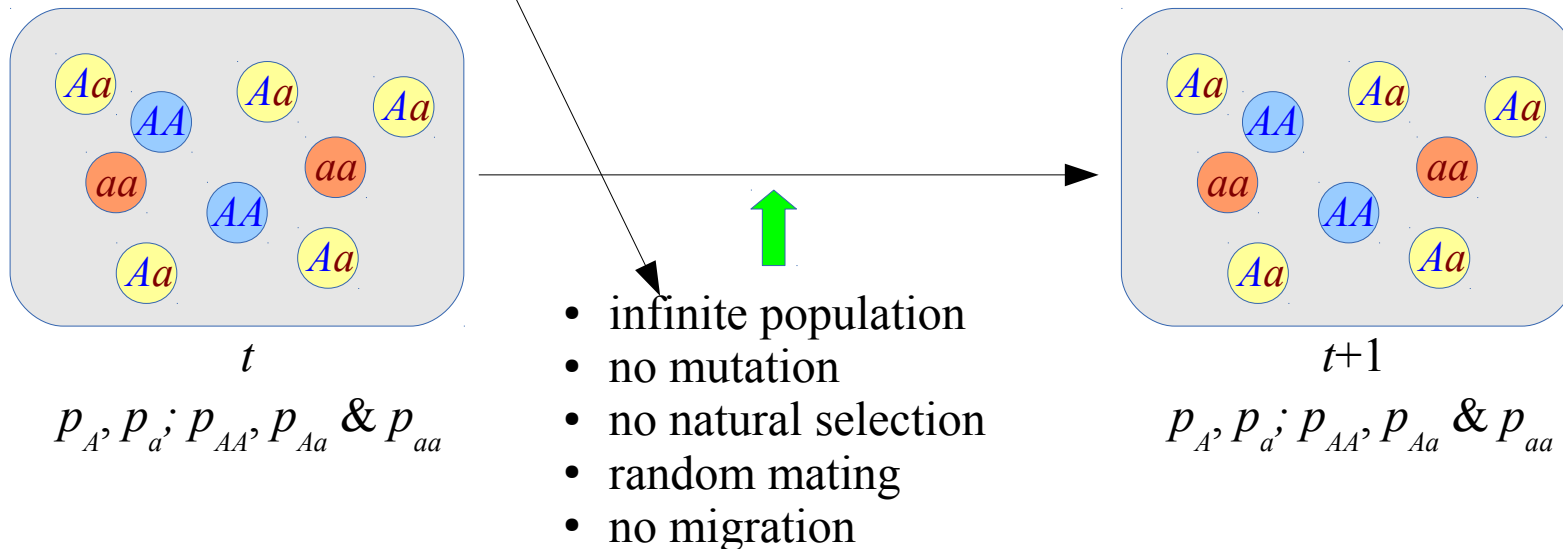

t

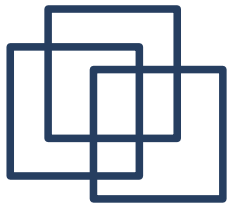$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

- infinite population
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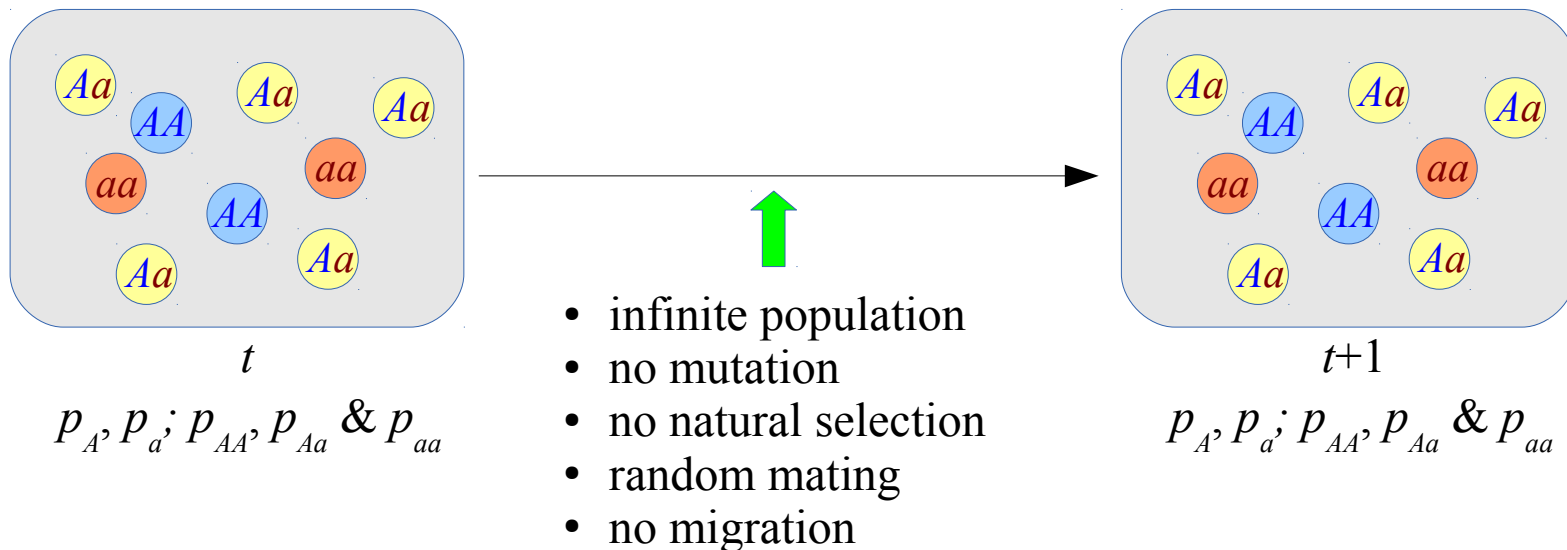- no migration
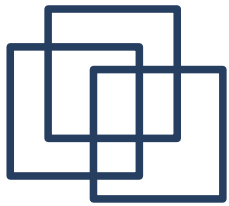
$t+1$

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

Time

Population and evolutionary genetics

- Hardy-Weinberg Equilibrium (HWE):

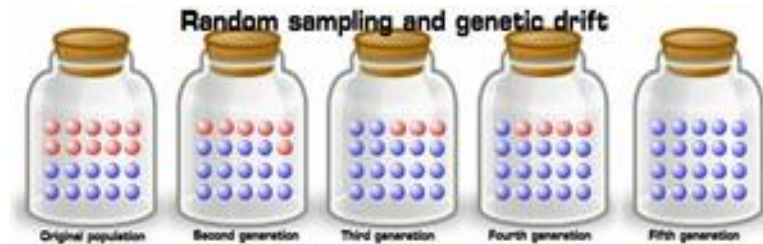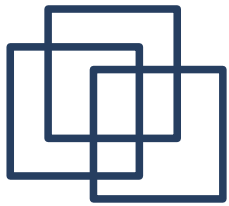 if nothing interesting happens → no change

  → simple null hypothesis

t

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

- infinite population
- no mutation
- no natural selection
- random mating
- no migration

$t+1$

$p_A, p_a; p_{AA}, p_{Aa}$ & p_{aa}

Time

Population and evolutionary genetics

- Finite populations $\rightarrow$ reproduction = random sampling

Random sampling and genetic drift

Population and evolutionary genetics

- Finite populations → reproduction = random sampling

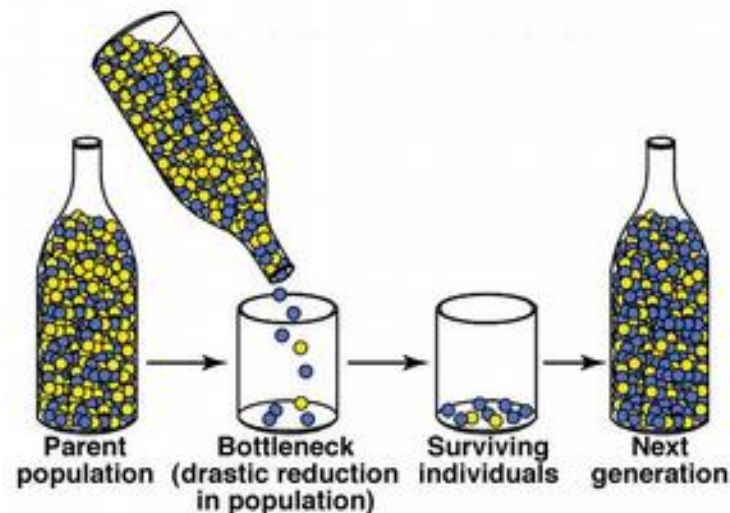- Bottlenecks → "founder effect" (e.g., SLI on Robinson Crusoe Island)

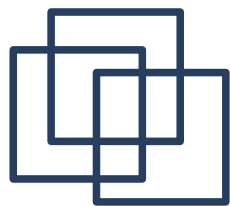Parent population | Bottleneck (drastic reduction in population) | Surviving individuals | Next generation
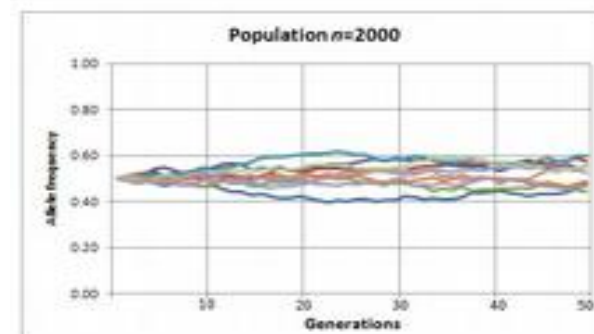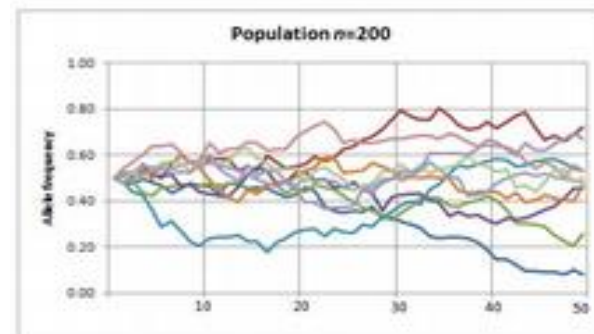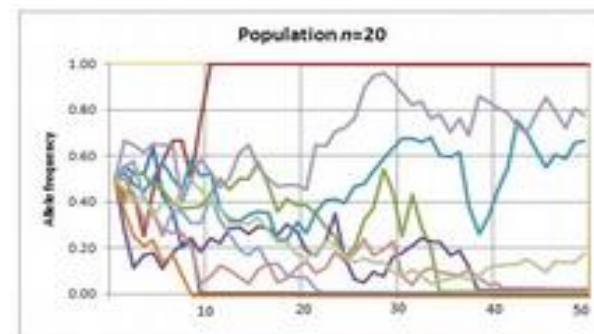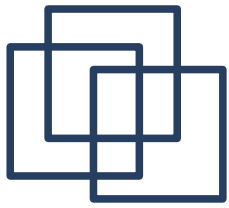
Photo courtesy of Miami University

Population and evolutionary genetics

- Finite populations → reproduction = random sampling

- Bottlenecks → "founder effect"

- (Effective) population size, N_e

Population and evolutionary genetics

- Finite populations → reproduction = random sampling

- Bottlenecks → "founder effect"

- (Effective) population size, N_e
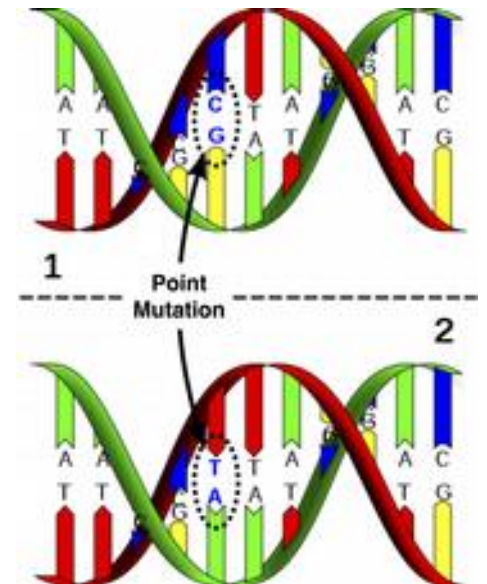
$$\rightarrow \text{reduces genetic diversity}$$
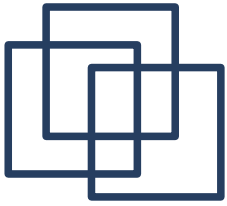
Population and evolutionary genetics

- Mutation → changes the genetic information (e.g., $A \to a$)

- Many types: point mutation

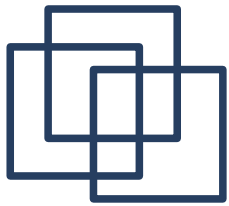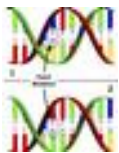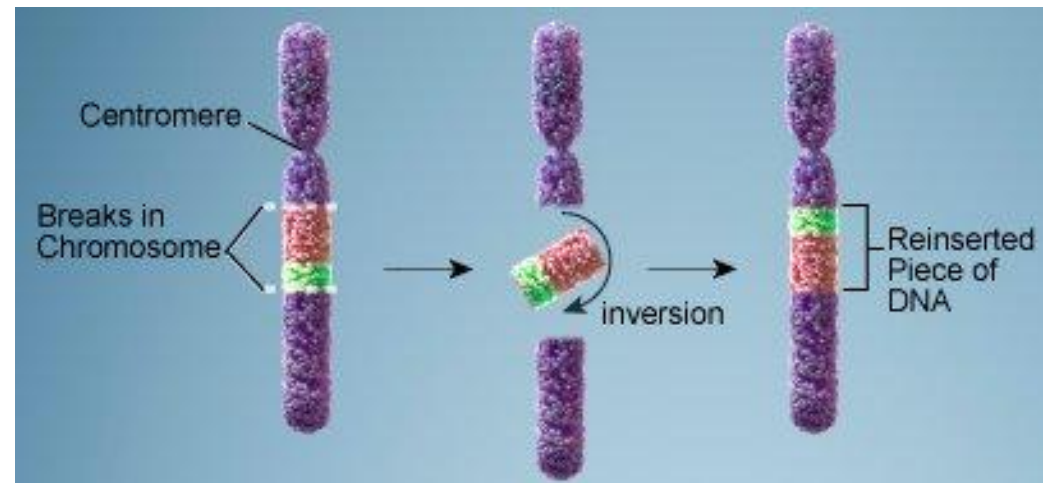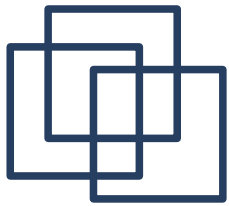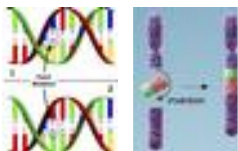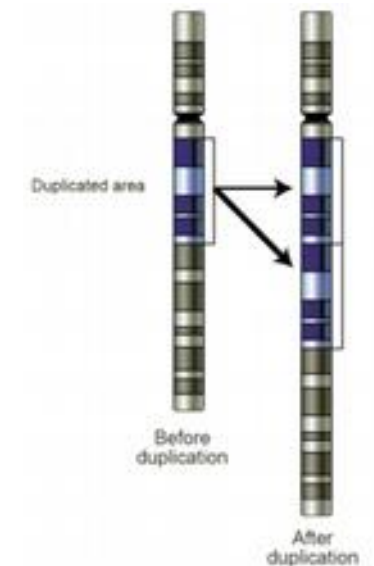# Population and evolutionary genetics

- Mutation → changes the genetic information (e.g., $A \rightarrow a$)
- Many types: point mutation, chromosomal rearrangements

Population and evolutionary genetics

- Mutation → changes the genetic information (e.g., $A \to a$)

- Many types: point mutation, chromosomal rearrangements, gene duplications
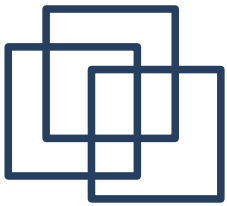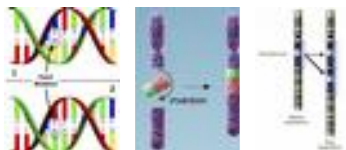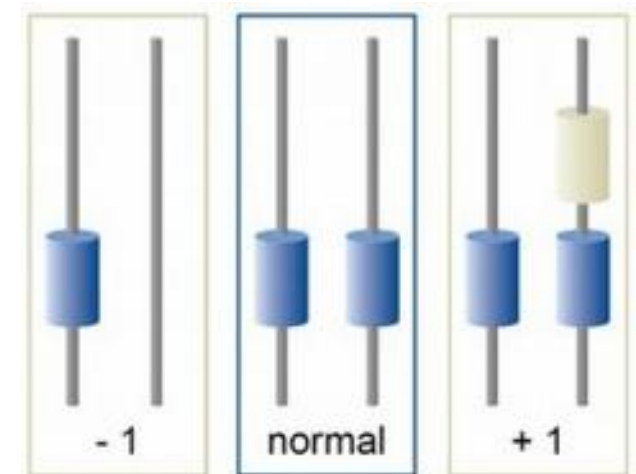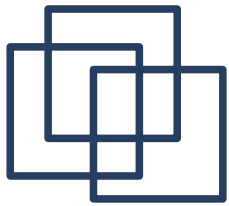

Duplicated area

Before duplication

After duplication

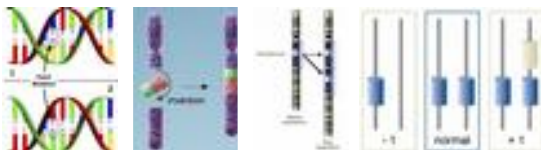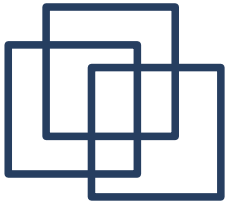# Population and evolutionary genetics

- Mutation → changes the genetic information (e.g., $A \to a$)

- Many types: point mutation, chromosomal rearrangements, gene duplications, CNVs...

Population and evolutionary genetics

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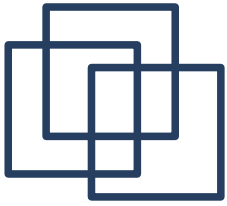
$$\to \text{ increases genetic diversity}$$

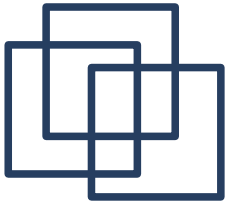- Selection → differential survival and reproduction

Population and evolutionary genetics

- Selection → differential survival and reproduction

- Types:

natural

Population and evolutionary genetics

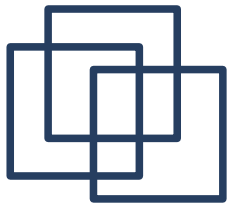- Selection → differential survival and reproduction

- Types:

natural   vs   sexual

Population and evolutionary genetics

- Selection → differential survival and reproduction
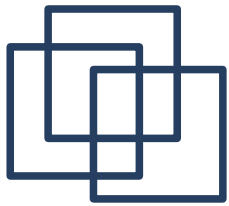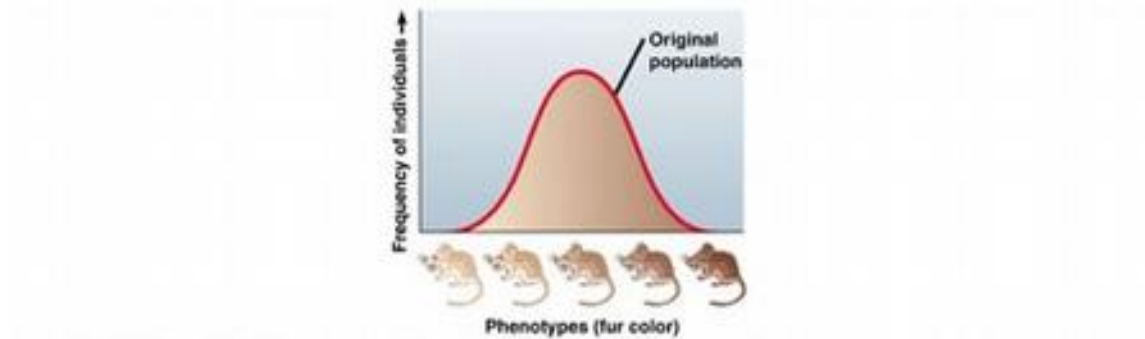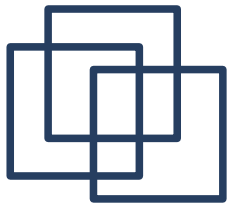- Types:

natural   vs   sexual   vs   artificial
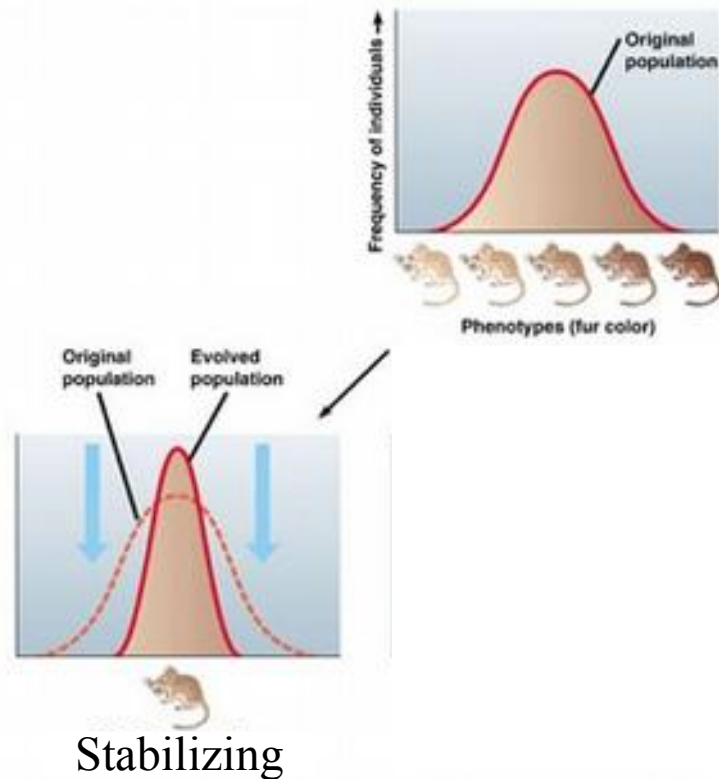
Population and evolutionary genetics

- Selection → differential survival and reproduction

- Types:

Population and evolutionary genetics

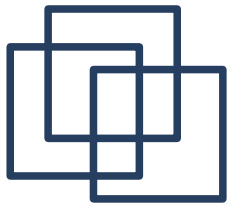- Selection → differential survival and reproduction

- Types:

Stabilizing

Population and evolutionary genetics

- Selection → differential survival and reproduction
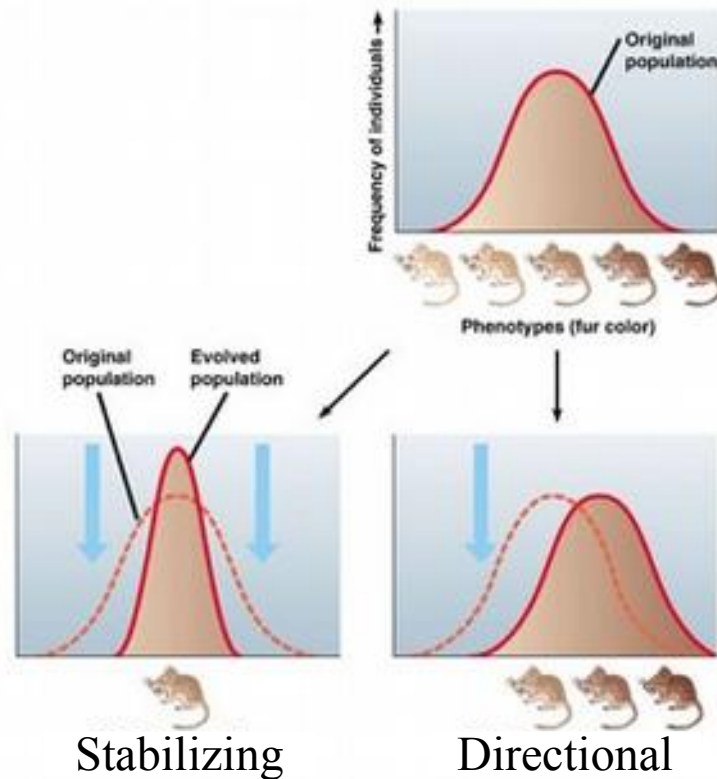- Types:

Population and evolutionary genetics

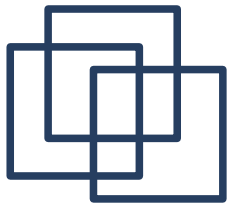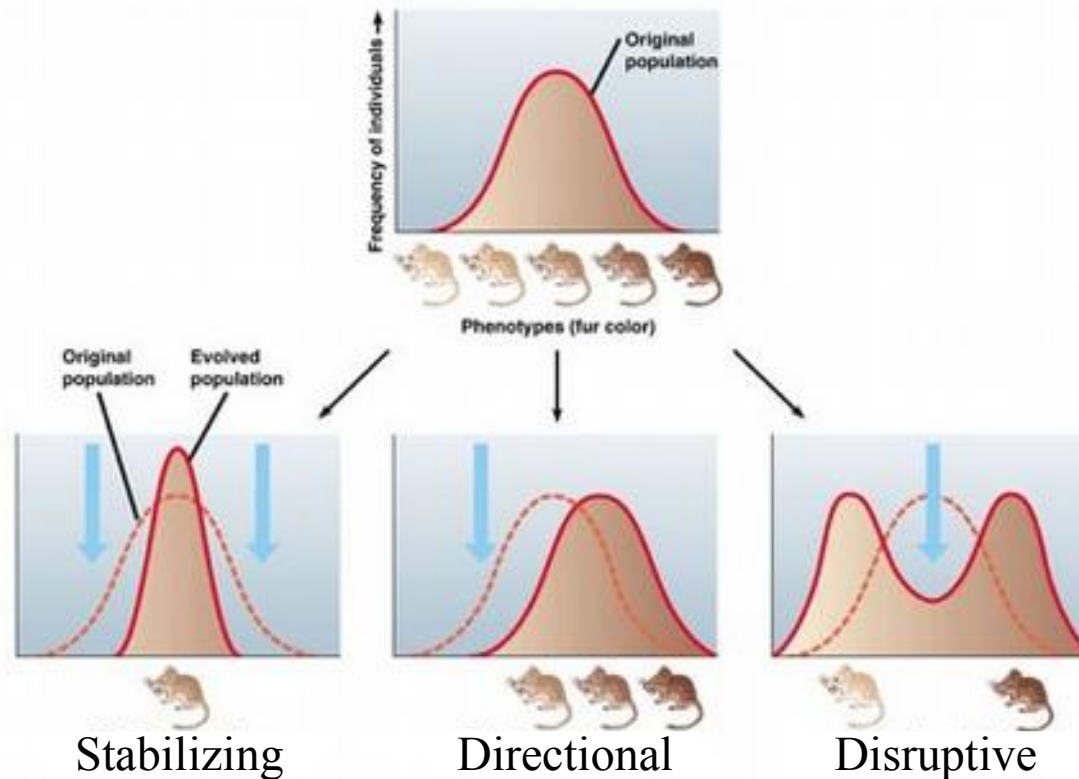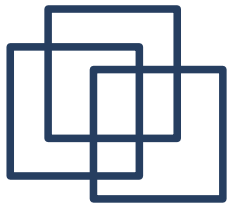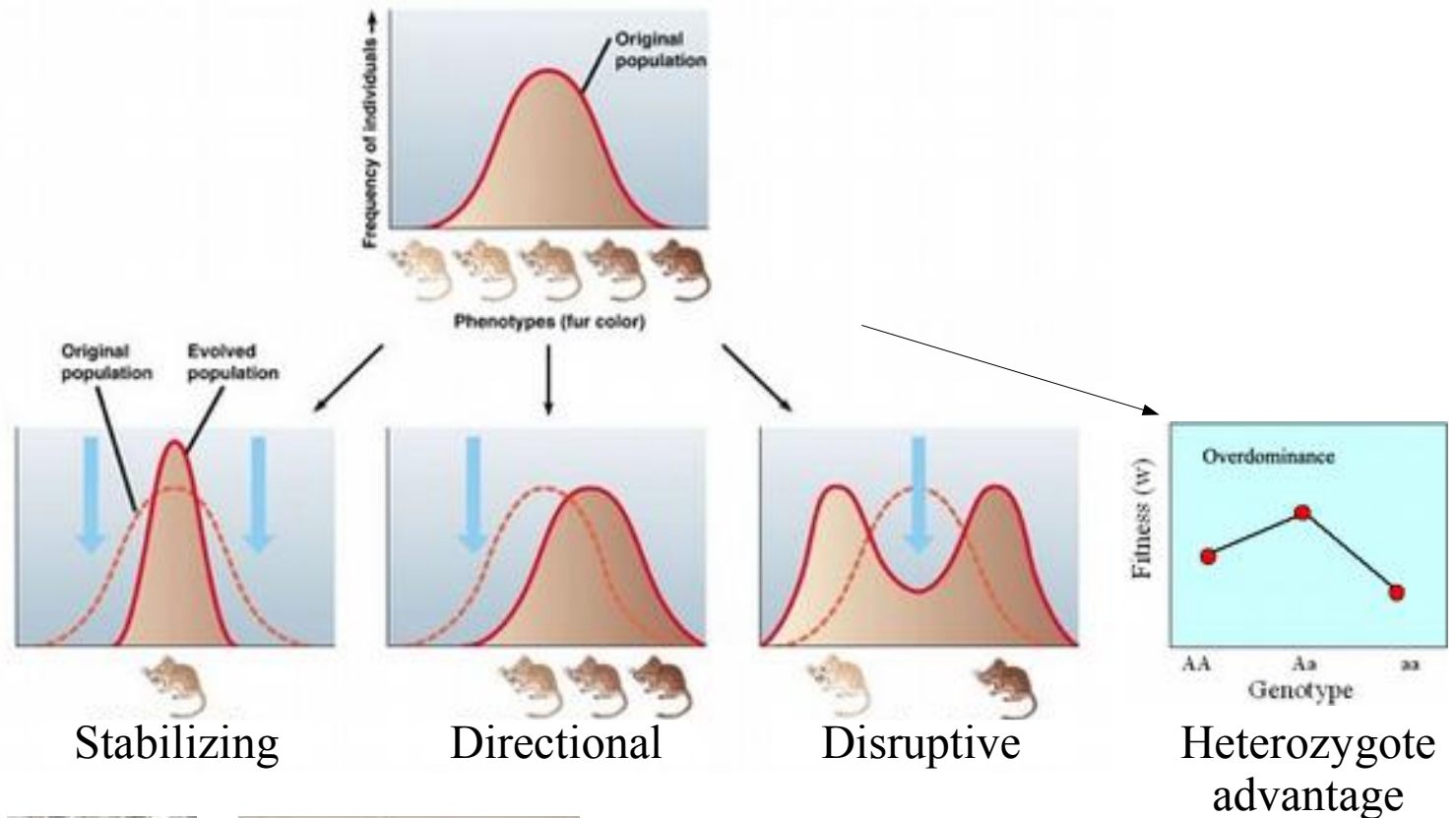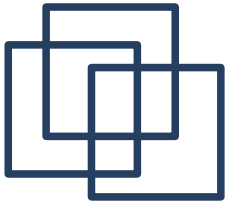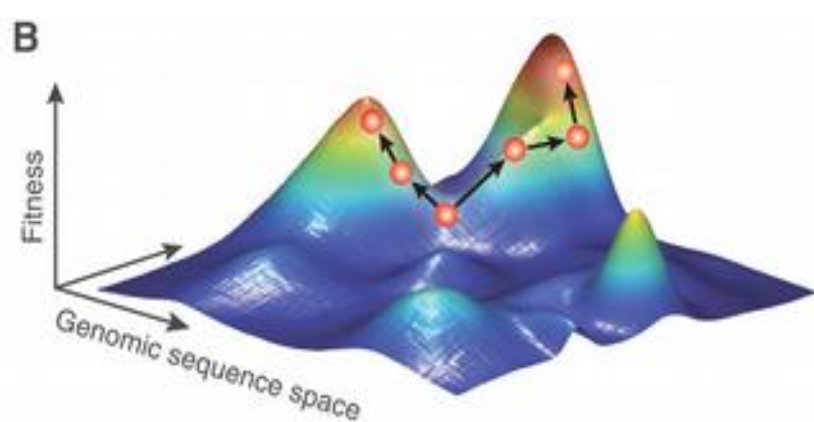- Selection → differential survival and reproduction
- Types:

Population and evolutionary genetics

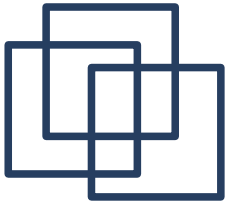- Selection → differential survival and reproduction
- Types:

Stabilizing   Directional   Disruptive   Heterozygote advantage
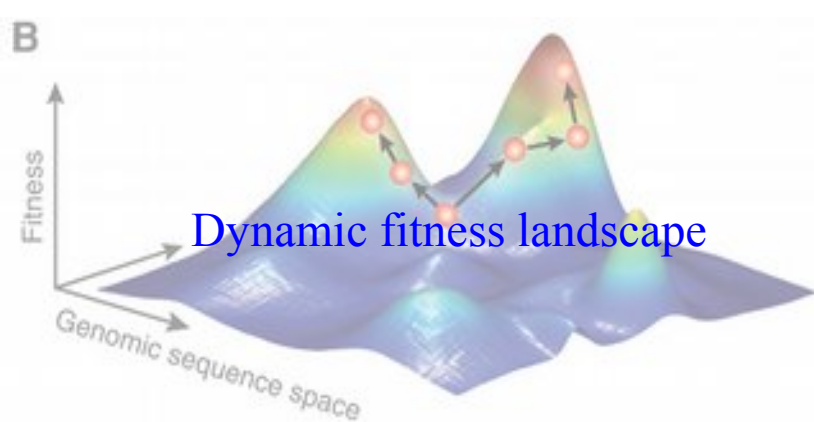
Population and evolutionary genetics

- Selection $\rightarrow$ differential survival and reproduction
- Walking the fitness landscape

Population and evolutionary genetics

- Selection → differential survival and reproduction

- Walking the fitness landscape

Dynamic fitness landscape

Population and evolutionary genetics

- Selection → differential survival and reproduction
- Walking the fitness landscape

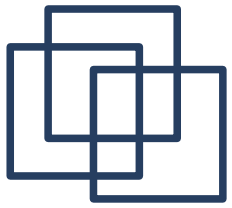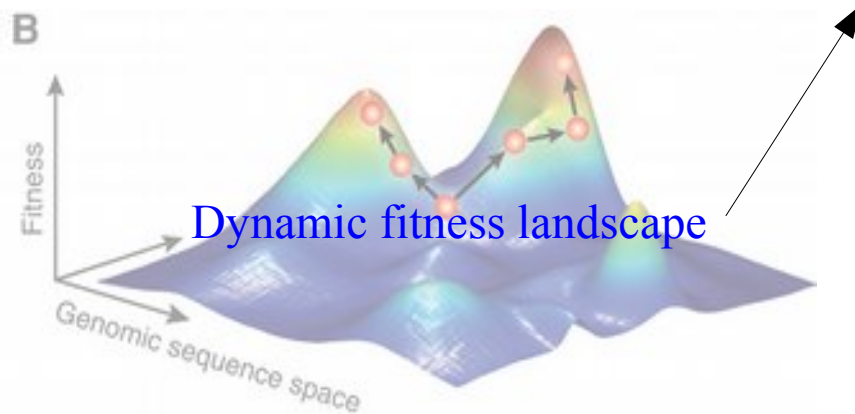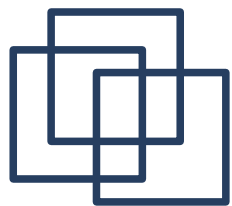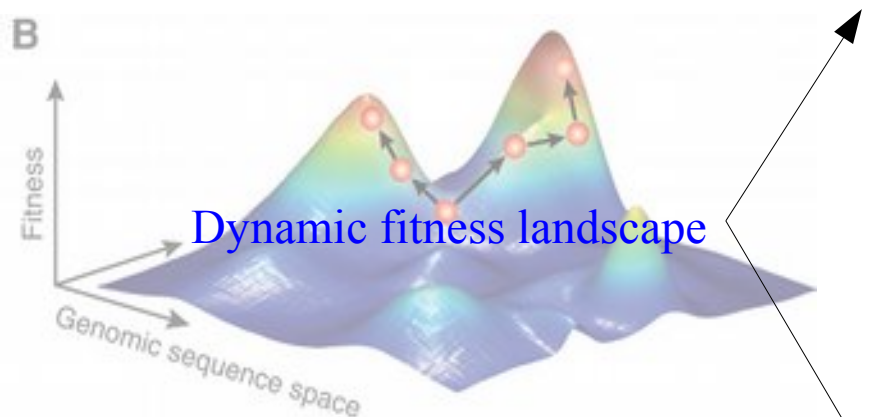Dynamic fitness landscape

Co-evolution & arms races

Population and evolutionary genetics

- Selection → differential survival and reproduction

- Walking the fitness landscape

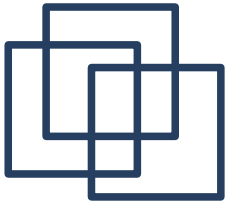Dynamic fitness landscape

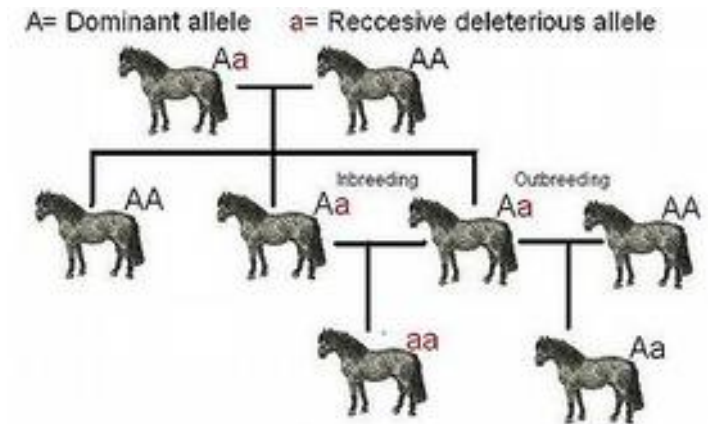Co-evolution & arms races

Niche construction
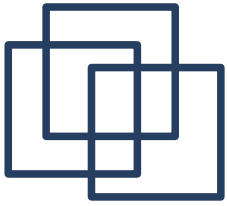
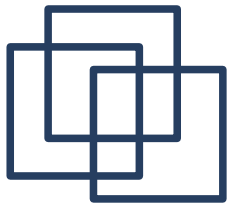# Population and evolutionary genetics

- Non-random mating

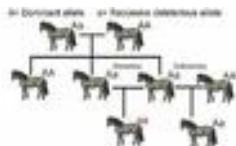# Population and evolutionary genetics

- **Non-random mating**

  – inbreeding

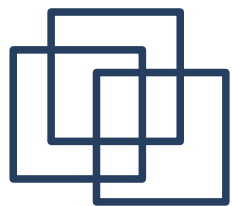# Population and evolutionary genetics

- Non-random mating

  - inbreeding

  - assortative mating

Population and evolutionary genetics

- **Non-random mating**

  - inbreeding

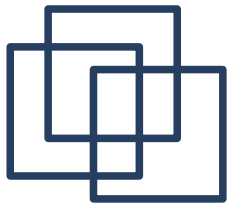 - assortative mating

  - population subdivision
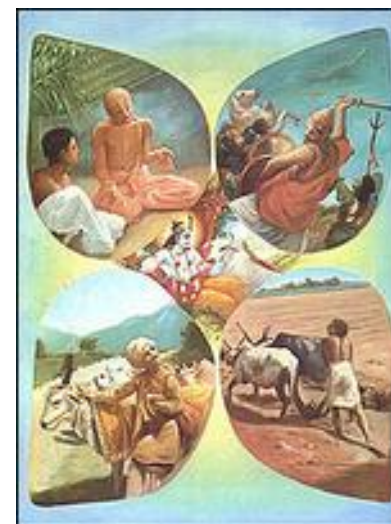
Population and evolutionary genetics

- **Non-random mating**

  - inbreeding

  - assortative mating

  - population subdivision

    → gene flow/admixture
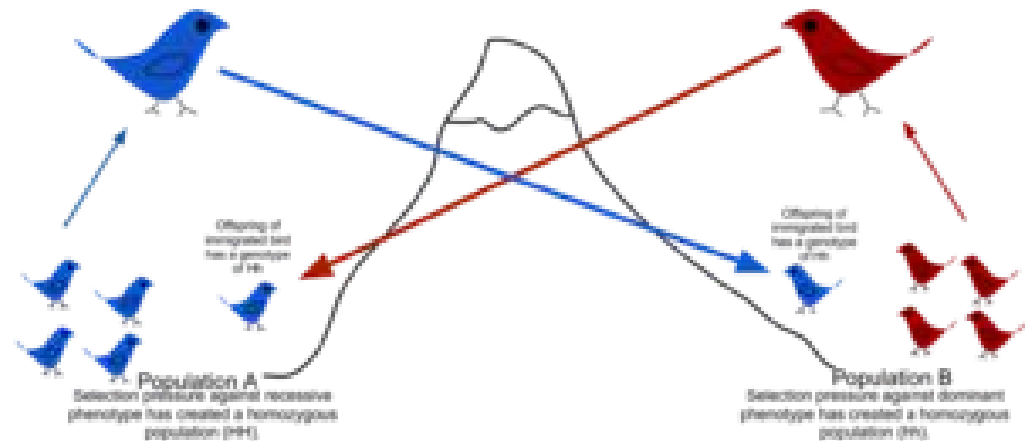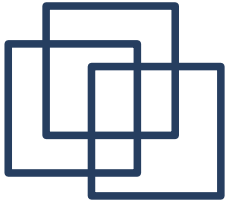
Human evolution
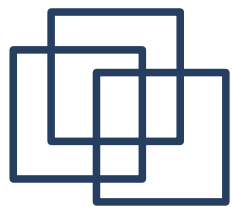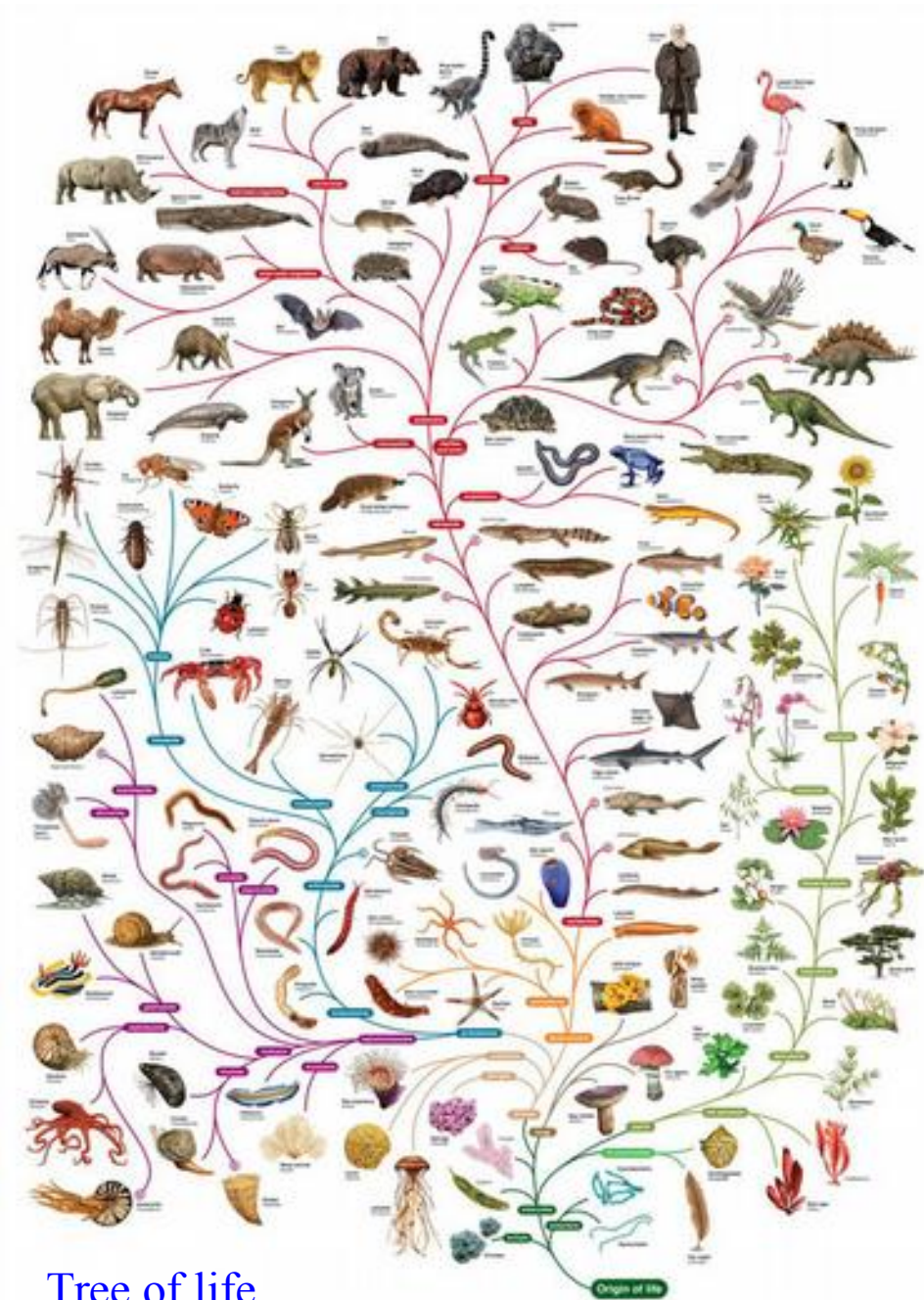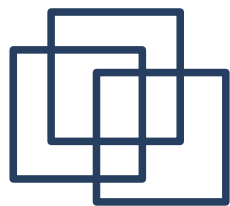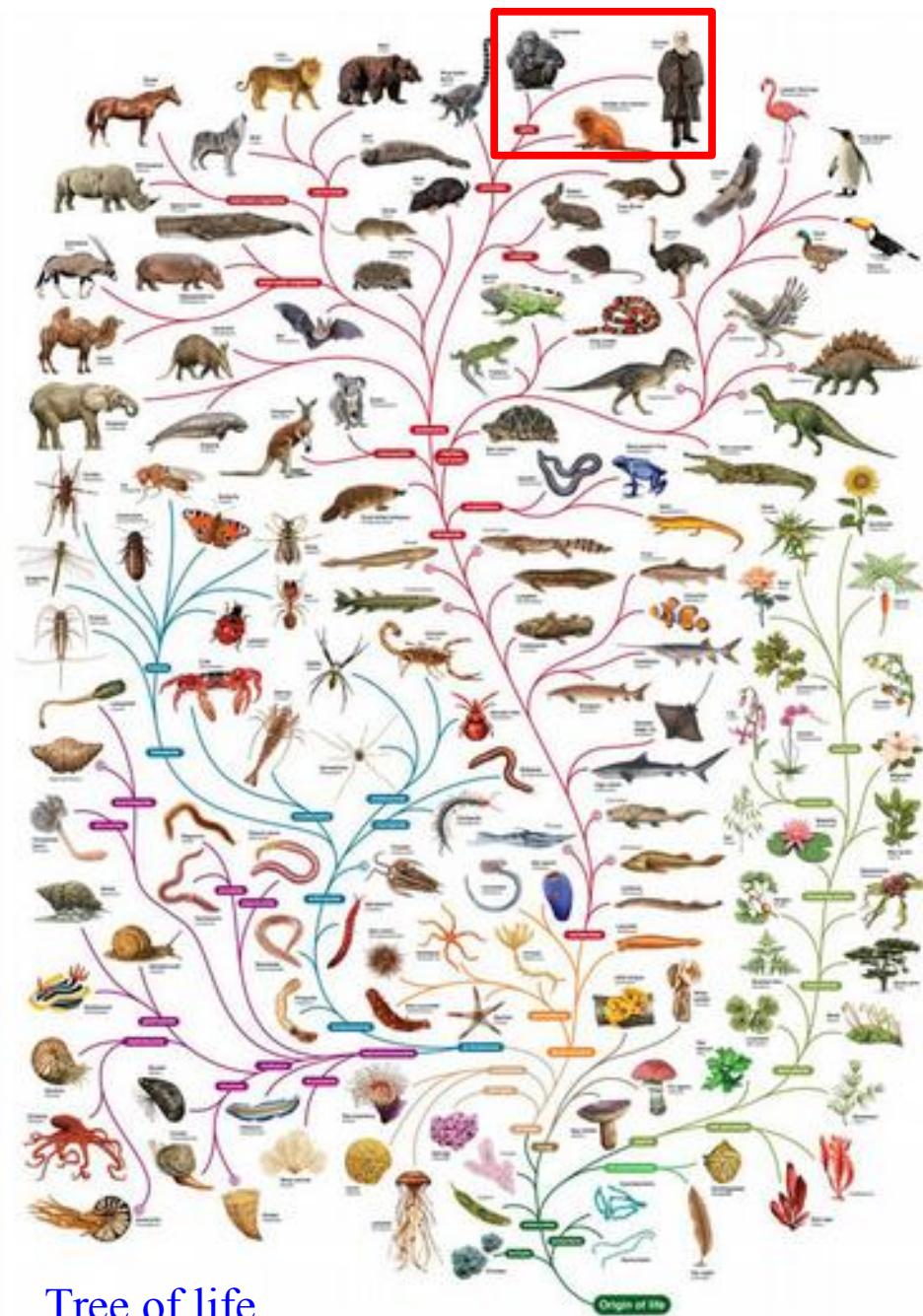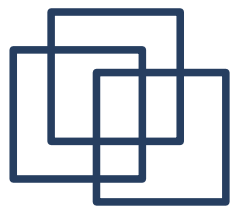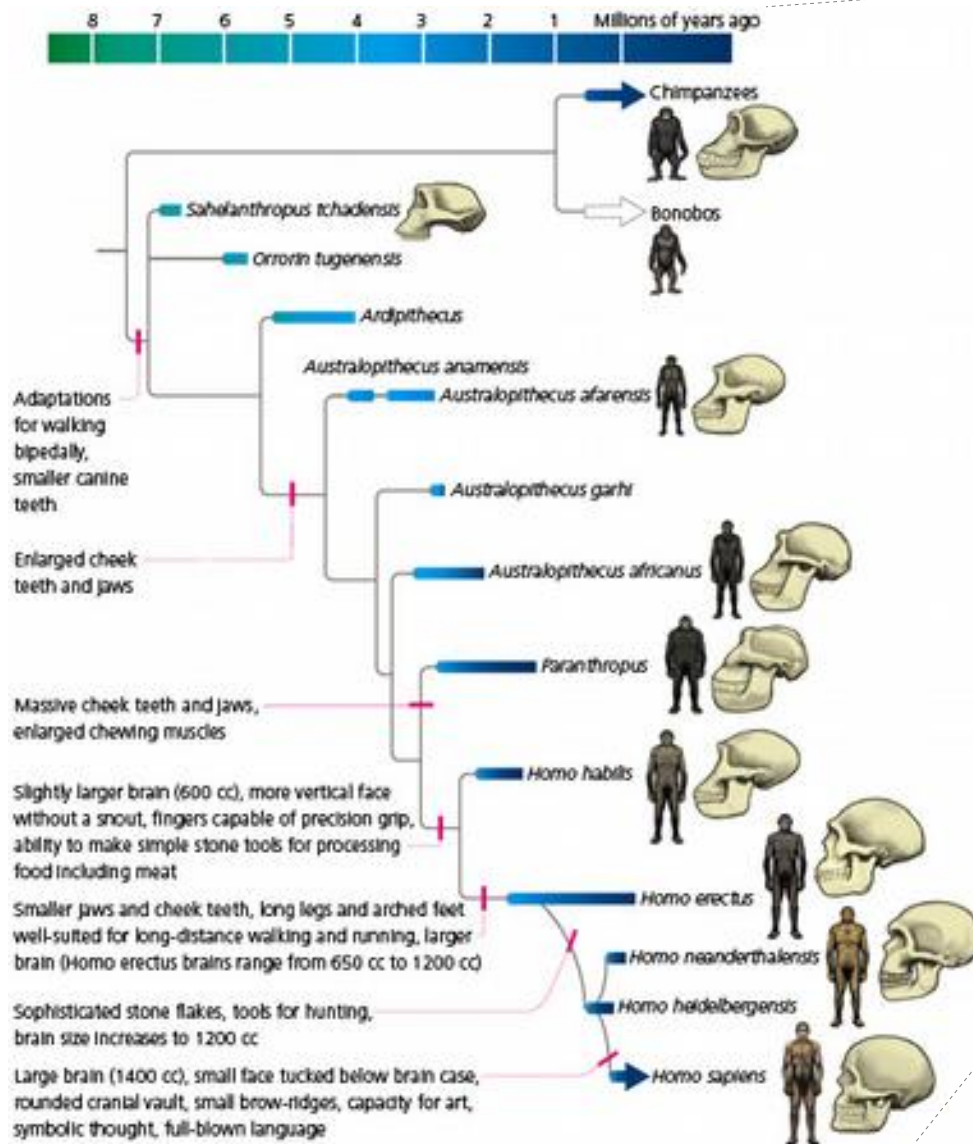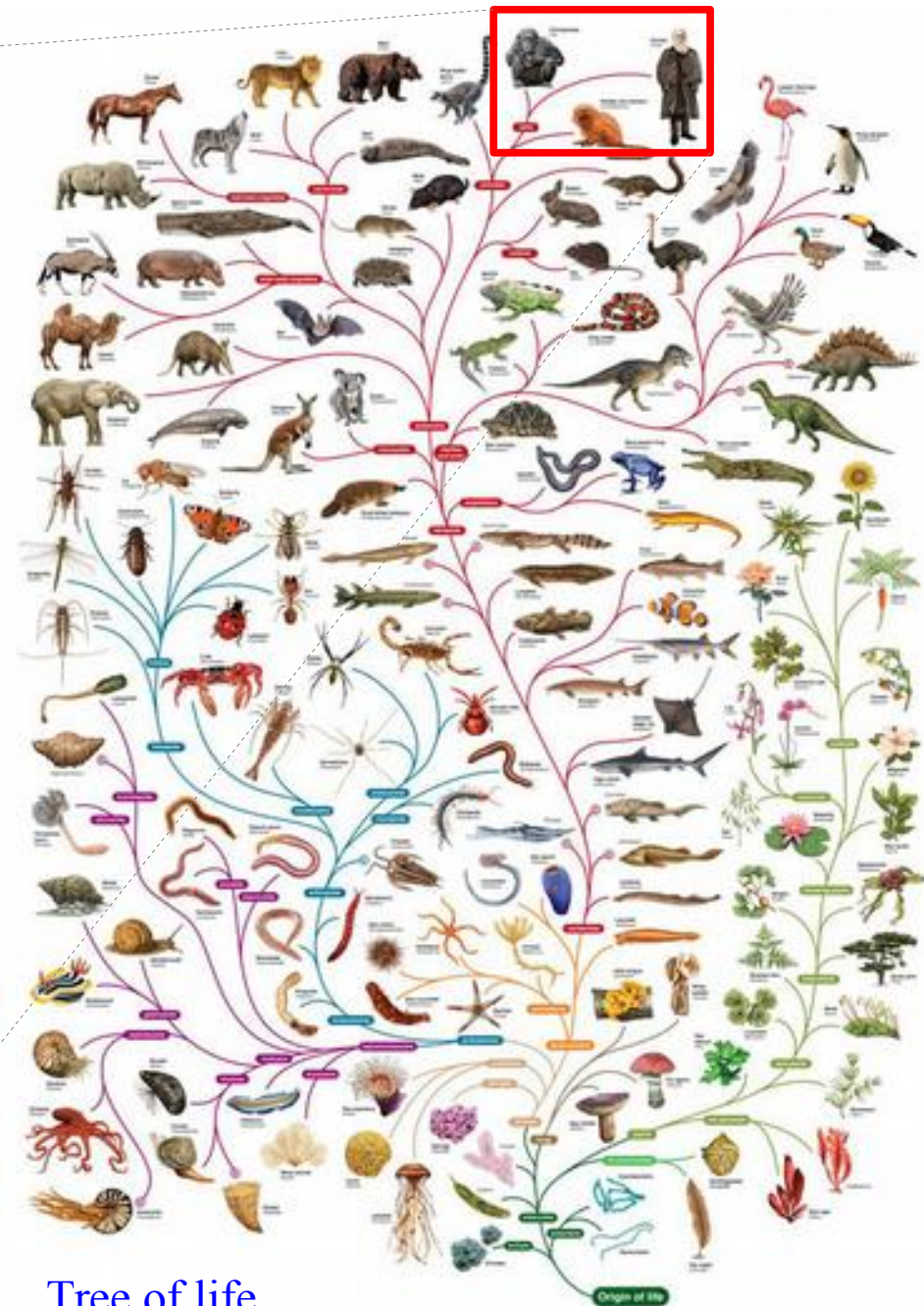
Human evolution

Tree of life

Human evolution

Tree of life

Human evolution

Human evolution

Tree of life
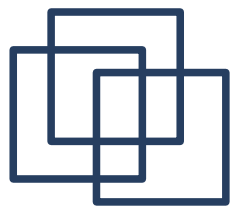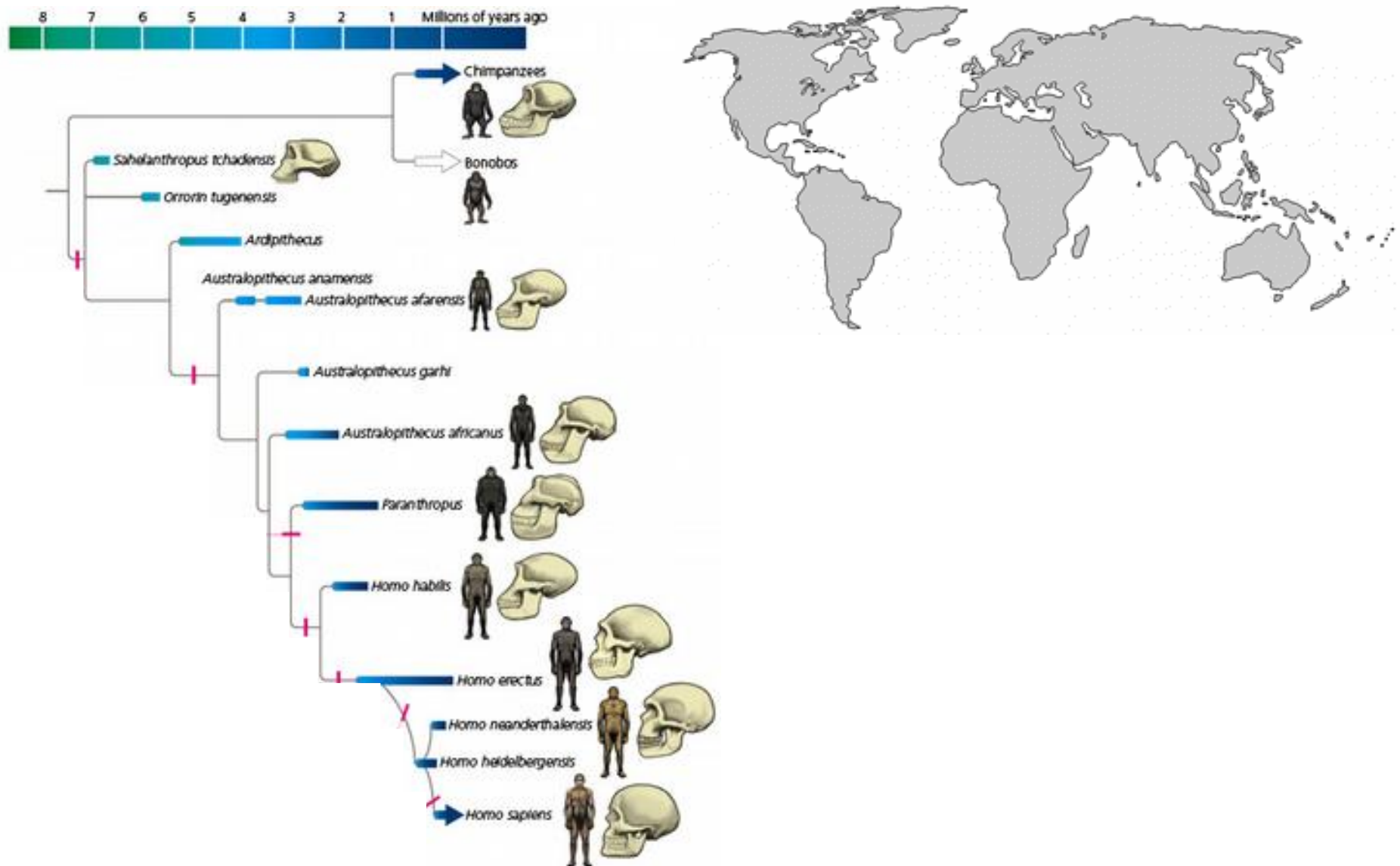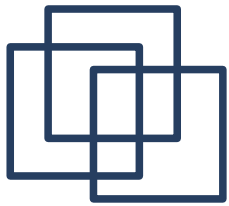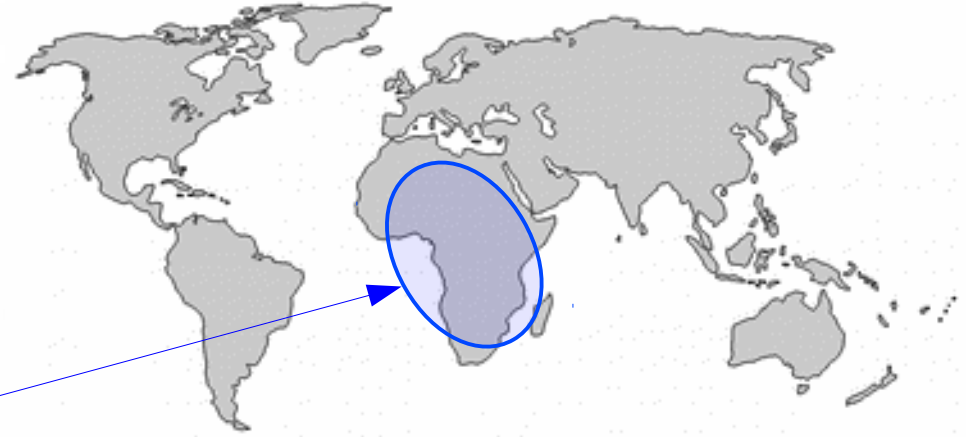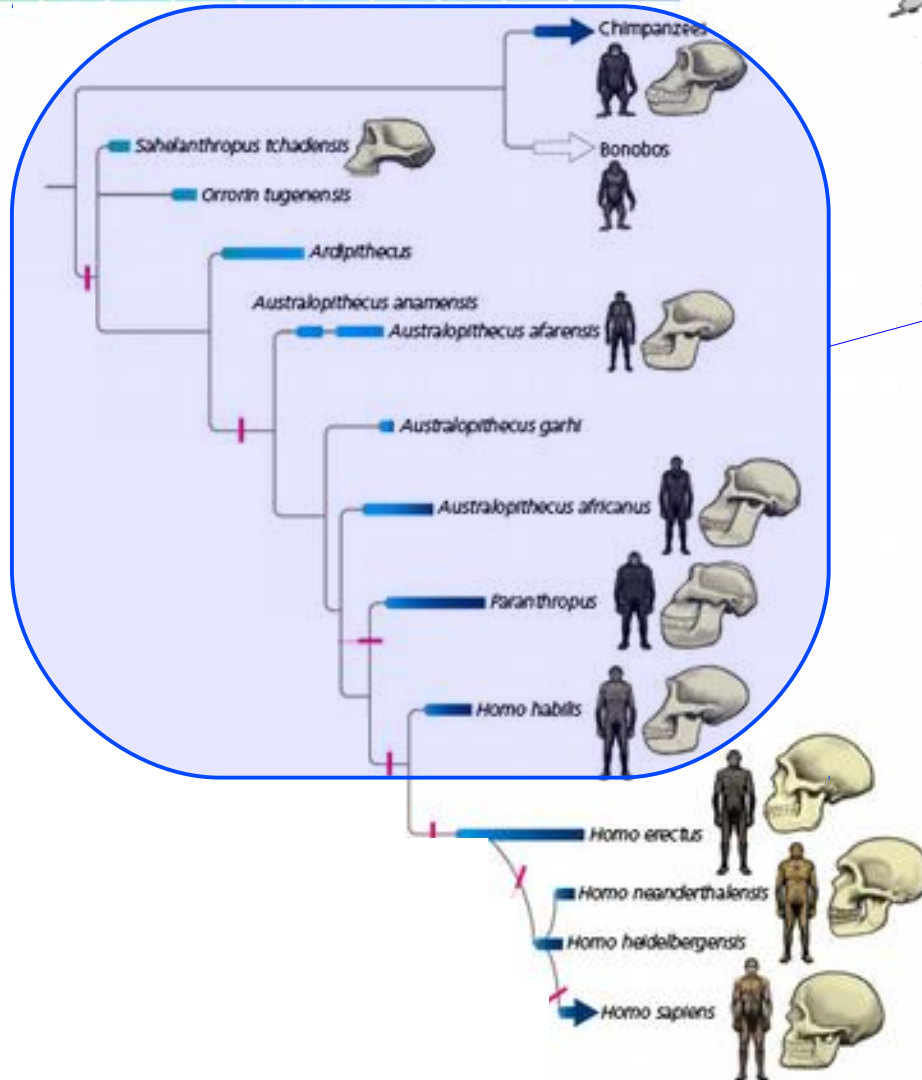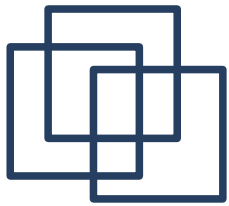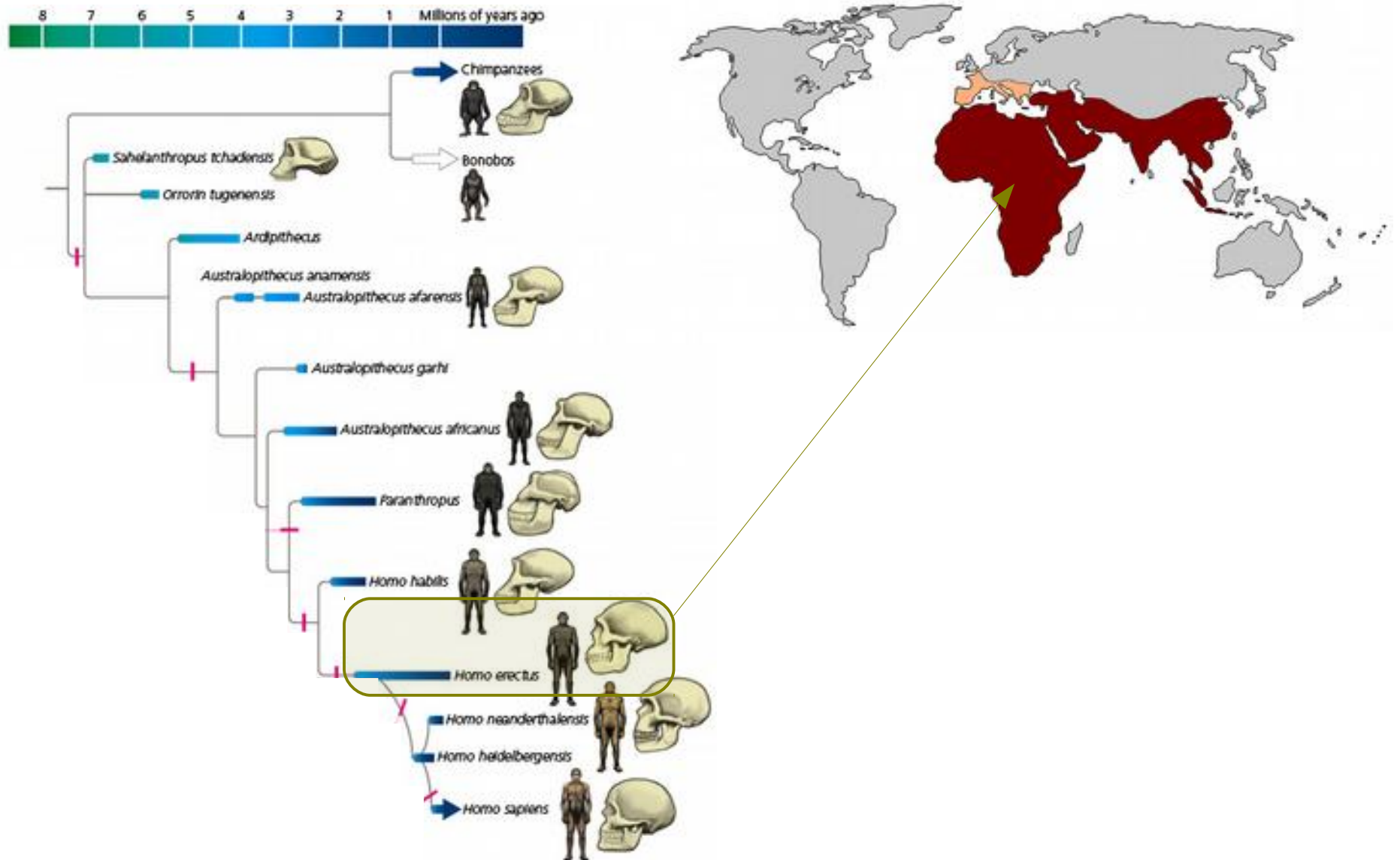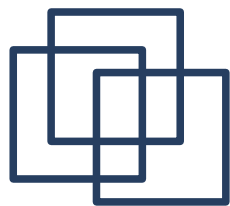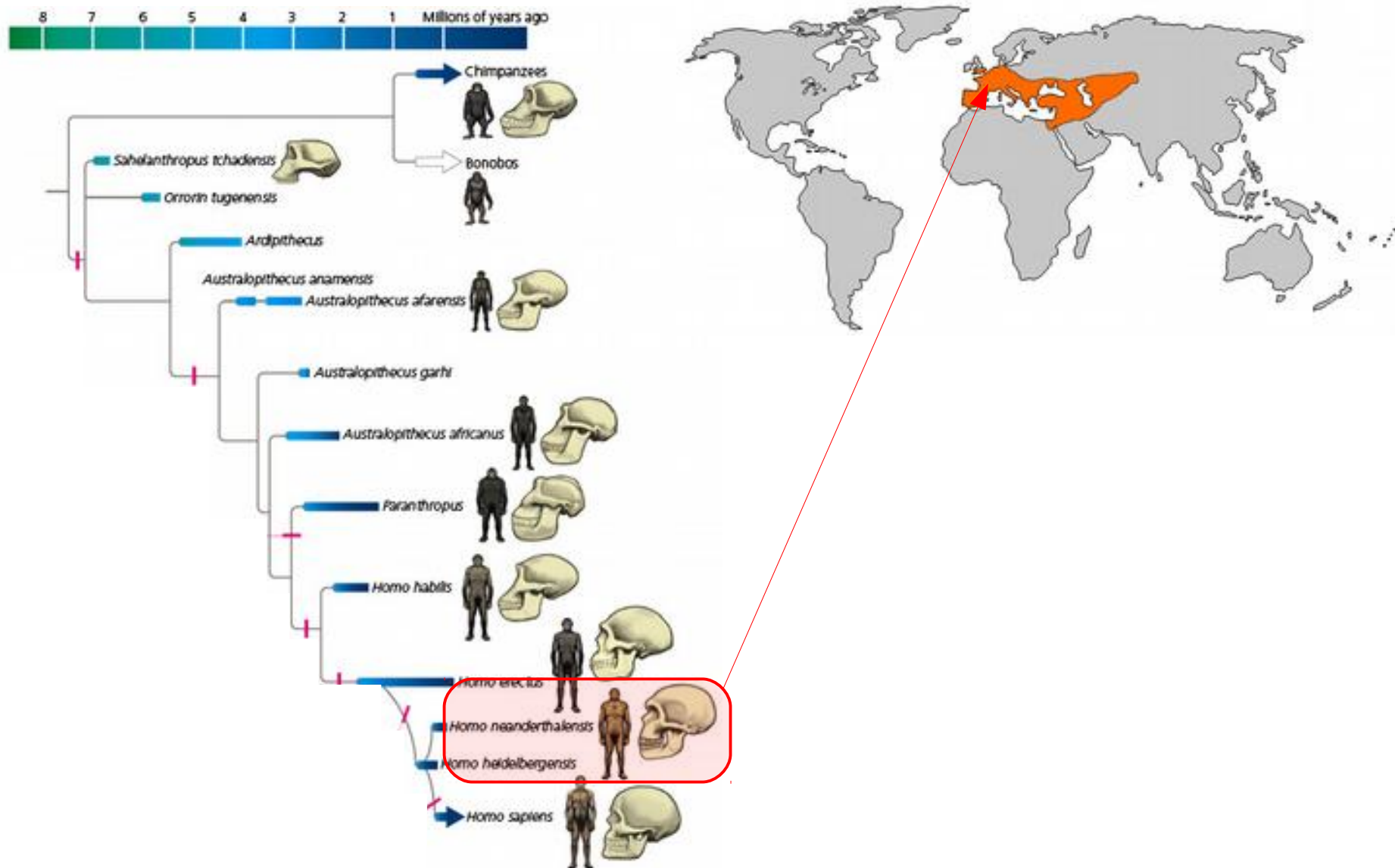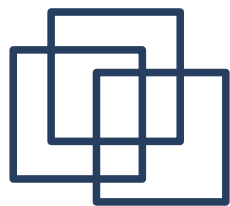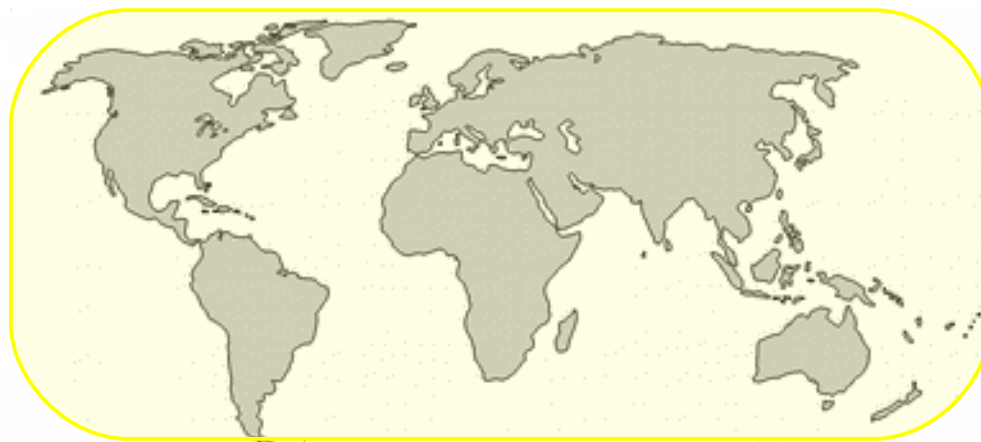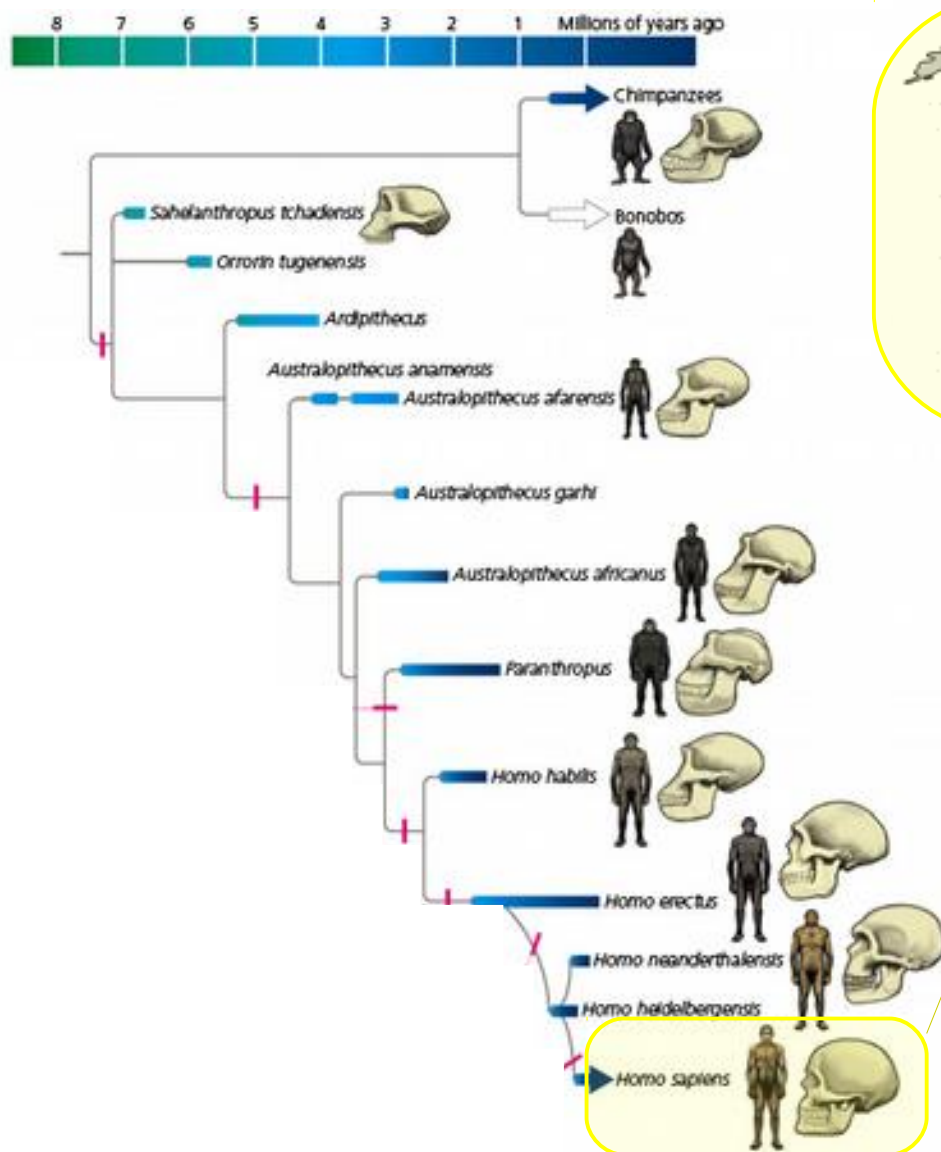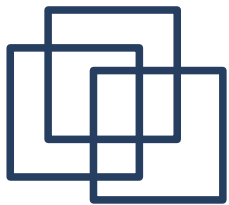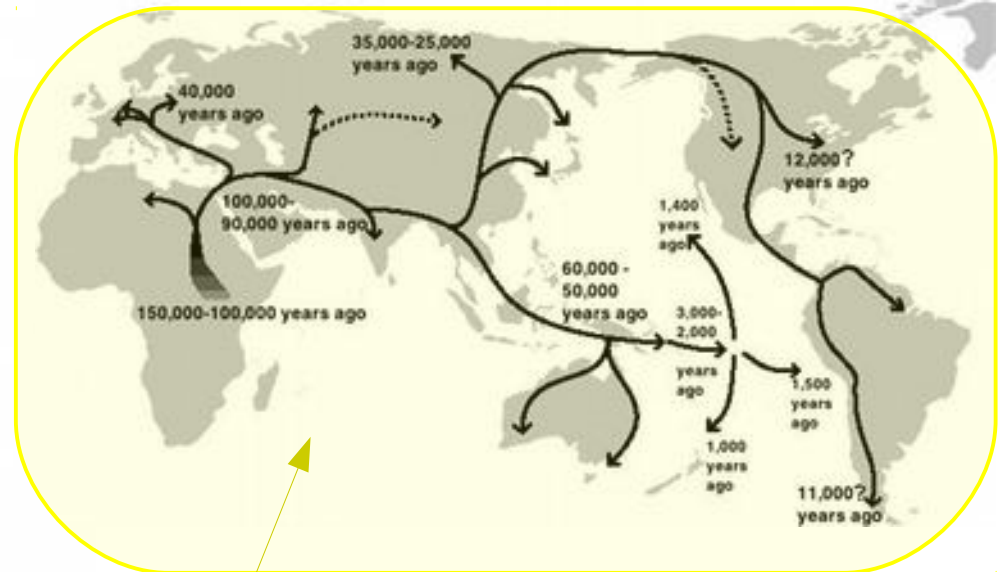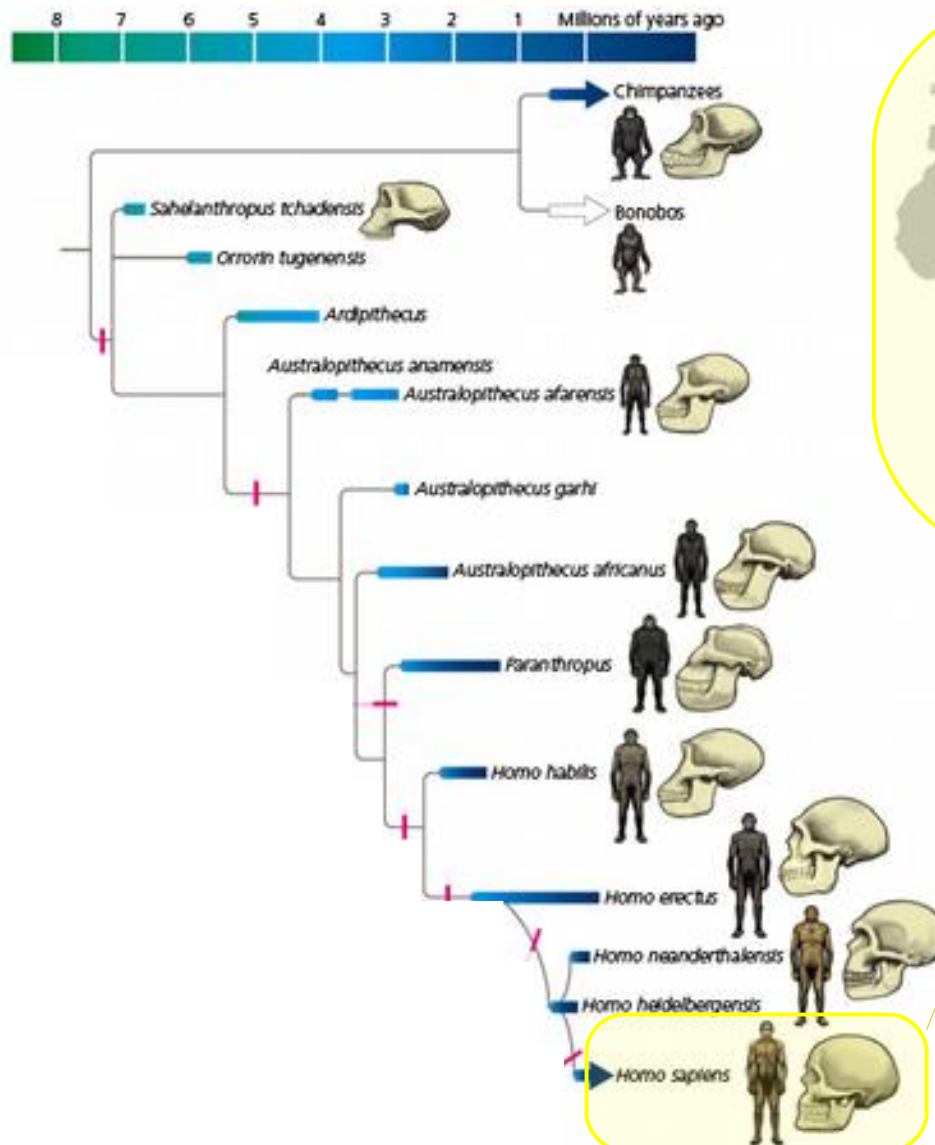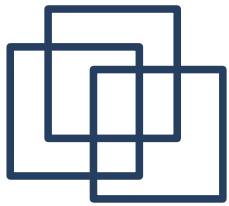
Human evolution
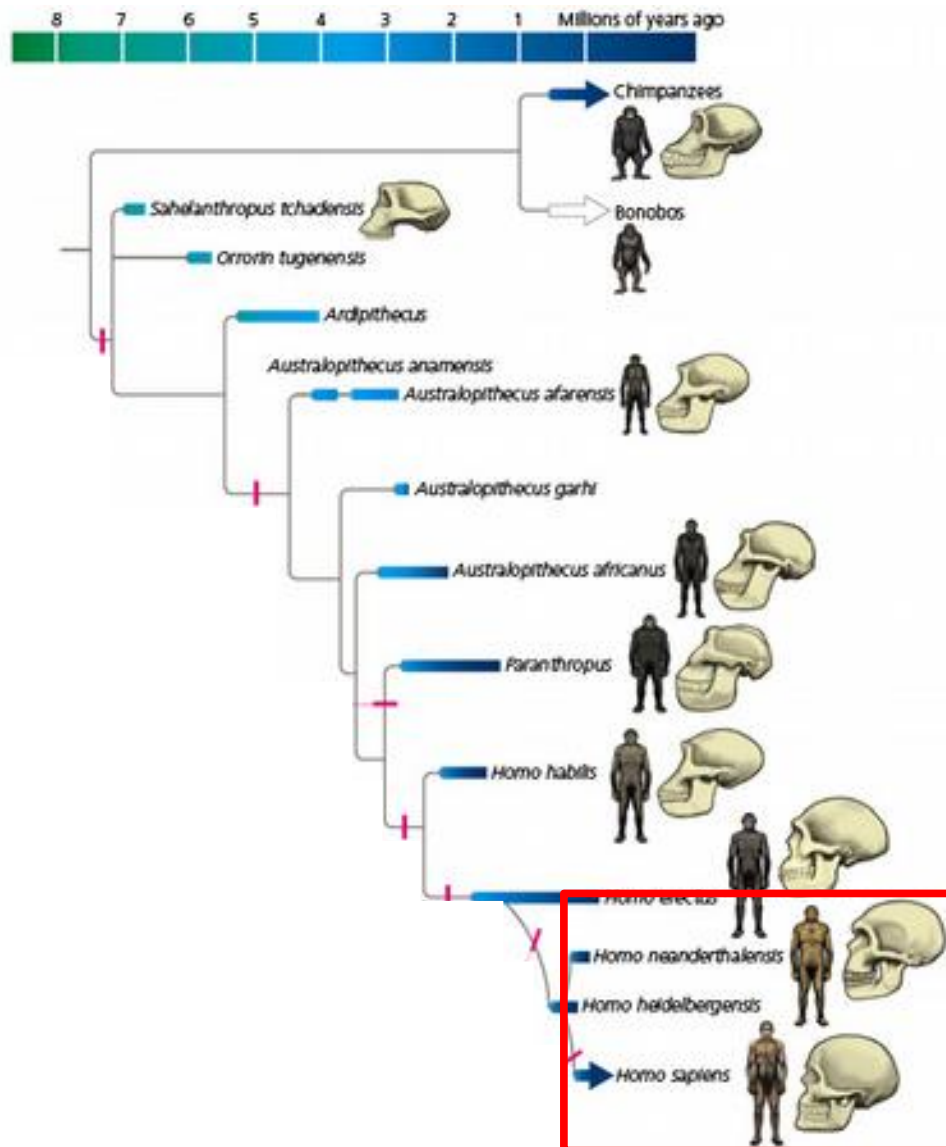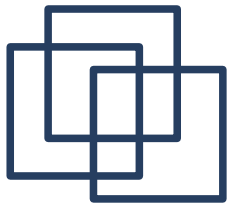
Human evolution

Human evolution
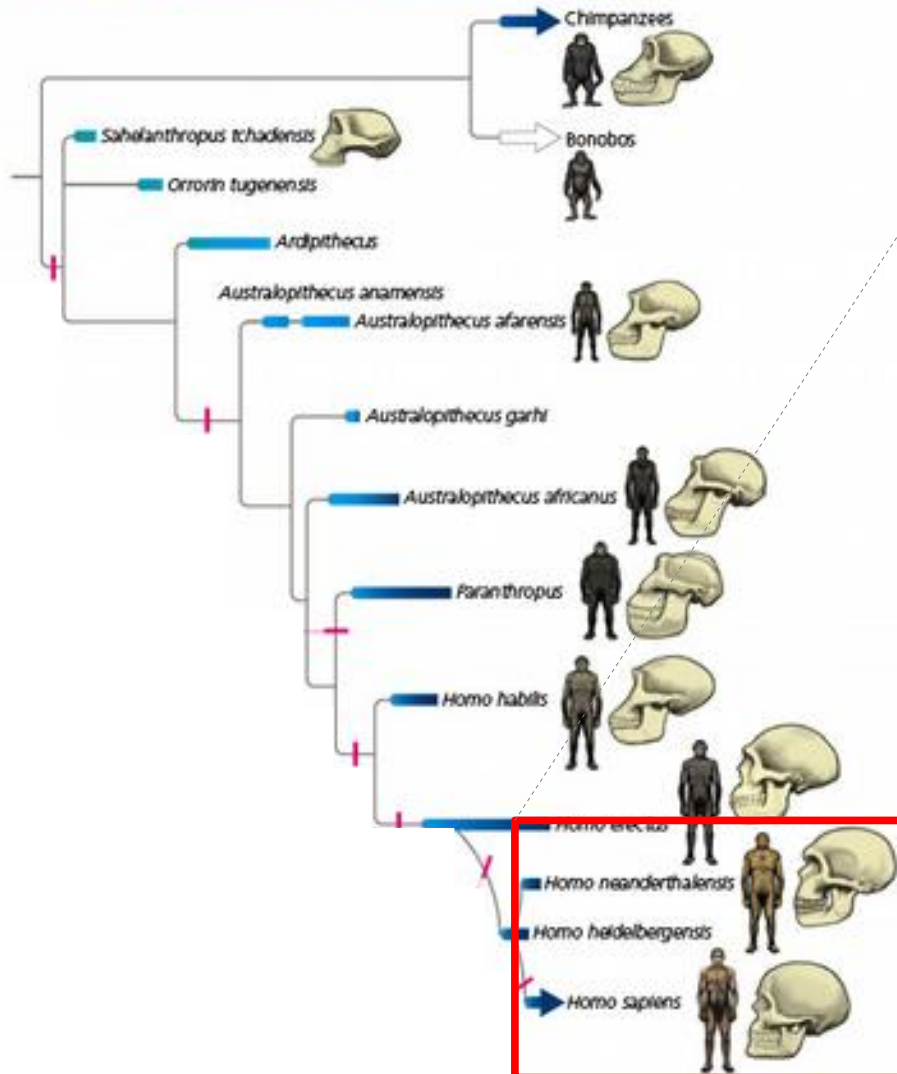
Human evolution

Human evolution
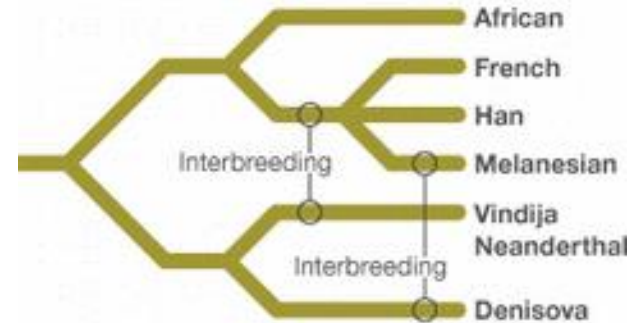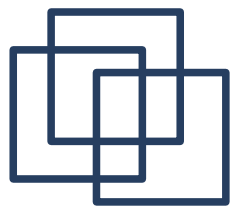
Human evolution
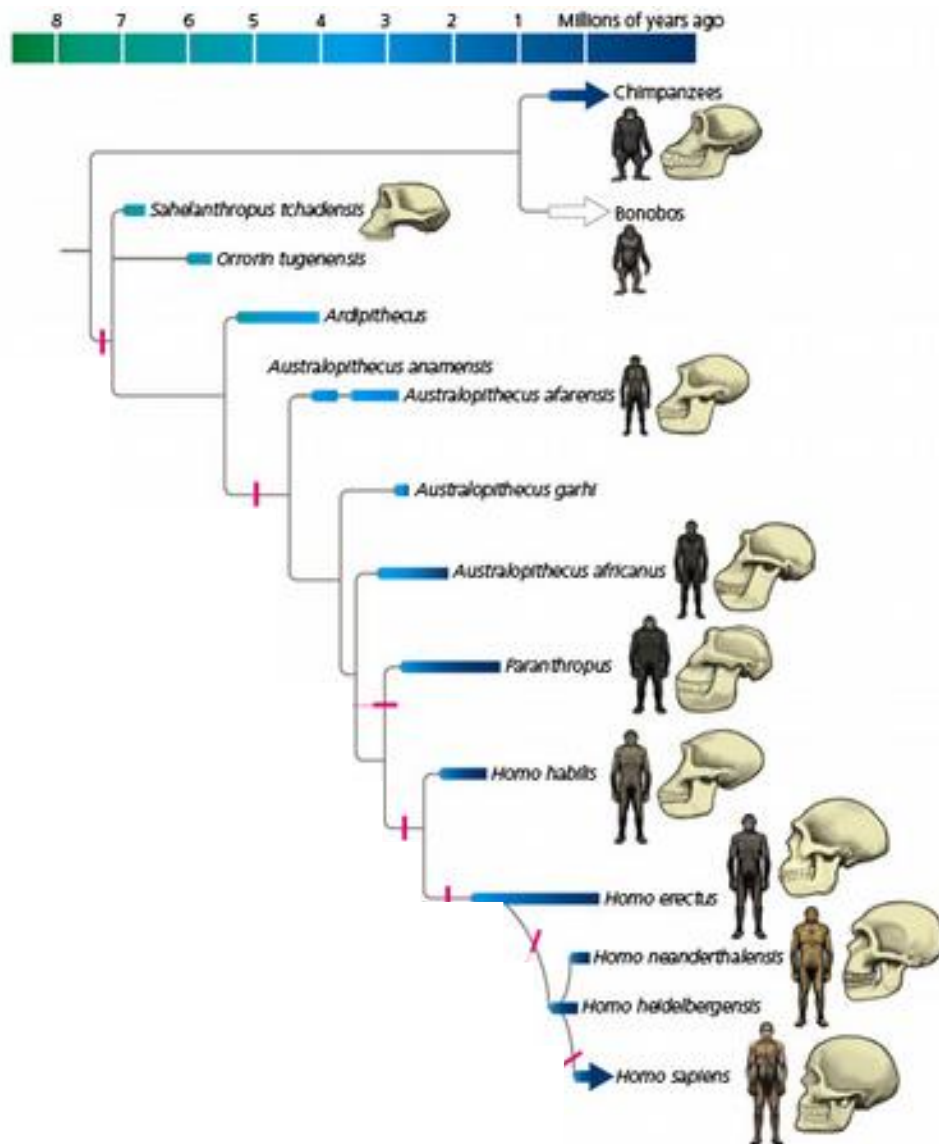
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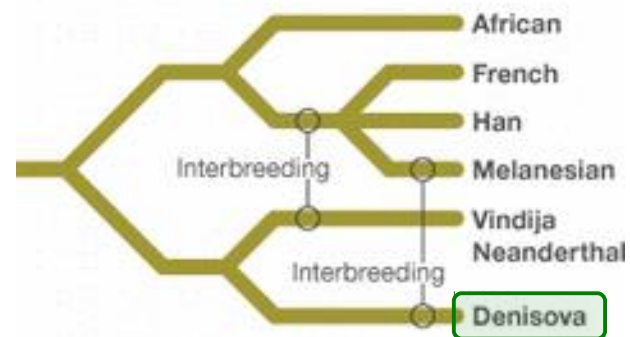
Human evolution
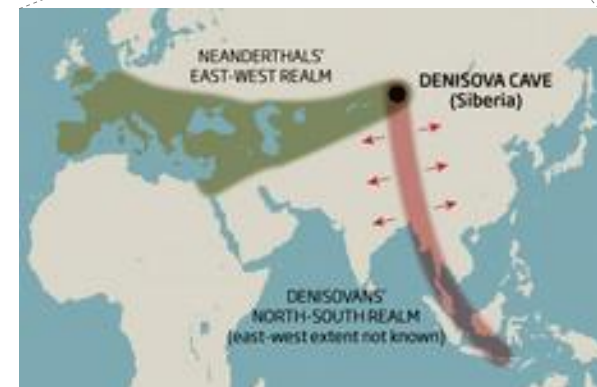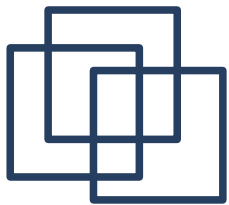
Human evolution
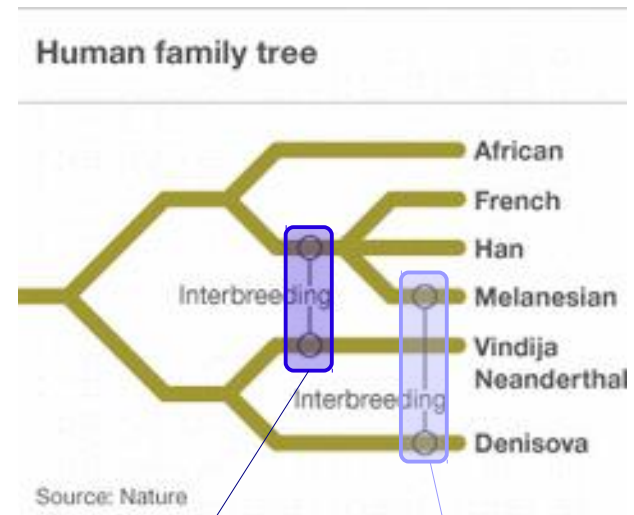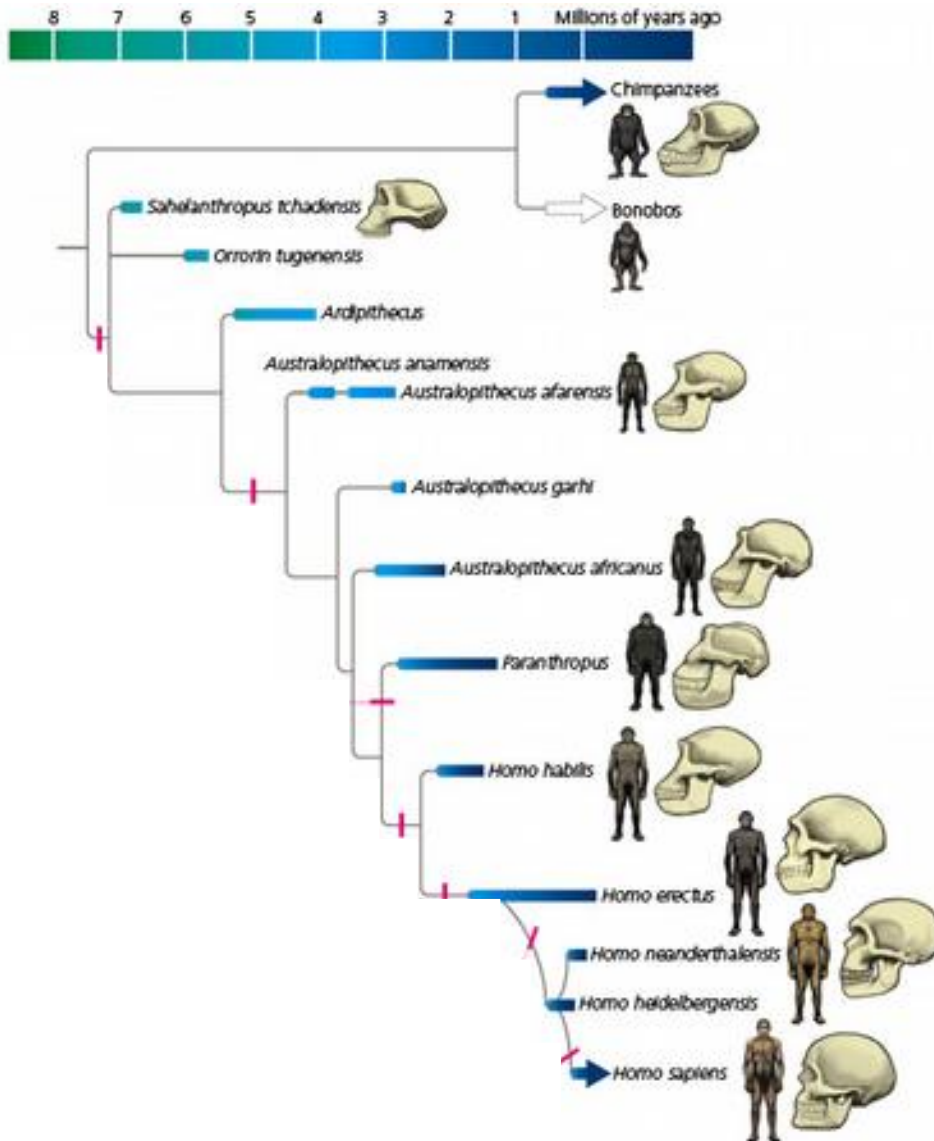
Human evolution

~4% Melanesians

1-4% (2.5%) non-Africans
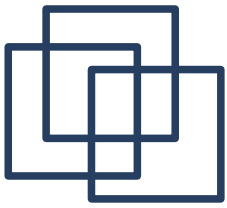
Human evolution

- "Modern" exon *FOXP2*
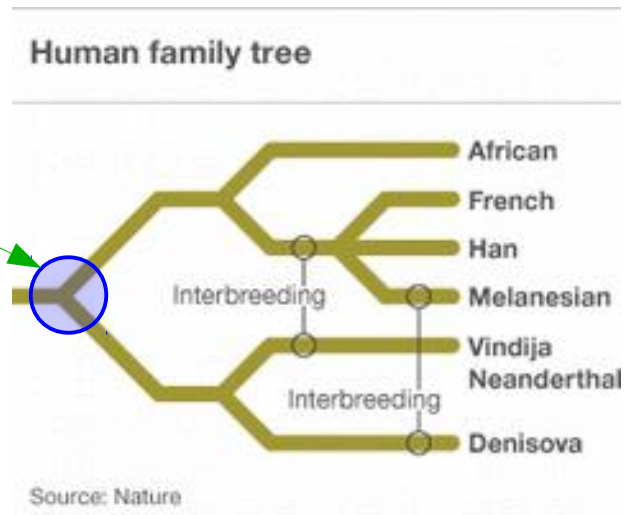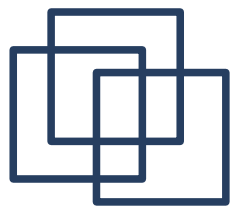
- Hyoid bone

- Audiograms

- Tools

- Symbolic culture (?)

=

Human family tree

African
French
Han
Melanesian
Vindija Neanderthal
Denisova

Interbreeding

Interbreeding

Source: Nature

Human evolution

- "Modern" exon *FOXP2*

- Hyoid bone

- Audiograms

- Tools

- Symbolic culture (?)

- Intron 8 *FOXP2*/*POU3F2*

- *CNTNAP2*

- *MCPH1*, *ASPM*, *DRD5*, *MEF2A*

- Survival

- Symbolic culture, tools...

Human family tree

African
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Han
Melanesian

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Neanderthal

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=

≠

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- "Modern" exon *FOXP2*
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$=$

- Intron 8 *FOXP2*/*POU3F2*
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Human family tree

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Interbreeding — Melanesian
Vindija Neanderthal
Interbreeding
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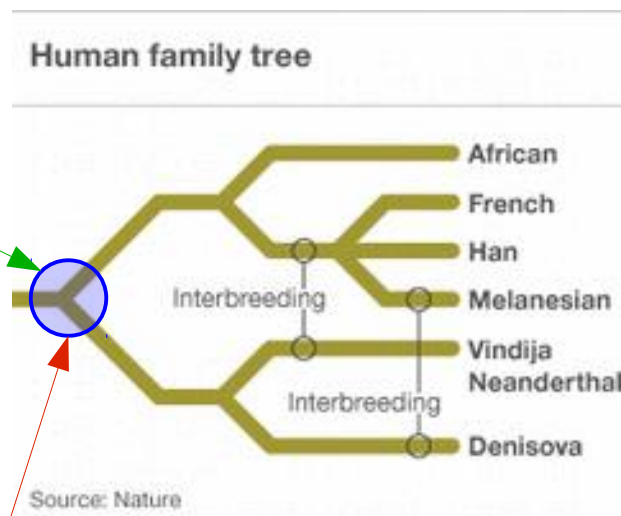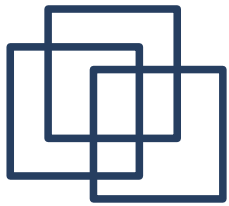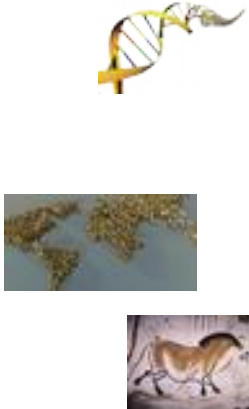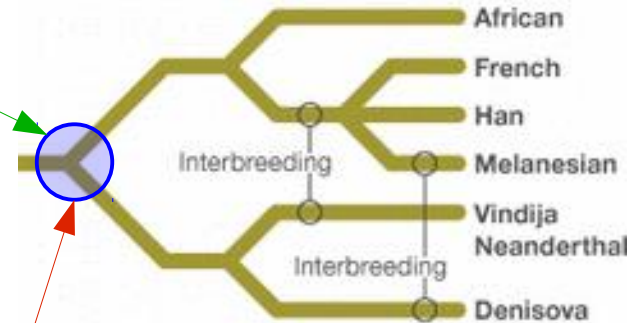
Source: Nature

Did "they" have language?
If so, what kind?
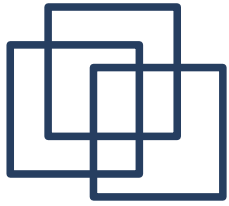
Human genetic and linguistic diversity

- Genetically quite similar to other animals:
    - 98% to chimps
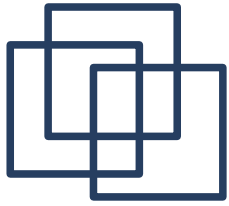
Human genetic and linguistic diversity

- Genetically quite similar to other animals:
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    - 85% to mice

Human genetic and linguistic diversity

- Genetically quite similar to other animals:

  - 98% to chimps

  - 85% to mice

  - 50% to fruit flies

98% 85%
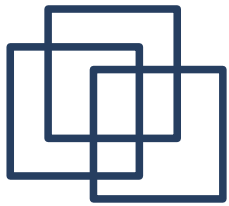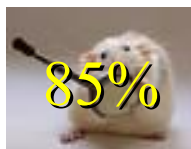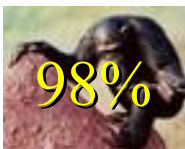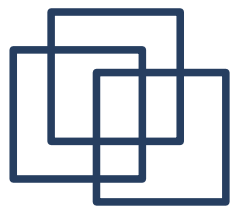
Human genetic and linguistic diversity

- Genetically quite similar to other animals:
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  - 85% to mice
  - 50% to fruit flies
- Most differences in regulatory networks and dynamics
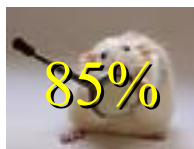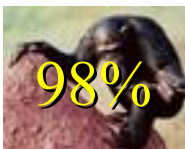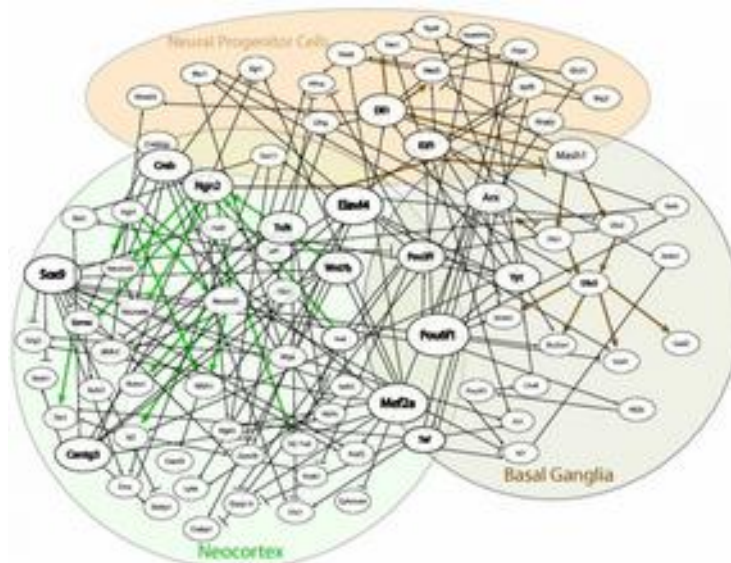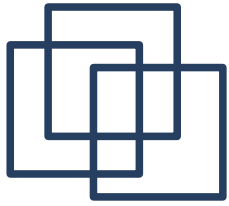
Human genetic and linguistic diversity

- Relatively genetically uniform species ← evolutionary history
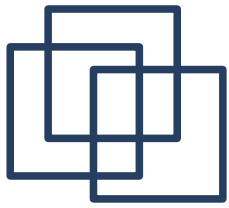
Human genetic and linguistic diversity

- Relatively genetically uniform species ← evolutionary history

- But we are no clones!

  ~ <0.5% nucleotide diversity b/w individuals (15 million bases)

Human genetic and linguistic diversity

- Relatively genetically uniform species ← evolutionary history

- But we are no clones!

- Distribution of diversity

  ~ 8% between : 7% within continents : 85% within local groups

Human genetic and linguistic diversity

- Relatively genetically uniform species ← evolutionary history

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  diversity decreases with distance from Africa

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phenotype

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phenotype genotype
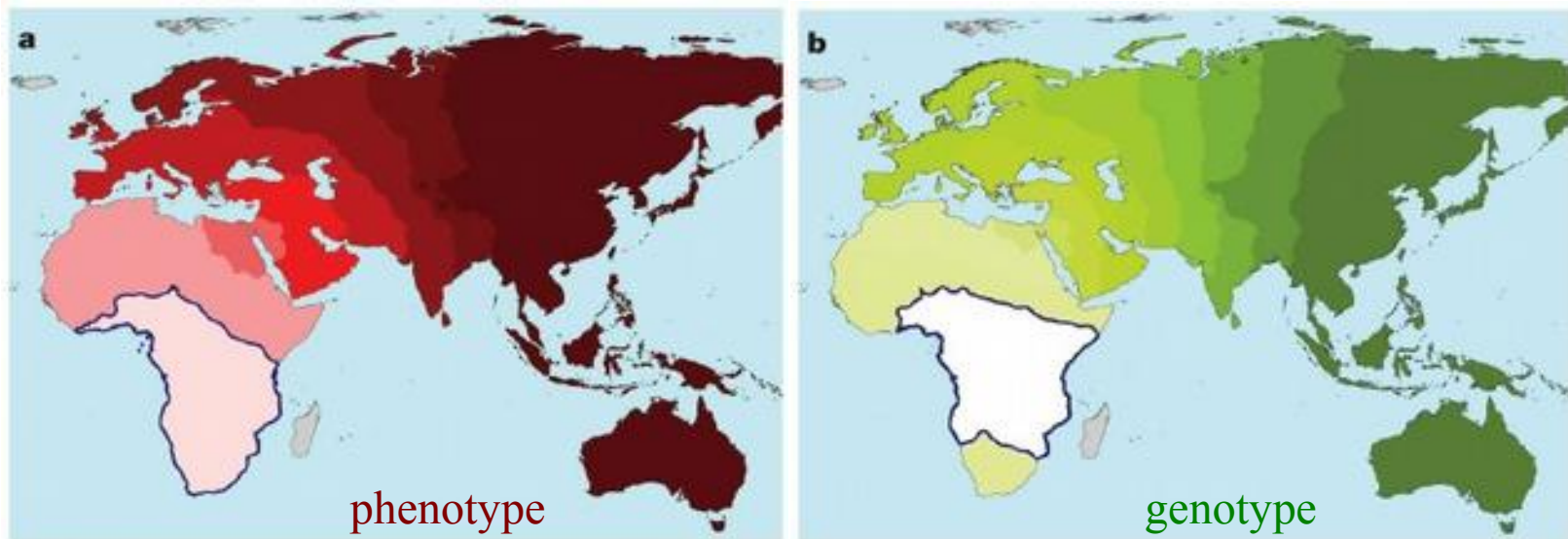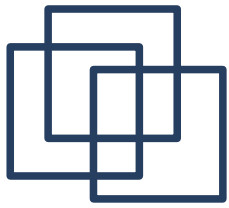
Human genetic and linguistic diversity

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  gradual clines as opposed to discontinuous boundaries
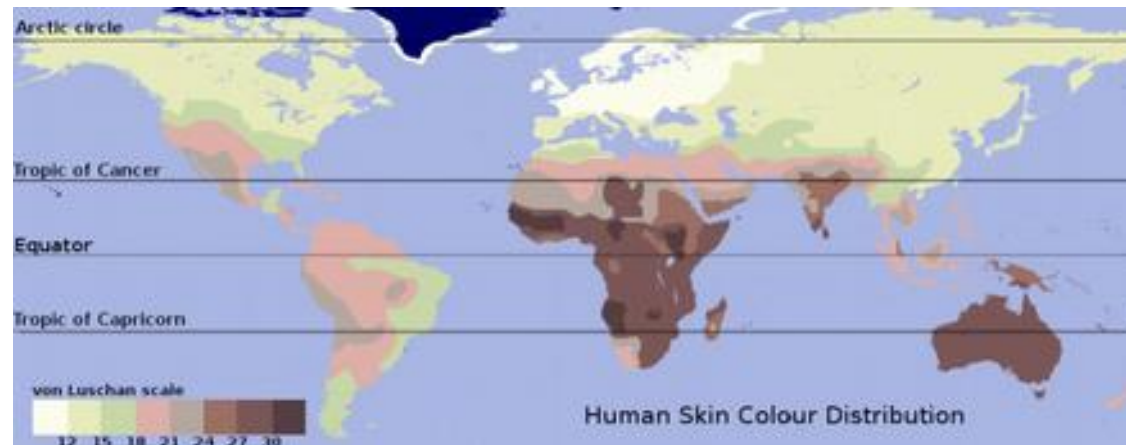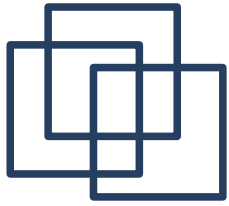
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  diversity decreases with distance from Africa

  gradual clines as opposed to discontinuous boundaries

  due to ubiquitous allele frequencies, less to private alleles:
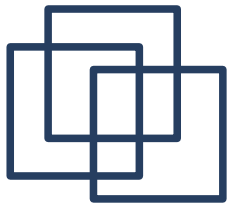
Human genetic and linguistic diversity
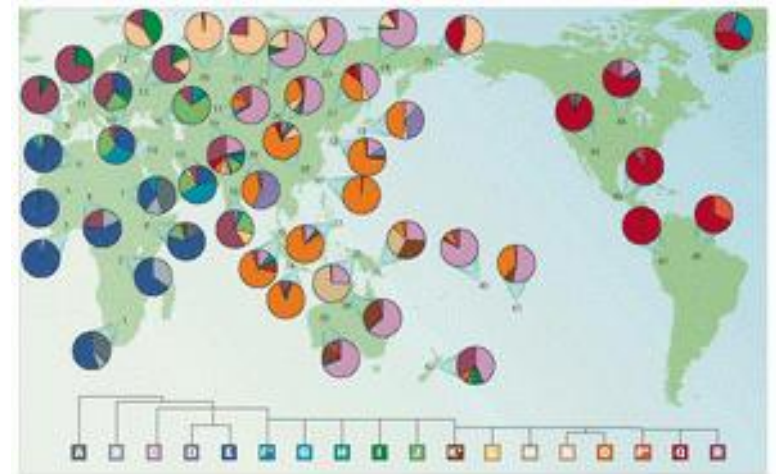
– continuous

Distribution of Y haplogroups
(Jobling & Tyler-Smith 2003)
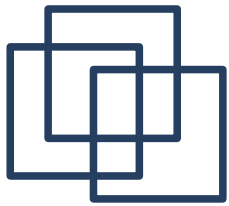
Human genetic and linguistic diversity

- continuous
- multiple loci
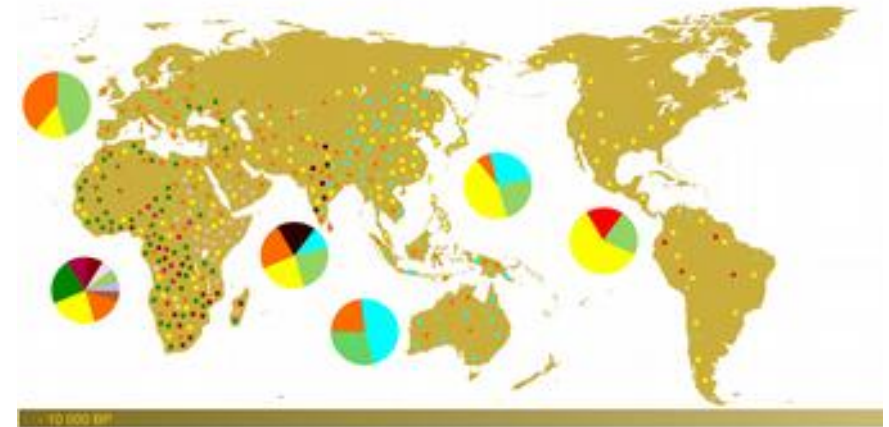
Human genetic diversity
(Barbujani & Colonna, 2010)
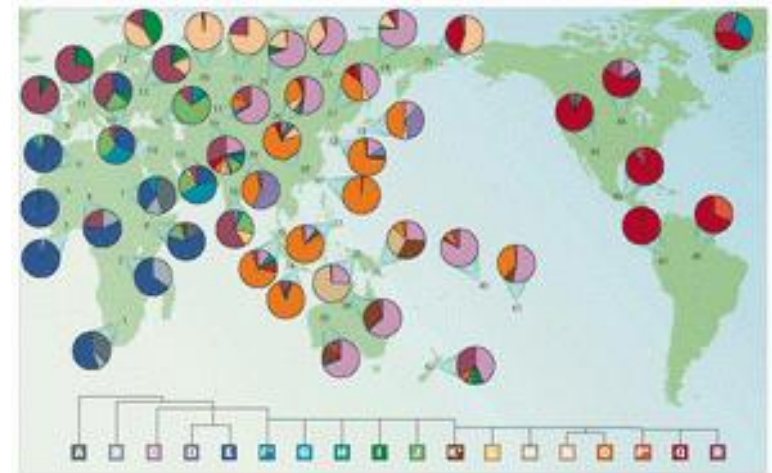
Distribution of Y haplogroups
(Jobling & Tyler-Smith 2003)

Human genetic and linguistic diversity

- continuous
- multiple loci
- geography

Human genetic diversity
(Barbujani & Colonna, 2010)

Genetic diversity Europe (500,000 SNPs)

Distribution of Y haplogroups
(Jobling & Tyler-Smith 2003)

Human genetic and linguistic diversity

- ~ **7000** languages
- > **100** language families
  - ~ **150** incl. language isolates

Human genetic and linguistic diversity

- ~ **7000** languages
- > **100** language families
  - ~ **150** incl. language isolates

- Language = **cultural evolutionary system**

- **vertical** inheritance → language families
- **horizontal** processes → language areas

Indian sprachbund
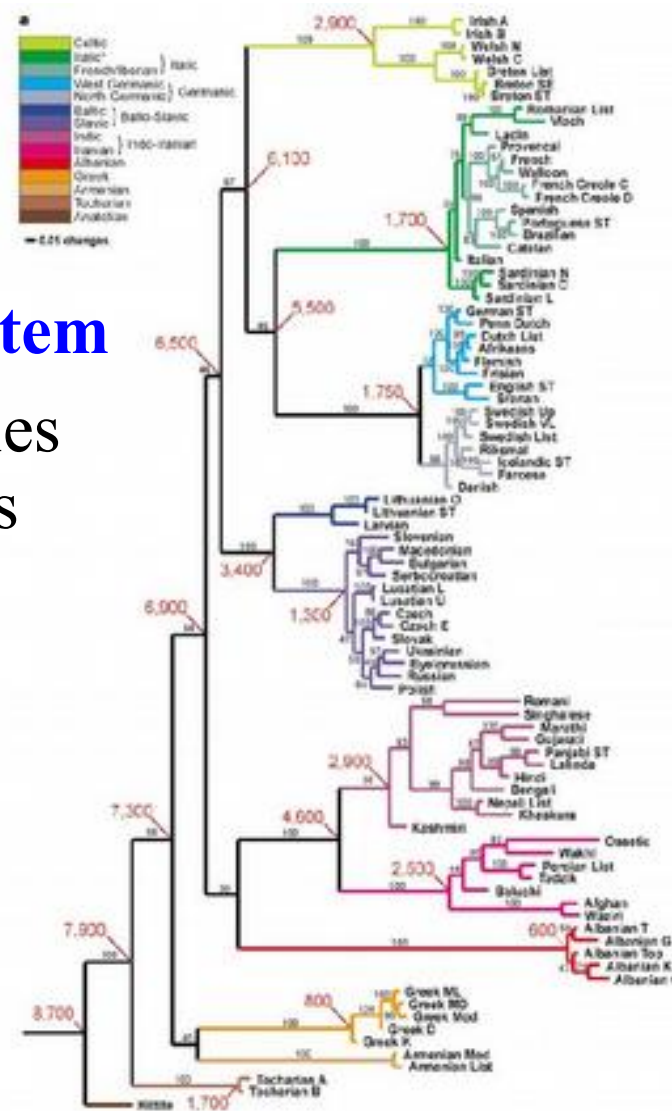
IE tree with ages, Gray & Atkinson (2003)
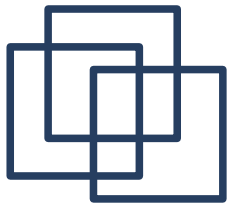
Human genetic and linguistic diversity

- Humans carry genes & languages
  → **historical processes** should create or break correlations

- Migrations

Viking longboat
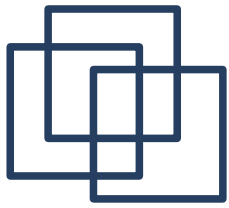
Human genetic and linguistic diversity

- Humans carry genes & languages
  → **historical processes** should create or break correlations

- Migrations
- Conquest

Roman legion
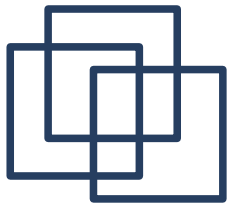
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Sultan Ahmed Mosque, Istambul
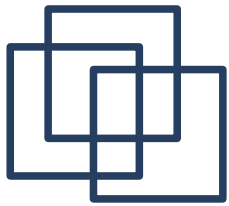
Human genetic and linguistic diversity

- Farming/language co-dispersal
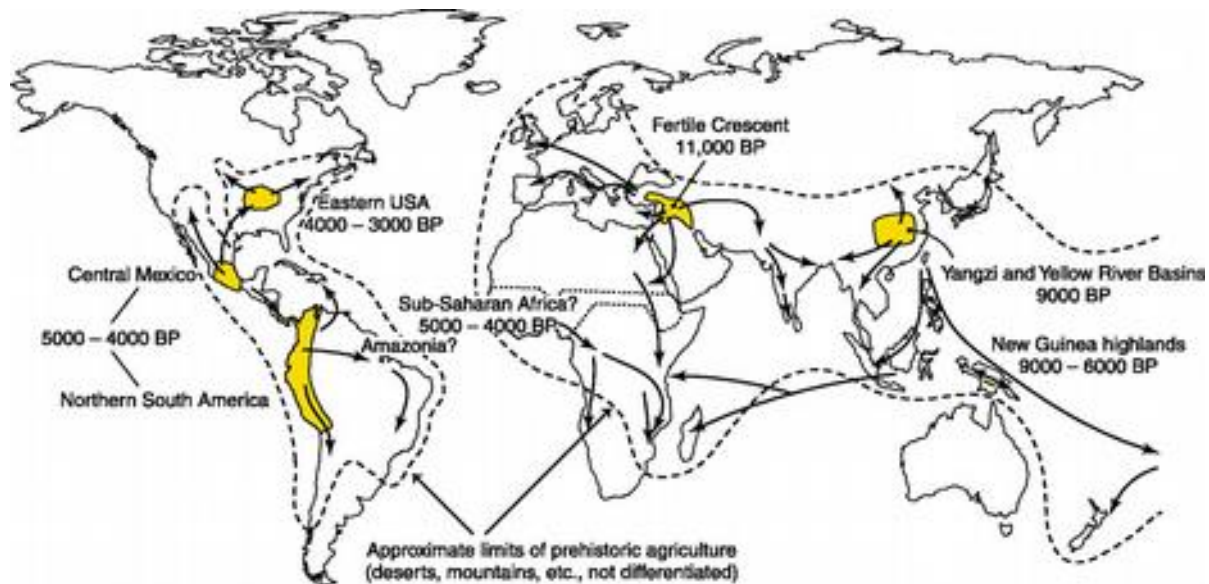
Proposed farming expansions Diamond & Bellwood (2003)
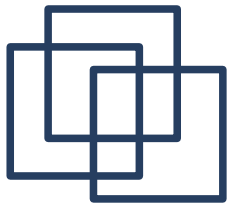
Human genetic and linguistic diversity
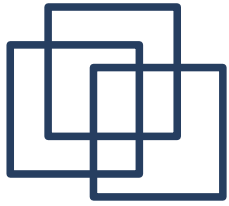
→ accidental correlations

Human genetic and linguistic diversity

→ accidental correlations
→ non-accidental correlations:

Human genetic and linguistic diversity

→ accidental correlations

→ non-accidental correlations:

• village sign languages

ABSL
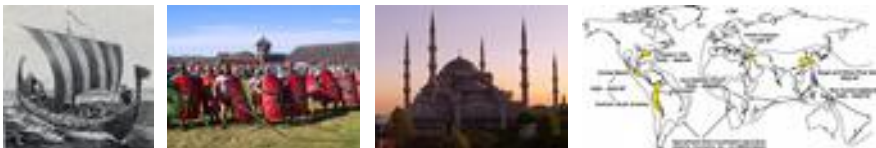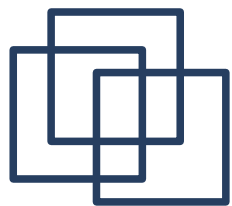
Kata Kolok
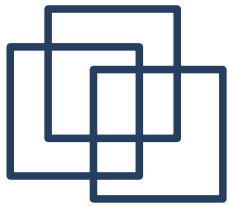
Human genetic and linguistic diversity

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→ non-accidental correlations:

- village sign languages
- tone, *ASPM*, *Microcephalin*

http://wals.info
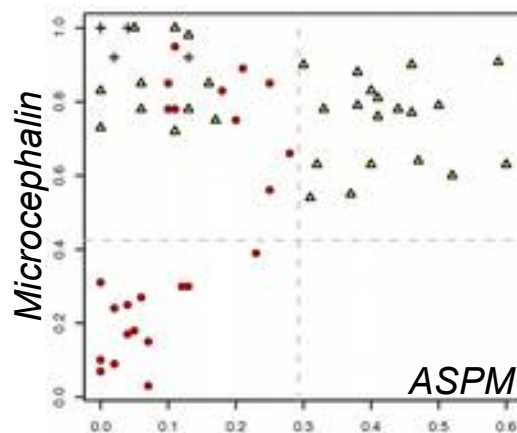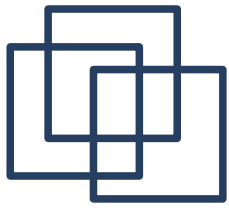
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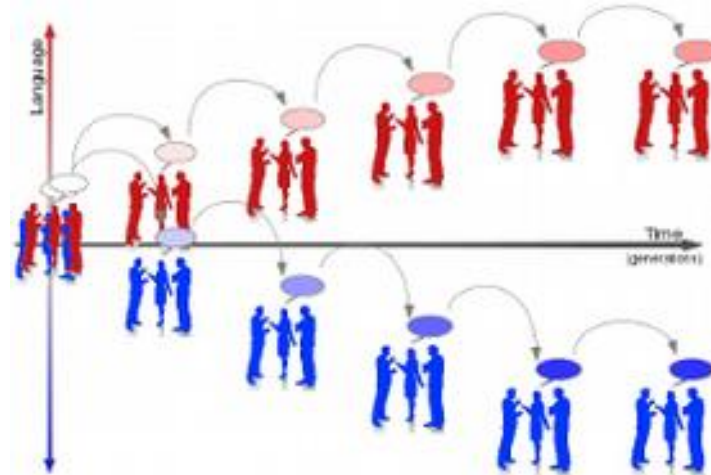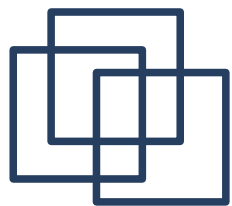
Niche construction & Gene-culture co-evolution
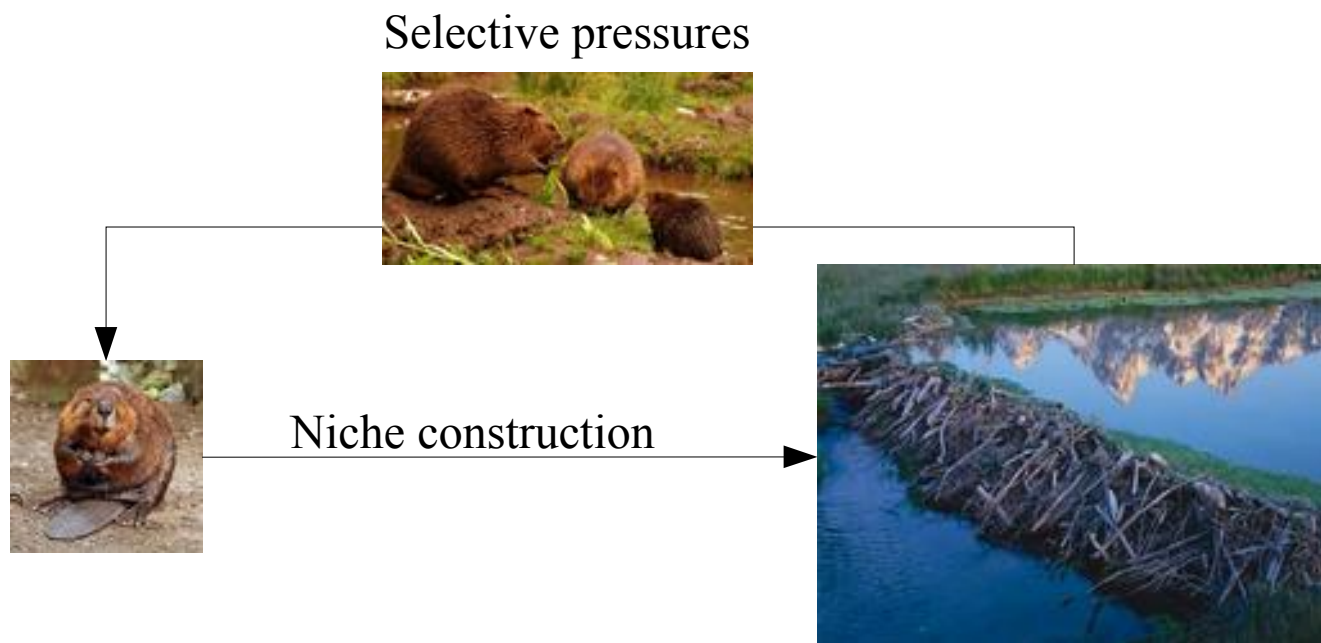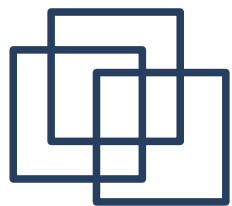
- Organisms construct their niches

Selective pressures

Niche construction

Niche construction & Gene-culture co-evolution

- Organisms construct their niches

- Culture shapes selective pressures:
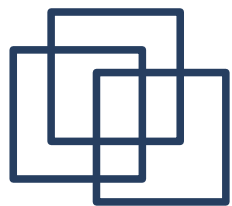
  - farming

Niche construction & Gene-culture co-evolution

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- Culture shapes selective pressures:

    - farming → immune system
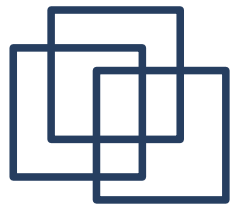
Niche construction & Gene-culture co-evolution

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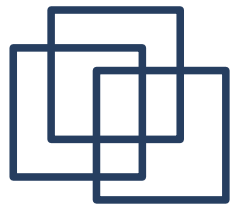
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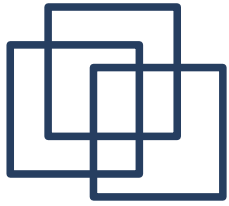
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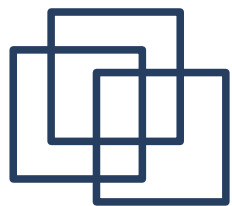
    - tool use → hand

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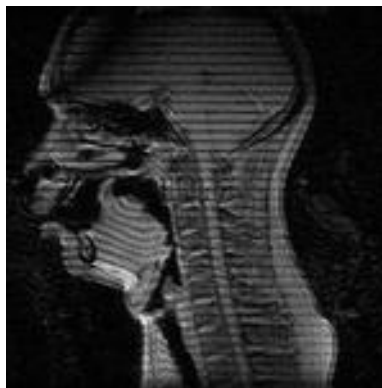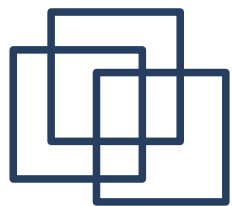
    – speech → vocal tract

Niche construction & Gene-culture co-evolution

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    - tool use → hand

    - speech → vocal tract

    - language → whole world
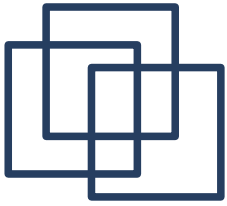
Conclusions

- Human evolution was really complicated → no simple story

- Complex relationships between genes and language

  → co-evolution and niche construction

- Ongoing evolution of the genetic architecture for language?

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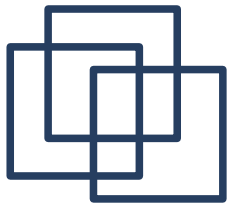
Suggested reading

- ## Books:
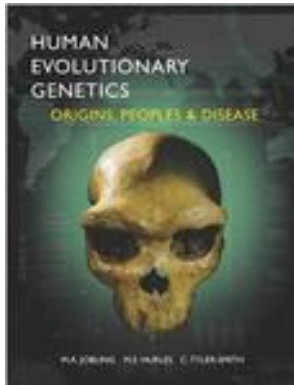
Jobling, M. A., Hurles, M., & Tyler-Smith, C. (2004). *Human Evolutionary Genetics: Origins, Peoples and Disease*. Garland Science: NY.
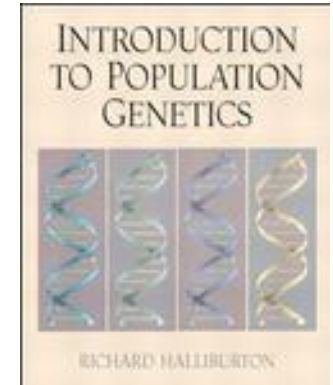
Halliburton, R. (2004). *Introduction to population genetics*. Pearson Eduction Inc.: Upper Saddle River.

- ## Population and evolutionary genetics:

Hurst, L. D. (2009). Fundamental concepts in genetics: genetics and the understanding of selection. *Nat Rev Genet* **10**:83–93. doi:10.1038/nrg2506

Charlesworth, B. (2009). Fundamental concepts in genetics: effective population size and patterns of molecular evolution and variation. *Nat Rev Genet* **10**:195–205. doi:10.1038/nrg2526

Jablonka, E., & Raz, G. (2009). Transgenerational epigenetic inheritance: prevalence, mechanisms, and implications for the study of heredity and evolution. *The Quarterly Review of Biology* **84**:131–176.

Koonin, E. V. (2009). Darwinian evolution in the light of genomics. *Nucleic Acids Res* **37**:1011–1034. doi:10.1093/nar/gkp089
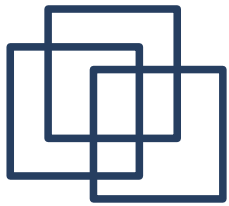
Suggested reading

- ## Human evolution:

Green, R. E., Krause, J., Briggs, A. W., Maricic, T., Stenzel, U., Kircher, M., … Fritz, M. H.-Y. (2010). A draft sequence of the Neandertal genome. *Science* **328**:710–722. doi:10.1126/science.1188021

Disotell, T. R. (2012). Archaic human genomics. *American Journal of Physical Anthropology* **149**:24–39. doi:10.1002/ajpa.22159

Meyer, M., Kircher, M., Gansauge, M.-T., Li, H., Racimo, F., Mallick, S., … Pääbo, S. (2012). A High-Coverage Genome Sequence from an Archaic Denisovan Individual. *Science* **338**:222–226. doi:10.1126/science.1224344

Dediu, D., & Levinson, S. C. (2013). On the antiquity of language: the reinterpretation of Neandertal linguistic capacities and its consequences. *Frontiers in Language Sciences* **4**:397. doi:10.3389/fpsyg.2013.00397

- ## Human diversity:

Barbujani, G., & Colonna, V. (2010). Human genome diversity: frequently asked questions. *Trends in Genetics* **26**:285–295. doi:10.1016/j.tig.2010.04.002

Novembre, J., Johnson, T., Bryc, K., Kutalik, Z., Boyko, A. R., Auton, A., … Bustamante, C. D. (2008). Genes mirror geography within Europe. *Nature* **456**:98–101. doi:10.1038/nature07331
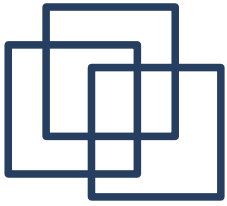
Suggested reading

- ## Linguistic and genetic diversity:

  Diamond, J., & Bellwood, P. (2003). Farmers and their languages: the first expansions. *Science* **300**:597–603.

  Dediu, D. (2011). Are languages really independent from genes? If not, what would a genetic bias affecting language diversity look like? *Hum Biol* **83**:279–296. doi:10.3378/027.083.0208

- ## Gene-culture co-evolution/niche construction:

  Laland, K. N., Odling-Smee, J., & Myles, S. (2010). How culture shaped the human genome: bringing genetics and the human sciences together. *Nat Rev Genet* **11**:137–148. doi:10.1038/nrg2734

  Fisher, S. E., & Ridley, M. (2013). Culture, Genes, and the Human Revolution. *Science* **340**:929–930. doi:10.1126/science.1236171

  Senghas, A. (2005). Language emergence: clues from a new bedouin sign. *Curr Biol* **15**:R463–R465. doi:10.1016/j.cub.2005.06.018