

The New Field of

Evo-Devo

How evolution re-wires genomes
to re-program development
to re-configure anatomy.

Lewis Held
Texas Tech



Outline

Anatomy \approx Geometry

Cellular Programming

Constraints on Evolution



Human Evolution

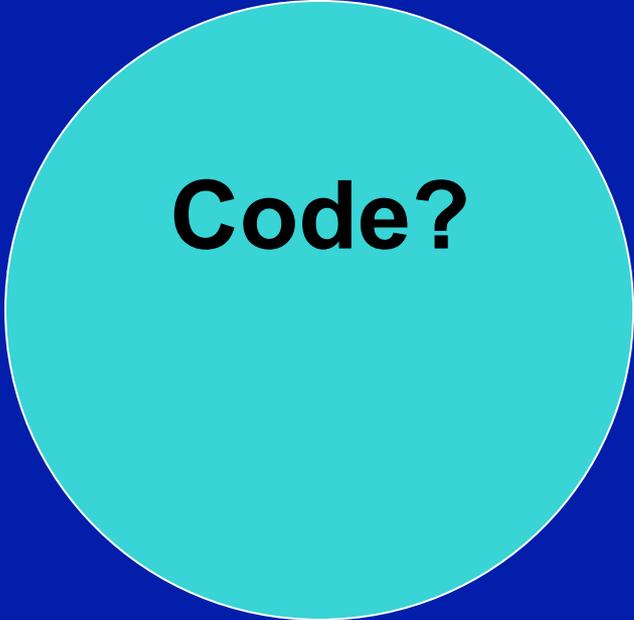
Why are we so naked?

How did we get so brainy?

Anatomy =
Coding of cells in the body

Geometry =
Coding of points in space

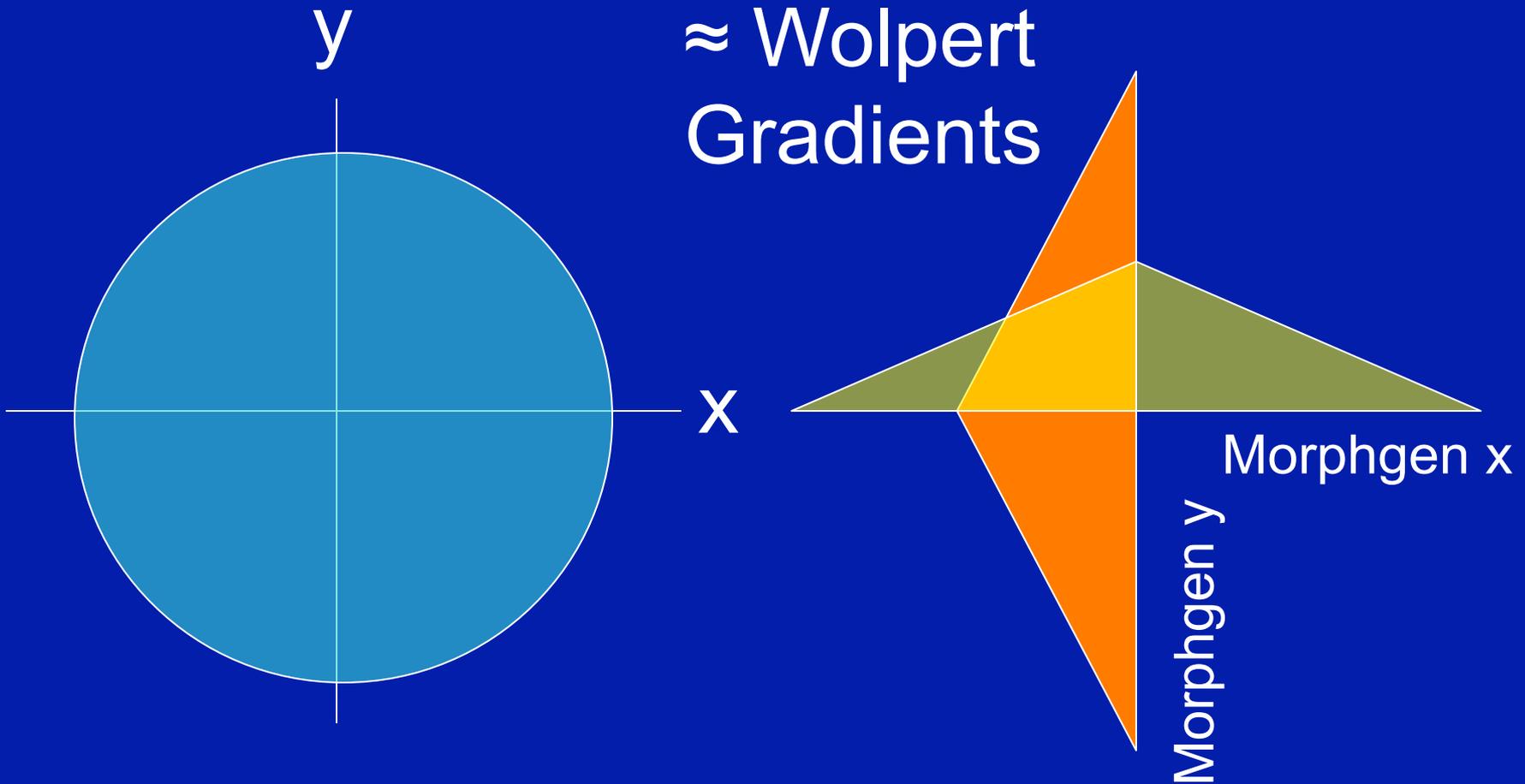
Example:
Circle



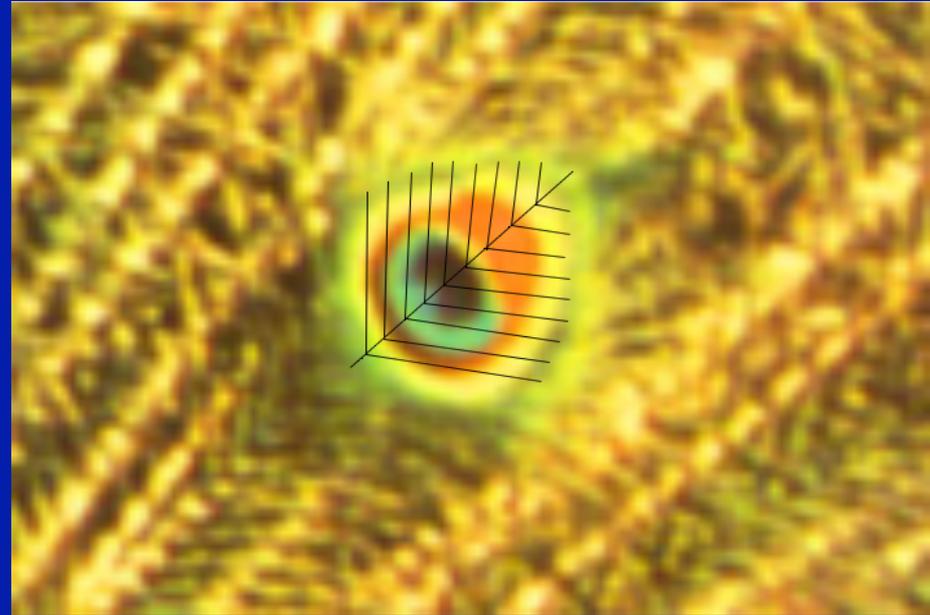
Code?

Code = Cartesian Coordinates?

Plot all (x,y) , where $x^2 + y^2 = r^2$.

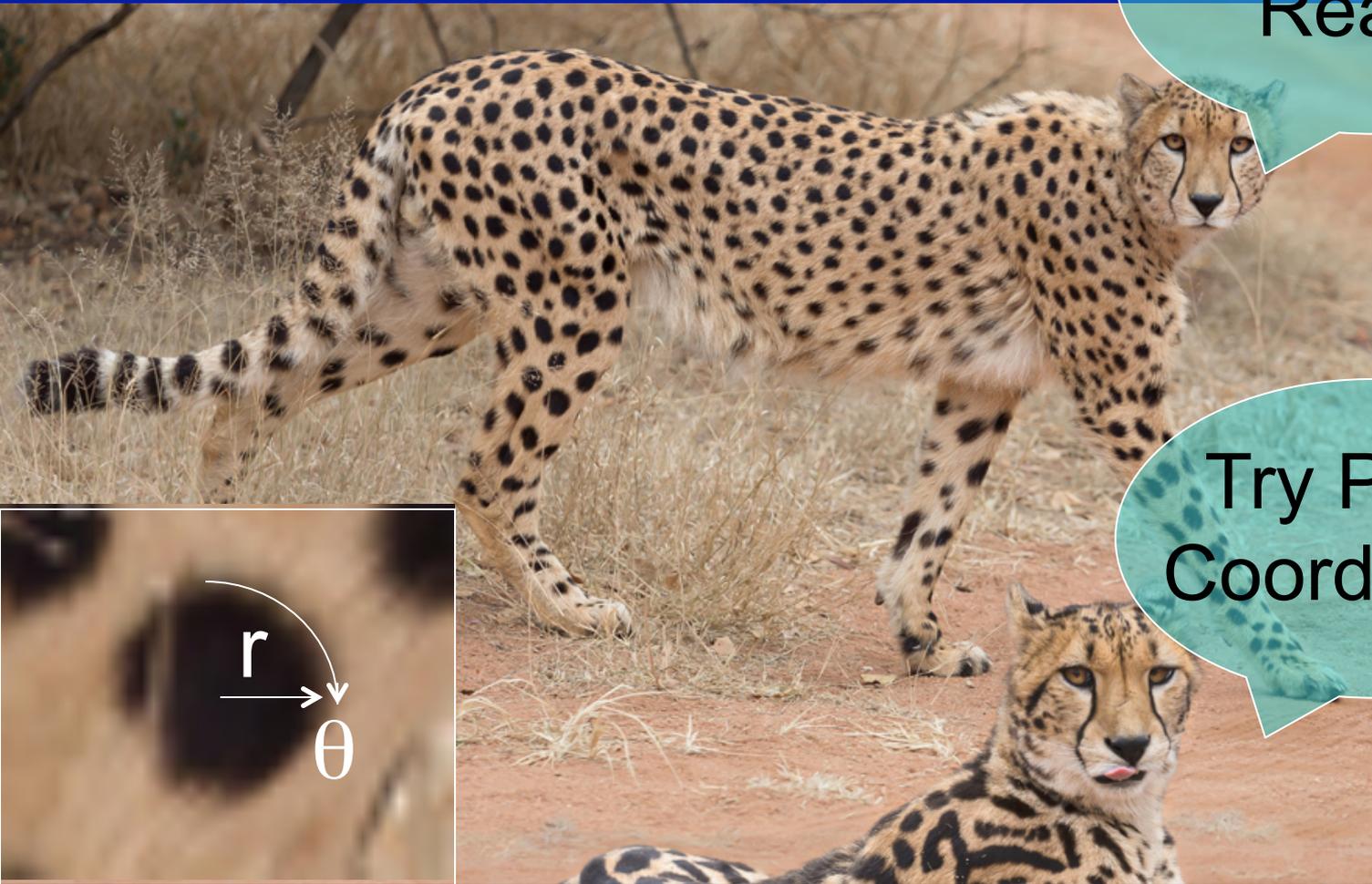


Hmmm ...



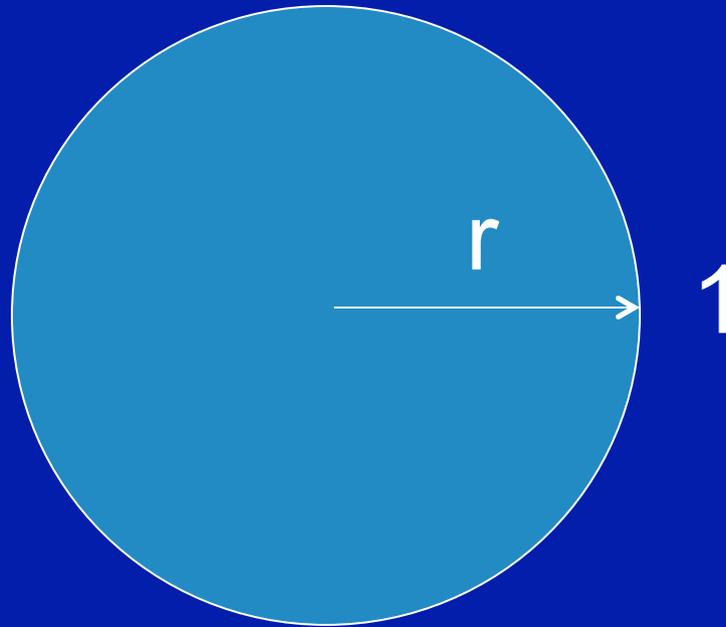
Cartesian?
Really?

Try Polar
Coordinates!



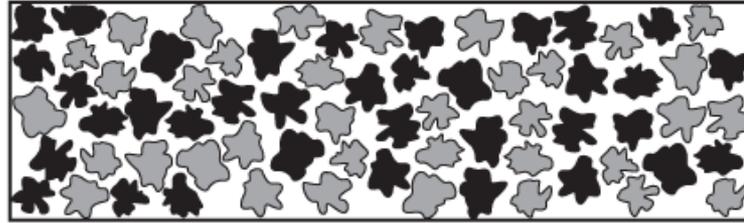
Code = Polar Coordinates?

Plot all (r, θ) , where $r < 1$.

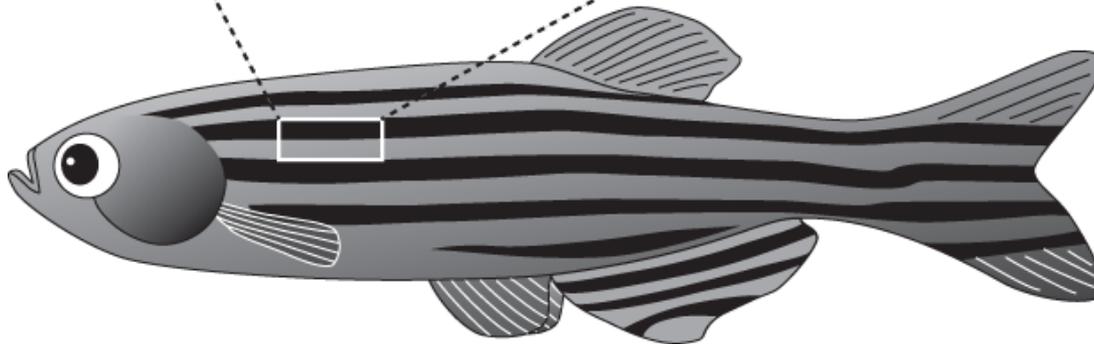
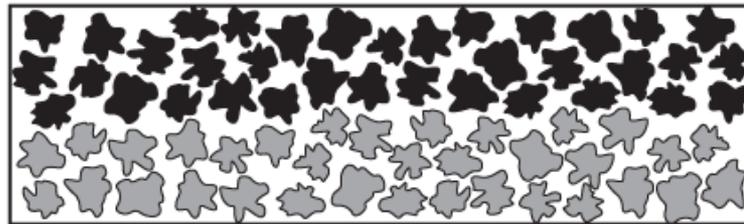


... but cells can be active (not just passive)!

Cell Rearrangement Mechanism

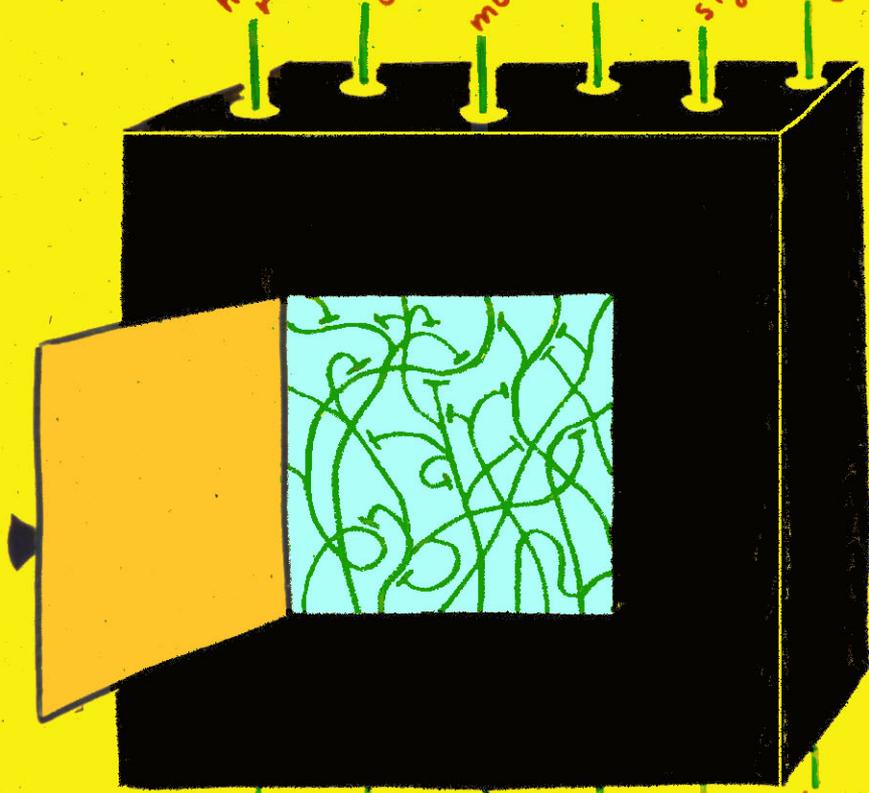


↓ segregation?



CELL
LEVEL

mitotic
rates
mitotic
orientations
movements
lineage
signaling
cell
death



repressors
activators
switches
knobs
clocks
sensors

GENE
LEVEL

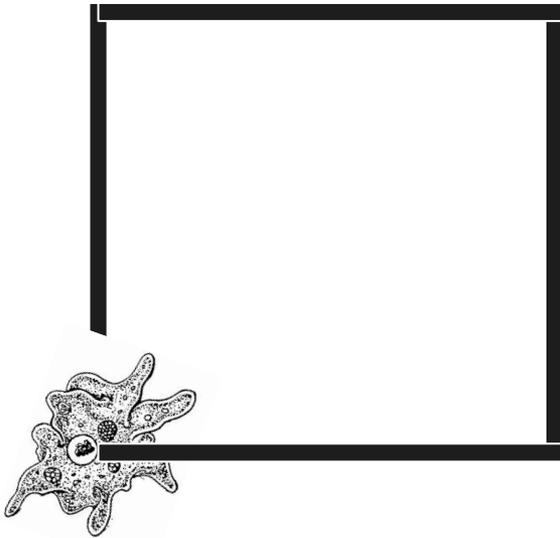
Cell Commands:

Square

Crawl forward 100 units.

Turn right 90 degrees.

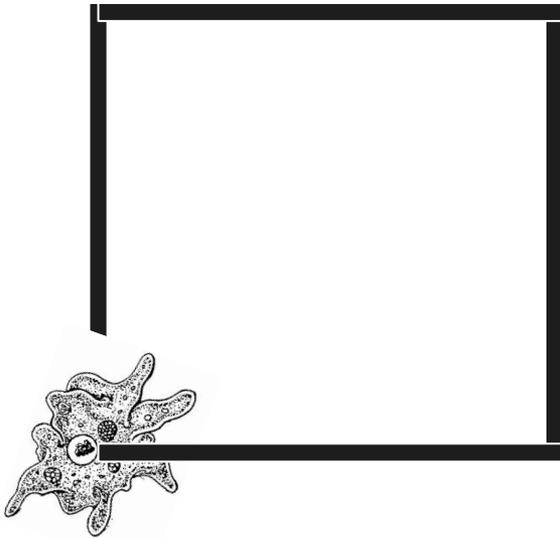
Repeat 4 times.



Cell Commands:

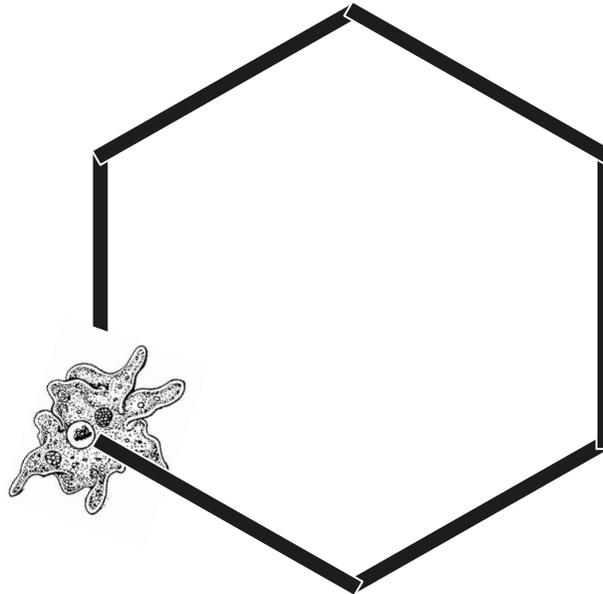
Square

Crawl forward 100 units.
Turn right 90 degrees.
Repeat 4 times.



Hexagon

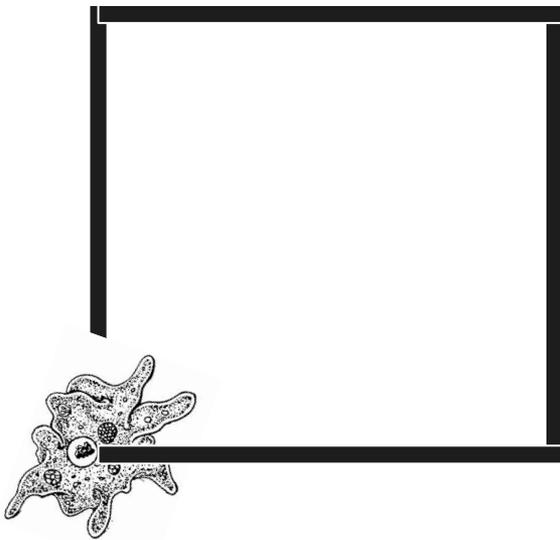
Crawl forward 60 units.
Turn right 60 degrees.
Repeat 6 times.



Cell Commands:

Square

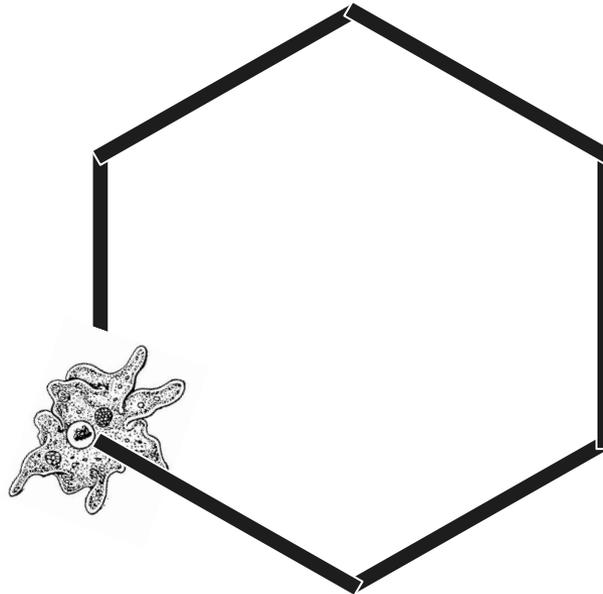
Crawl forward 100 units.
Turn right 90 degrees.
Repeat 4 times.



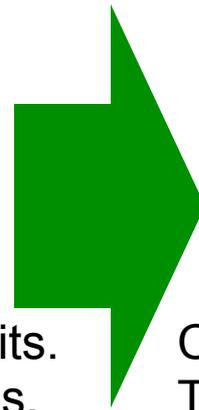
(4, 100)

Hexagon

Crawl forward 60 units.
Turn right 60 degrees.
Repeat 6 times.

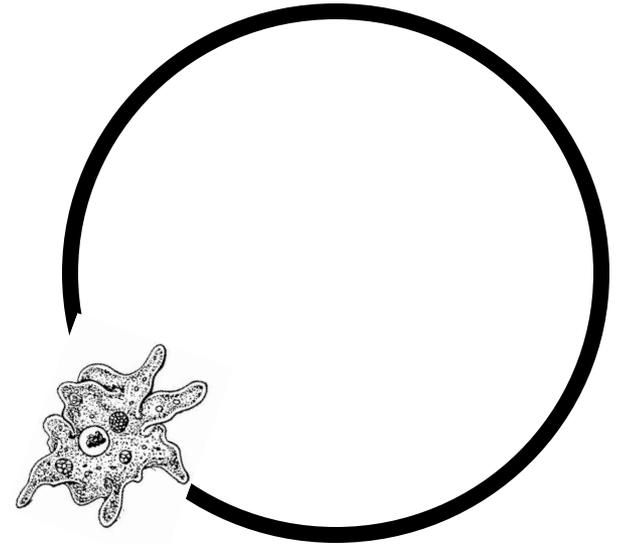


(6, 60)



Polygon (n, s)

Crawl forward s units.
Turn right $360/n$ degrees.
Repeat n times.



(100, 1)

Cell Commands:

Dev. Program

Input Variables

Square

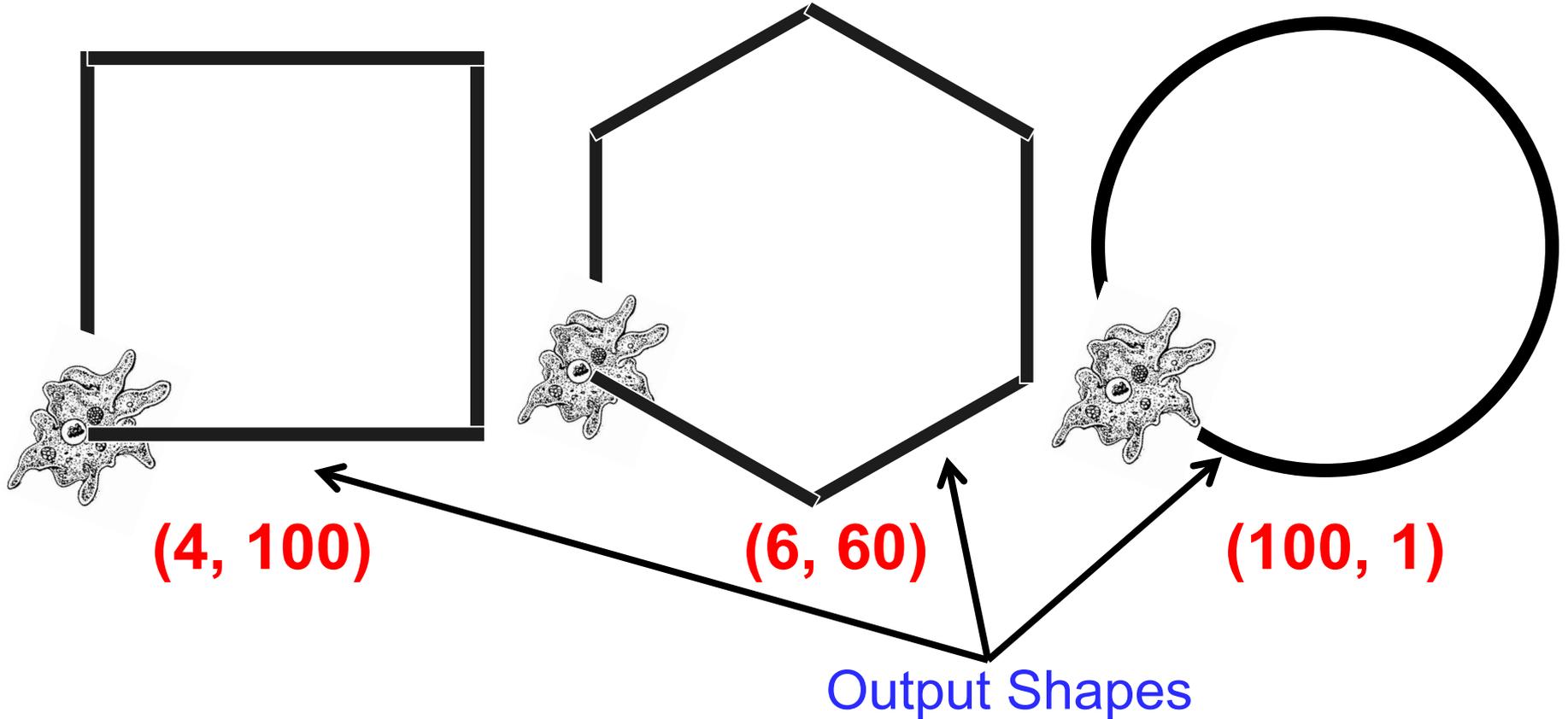
Crawl forward 100 units.
Turn right 90 degrees.
Repeat 4 times.

Hexagon

Crawl forward 60 units.
Turn right 60 degrees.
Repeat 6 times.

Polygon (n, s)

Crawl forward s units.
Turn right $360/n$ degrees.
Repeat n times.



$(4, 100)$

$(6, 60)$

$(100, 1)$

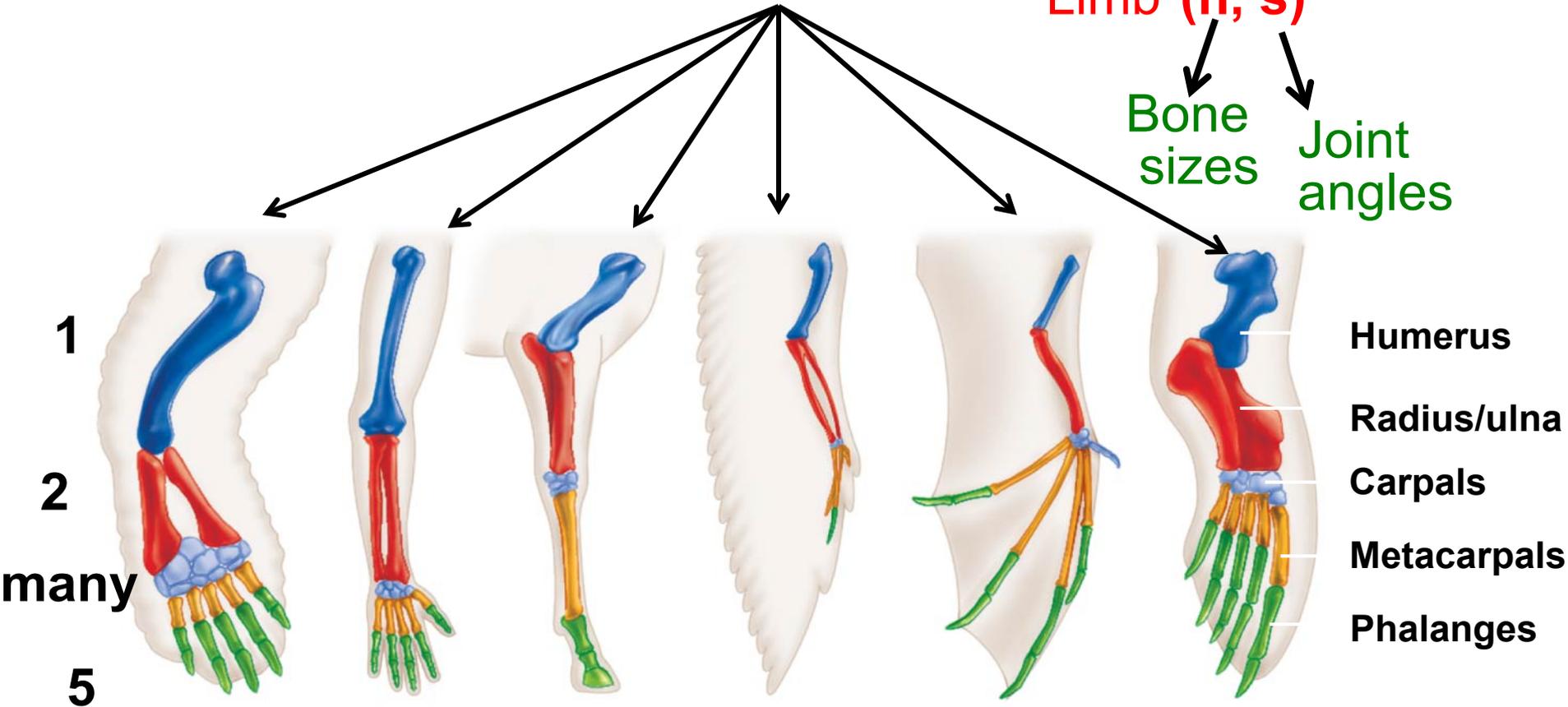
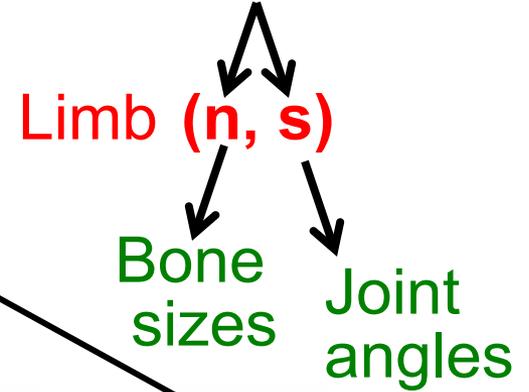
Output Shapes

Homology = Similarity due to  **Descent ... with modification**

Dev. Program

Input Variables

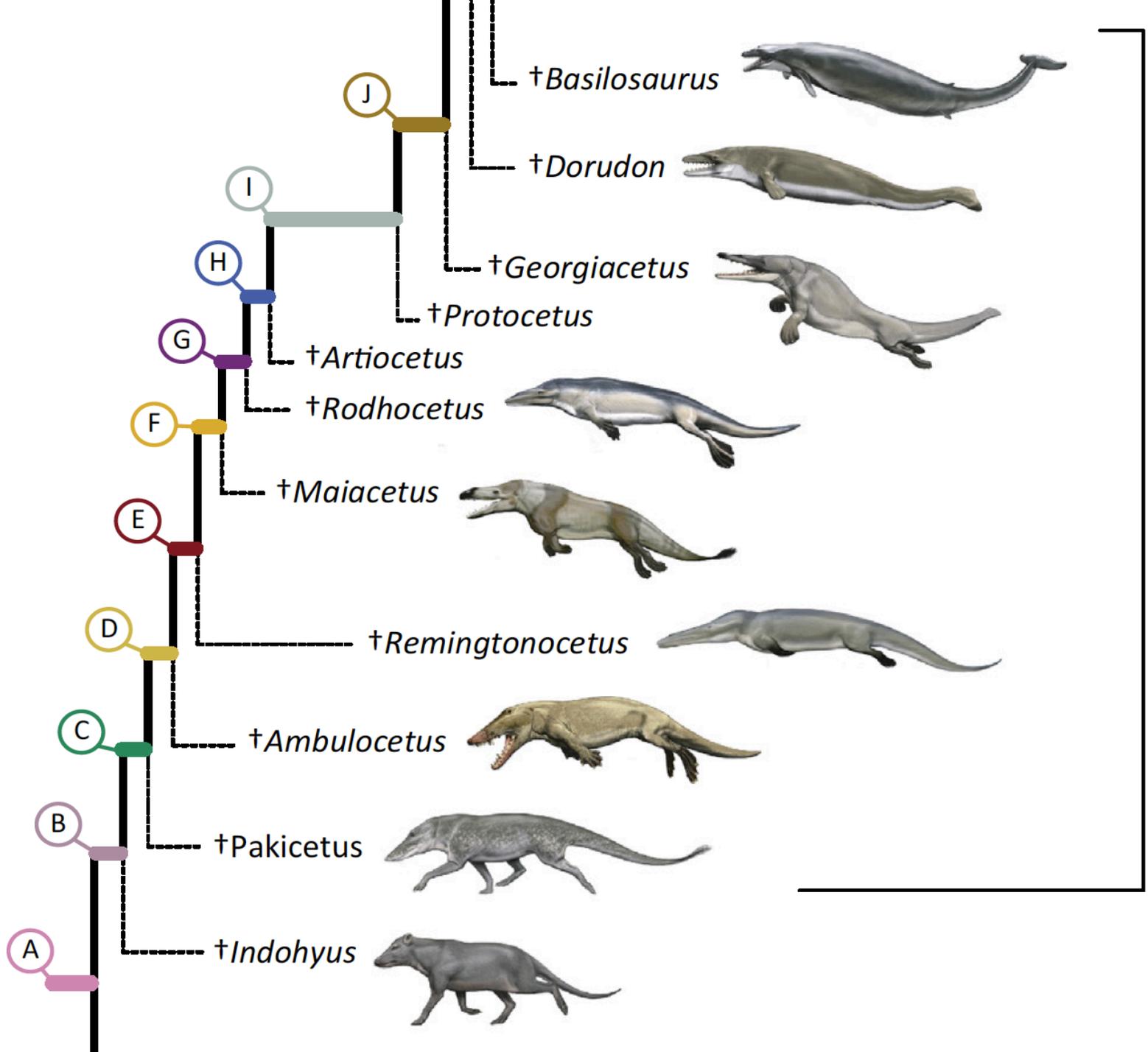
Cell Commands: 1:2:many:5



Output Anatomies

**Q: How does evolution
rewire the genome?**

**A: By changing the inputs
to a conserved program,
which changes anatomy.**



Molecular evolution tracks macroevolutionary transitions in Cetacea

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¹Center for Molecular Medicine and Genetics, Wayne State University School of Medicine, Detroit, MI 48201, USA

²Department of Biology, University of California-Riverside, Riverside, CA 92521, USA

Cetacea (whales, dolphins, and porpoises) is a model group for investigating the molecular signature of macroevolutionary transitions. Recent research has begun to reveal the molecular underpinnings of the remarkable anatomical and behavioral transformation in this clade. This shift from terrestrial to aquatic environments is arguably the best-understood major morphological transition in vertebrate evolution. The ancestral body plan and physiology were extensively modified and, in many cases, these crucial changes are recorded in cetacean genomes. Recent studies have highlighted cetaceans as central to understanding adaptive molecular convergence and pseudogene formation. Here, we review current research in cetacean molecular evolution and the potential of Cetacea as a model for the study of other macroevolutionary transitions from a genomic perspective.

A poster child for macroevolution

The evolution of cetaceans (see [Glossary](#)) has emerged as a poster child for macroevolution, and is one of the best-

Cetacea offers notable advantages relative to other taxa that also have experienced major anatomical and ecological reorganizations. Phylogenetic hypotheses for extinct and extant taxa are well developed ([Figure 1](#)), and include molecular divergence times among extant species with integration of extensive fossil data [[4–8](#)]. Semiaquatic hippopotamids (hippopotamuses), potential extant ‘intermediate forms,’ share multiple aquatic traits with cetaceans ([Figure 2A,B](#)) and might provide critical genetic insights into the early evolution of Cetacea ([Figure 1](#)).

Glossary

Cetaceans: a clade of mammals that includes whales, dolphins, and porpoises; the approximately 90 extant species are characterized by a wholly aquatic lifestyle.

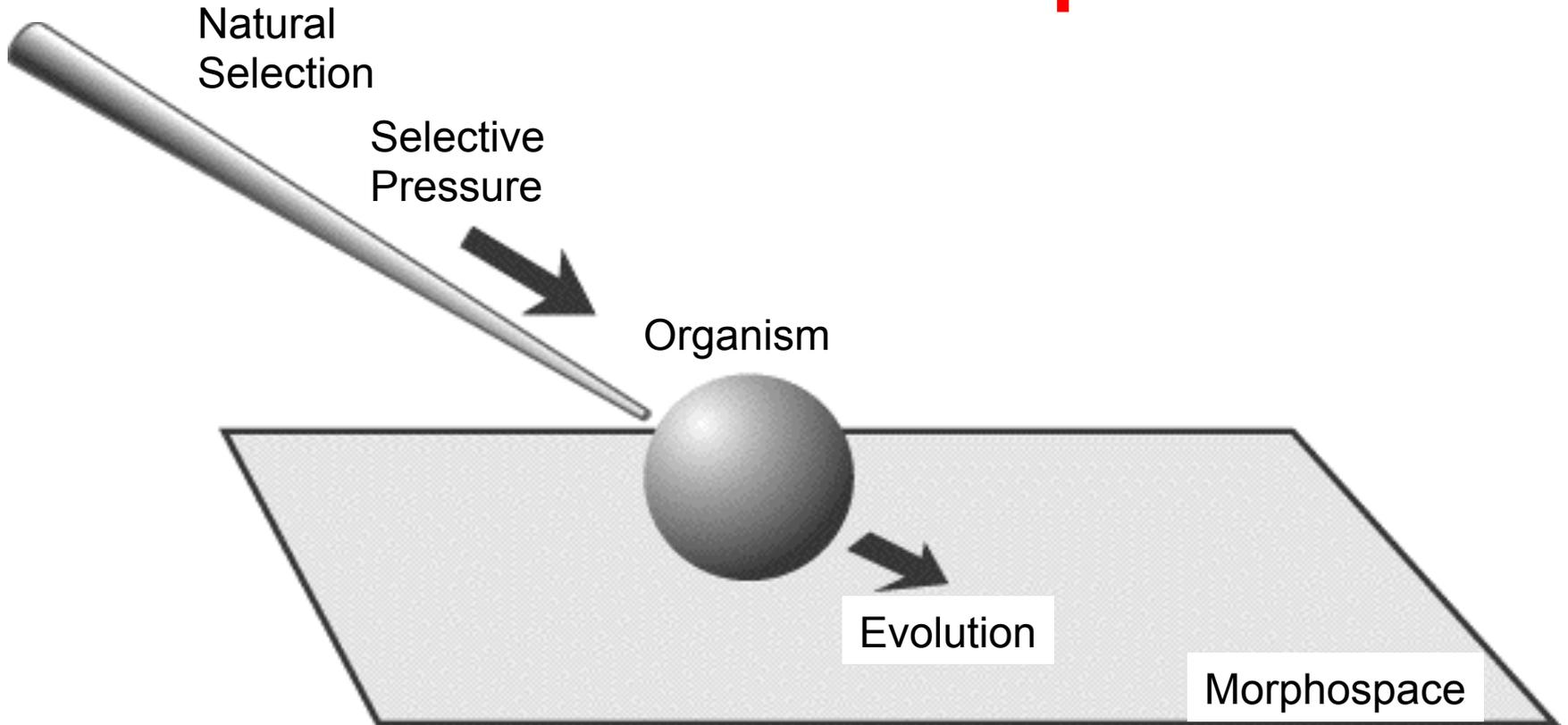
Cetartiodactyls: a clade of mammals that includes camels, pigs, peccaries, cattle, antelope, deer, chevrotains, giraffes, hippopotamuses, and cetaceans.

Convergent recruitment: evolutionary change whereby an orthologous gene is utilized in two or more independent lineages to effect a similar phenotypic outcome in each lineage.

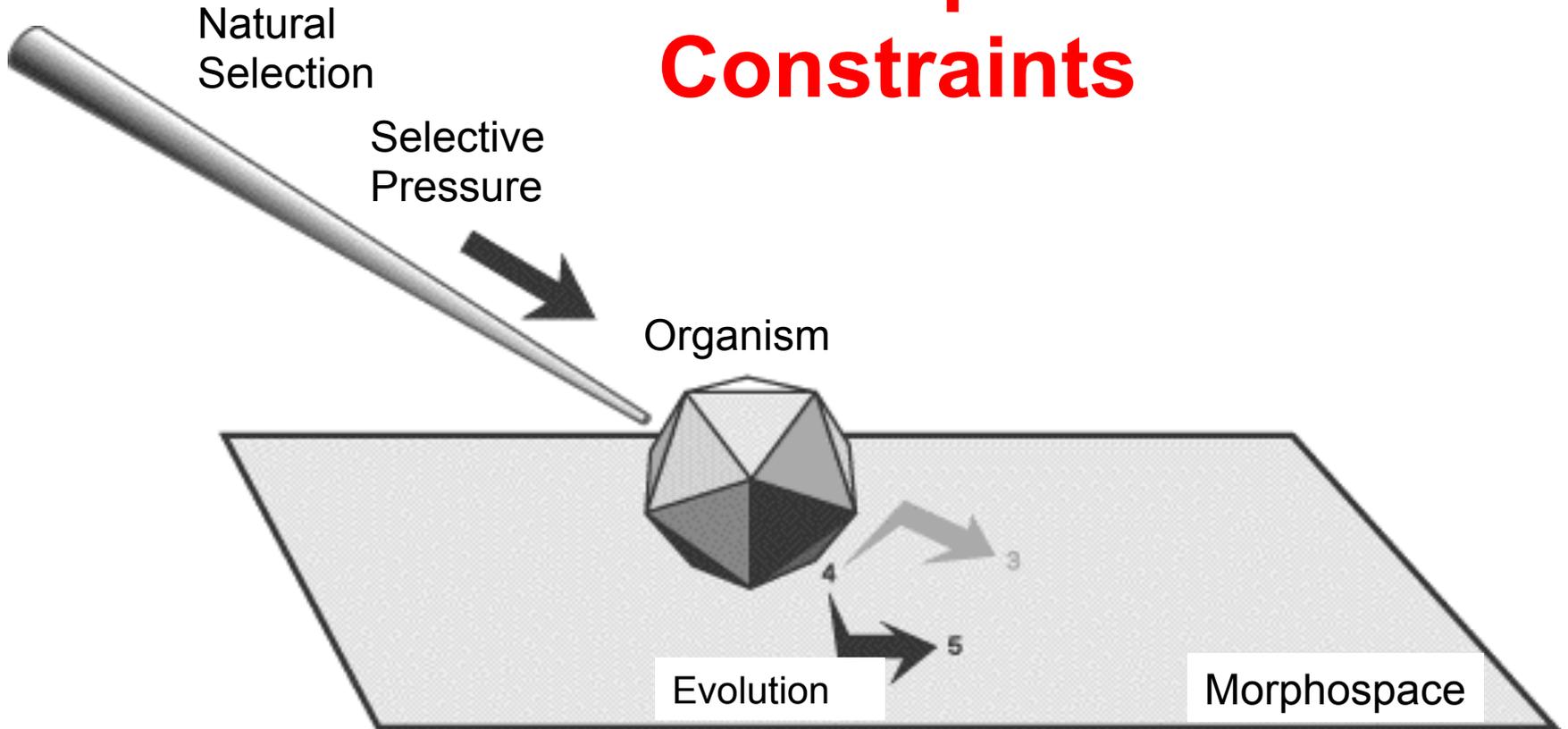
Convergent substitution: independent molecular evolution of the same state at a homologous position of the same gene and/or locus in two or more evolutionary lineages, affecting similar phenotypic change in each lineage.

**Why are some traits conserved ...
while others change very easily?**

Misconception



Developmental Constraints





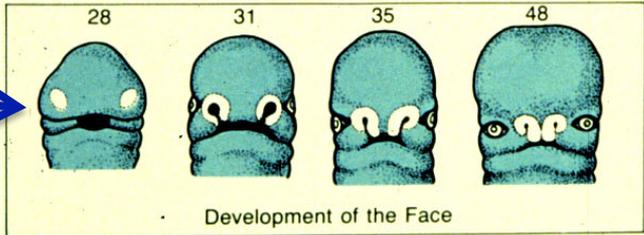
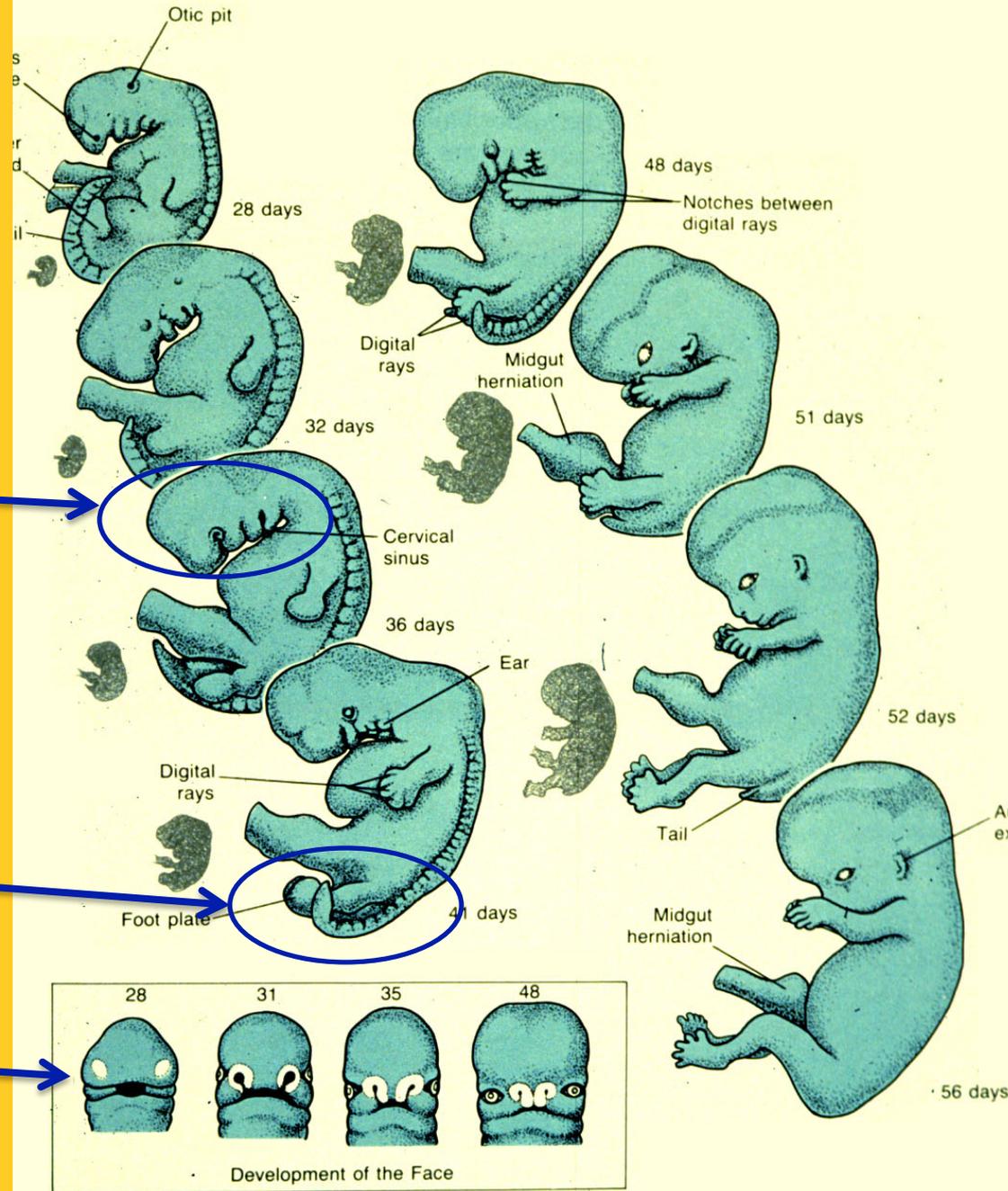
“Geep”

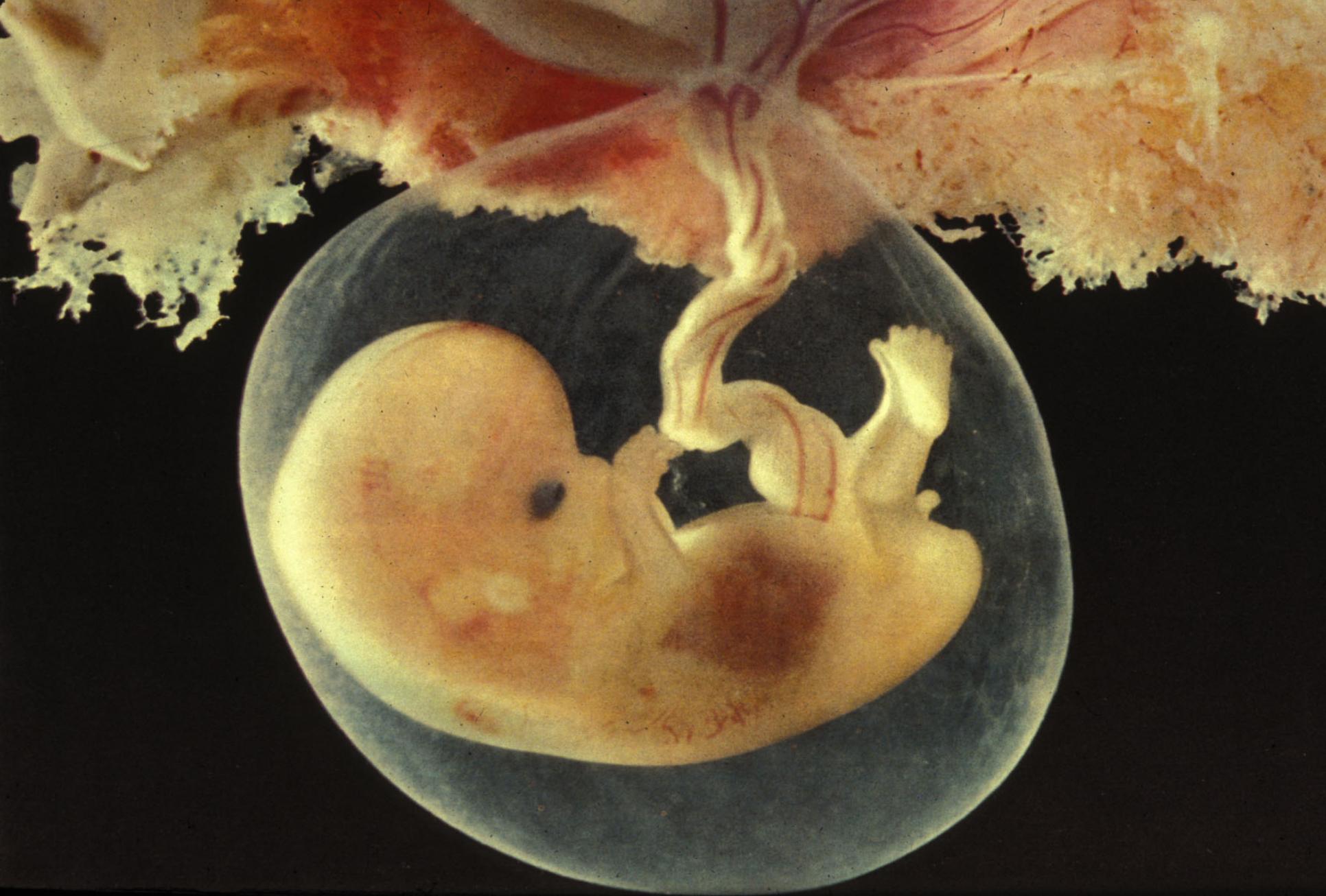
Ontogeny Recapitulates Phylogeny

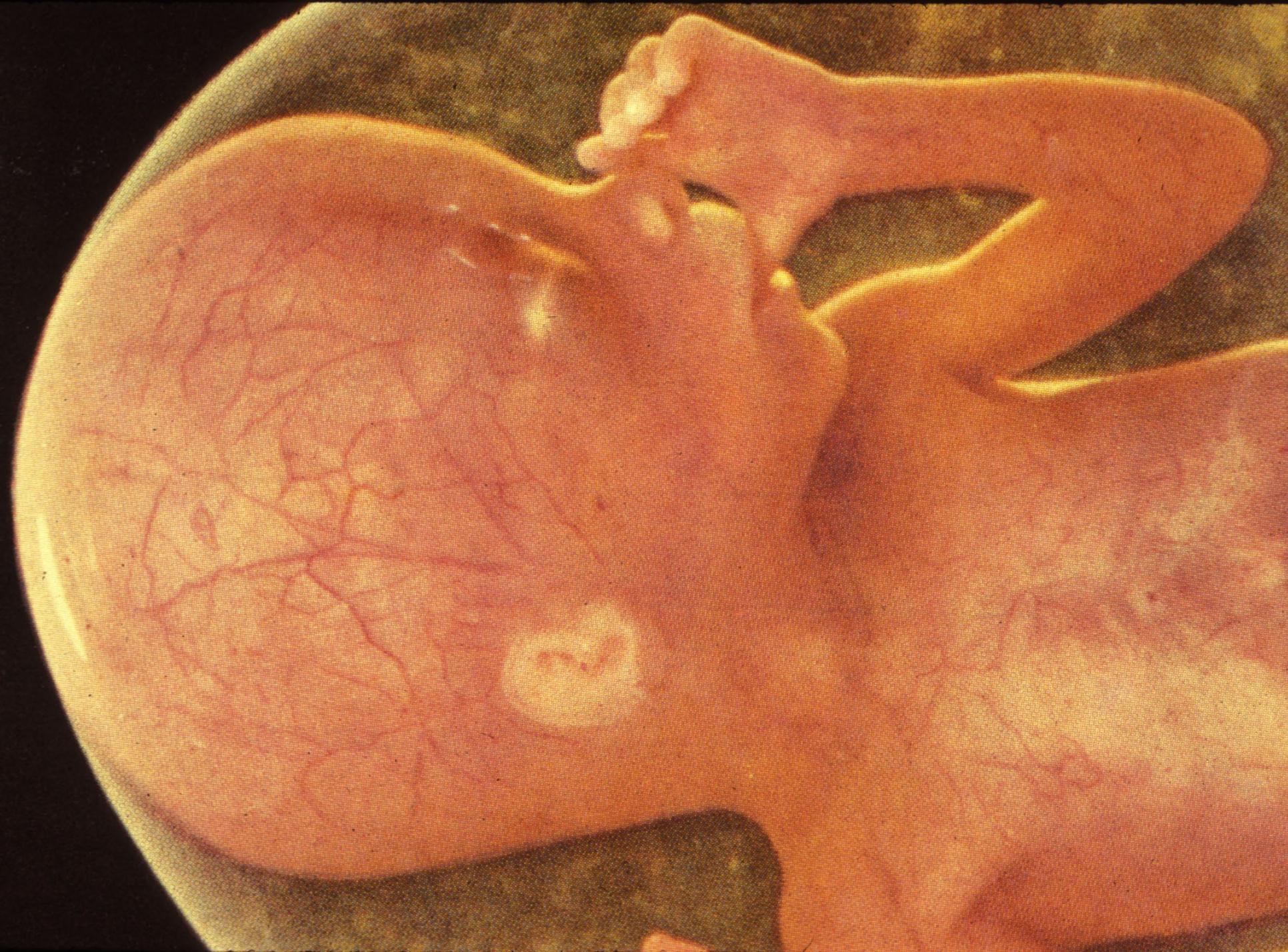
Gill slits (like a fish)

Tail (like a monkey)

Eyes on side (\approx fish)



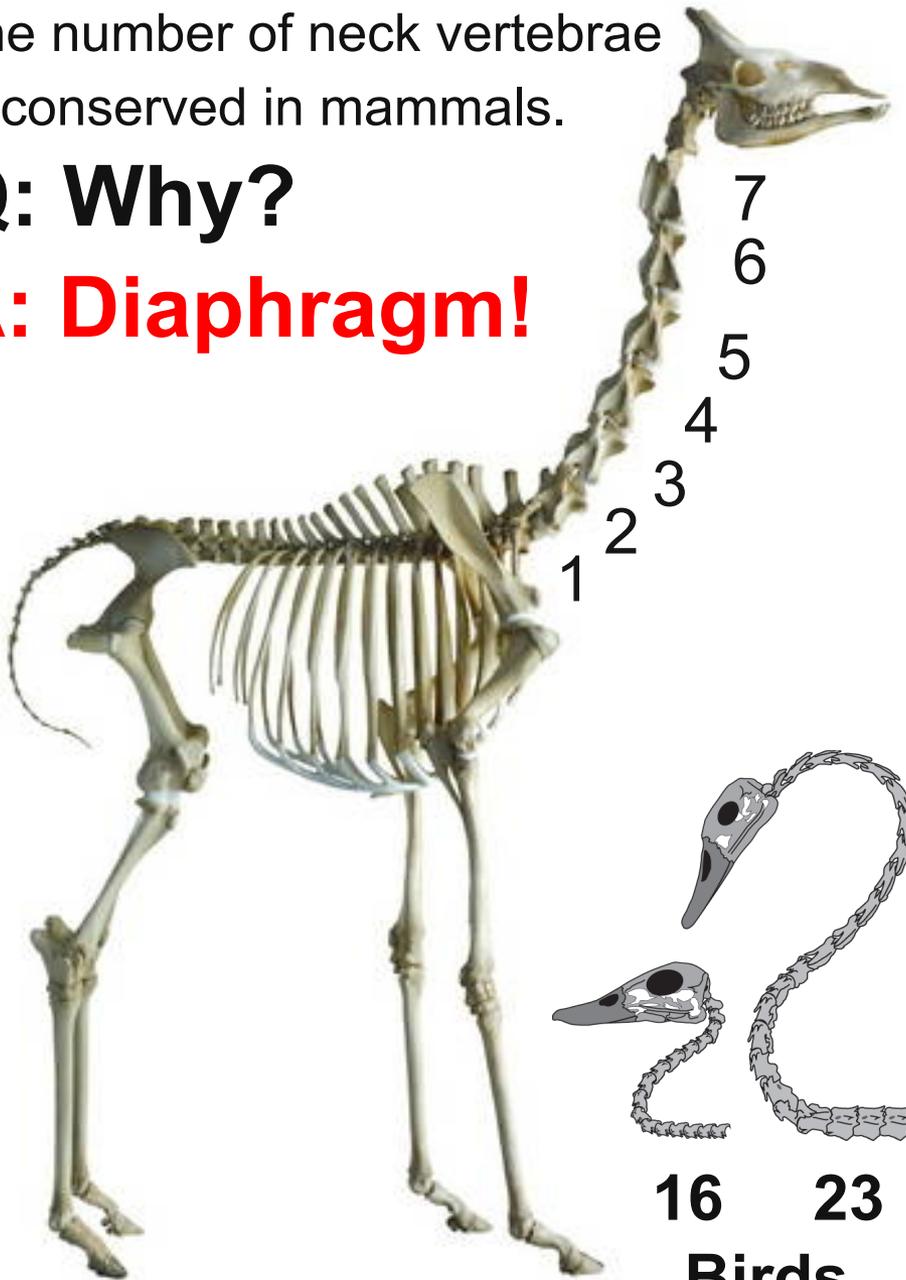




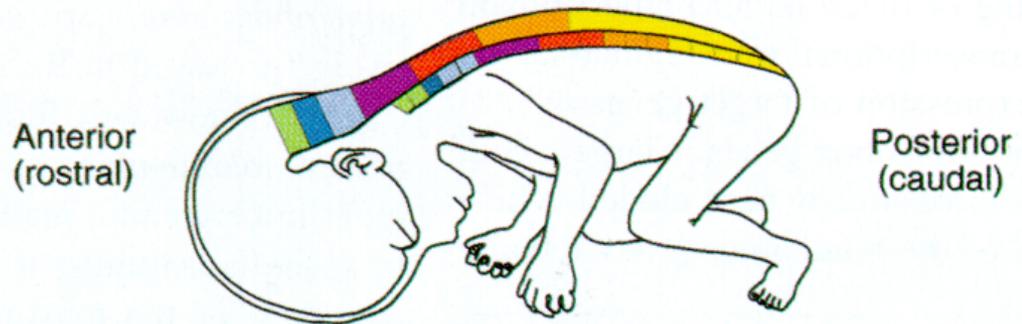
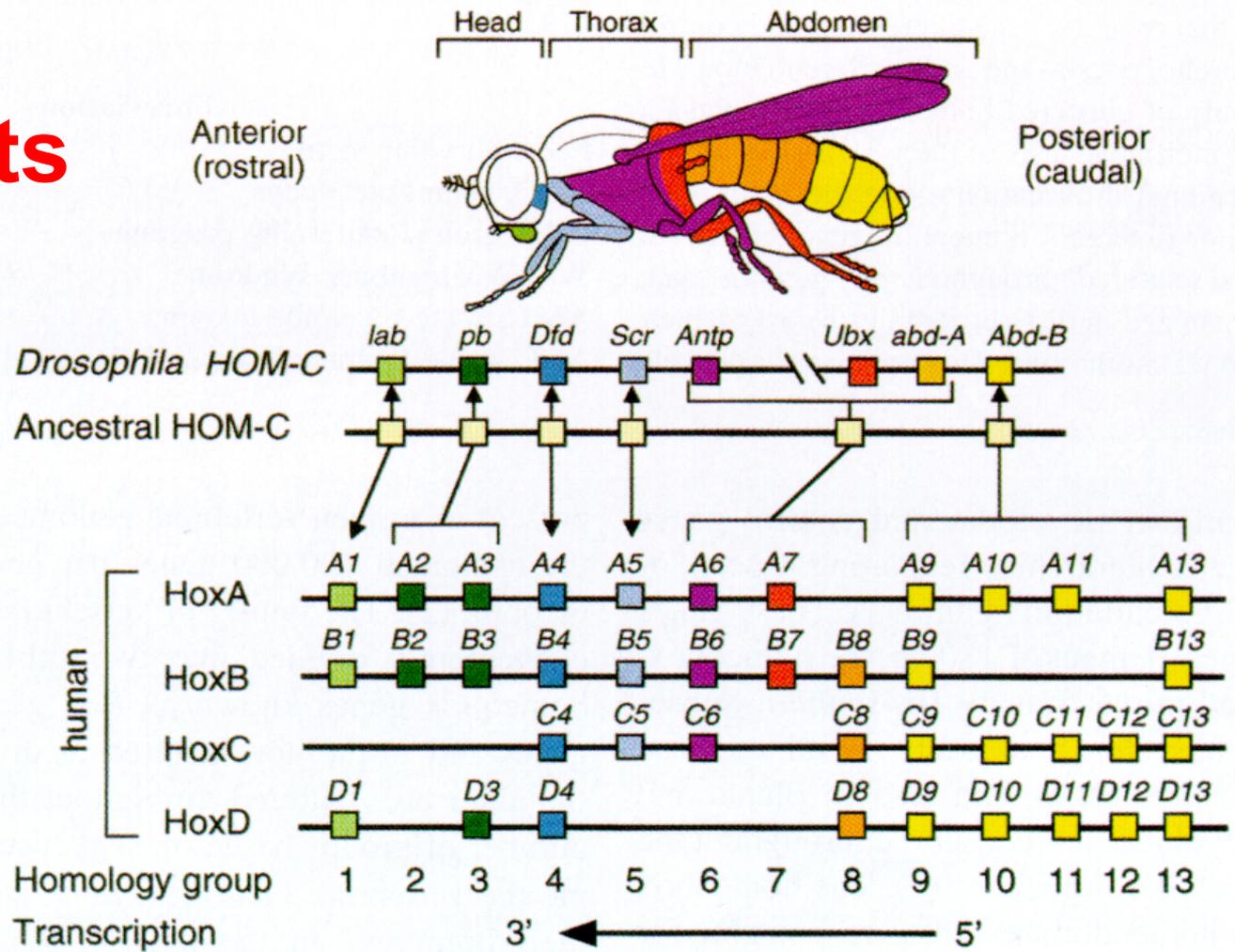
The number of neck vertebrae
Is conserved in mammals.

Q: Why?

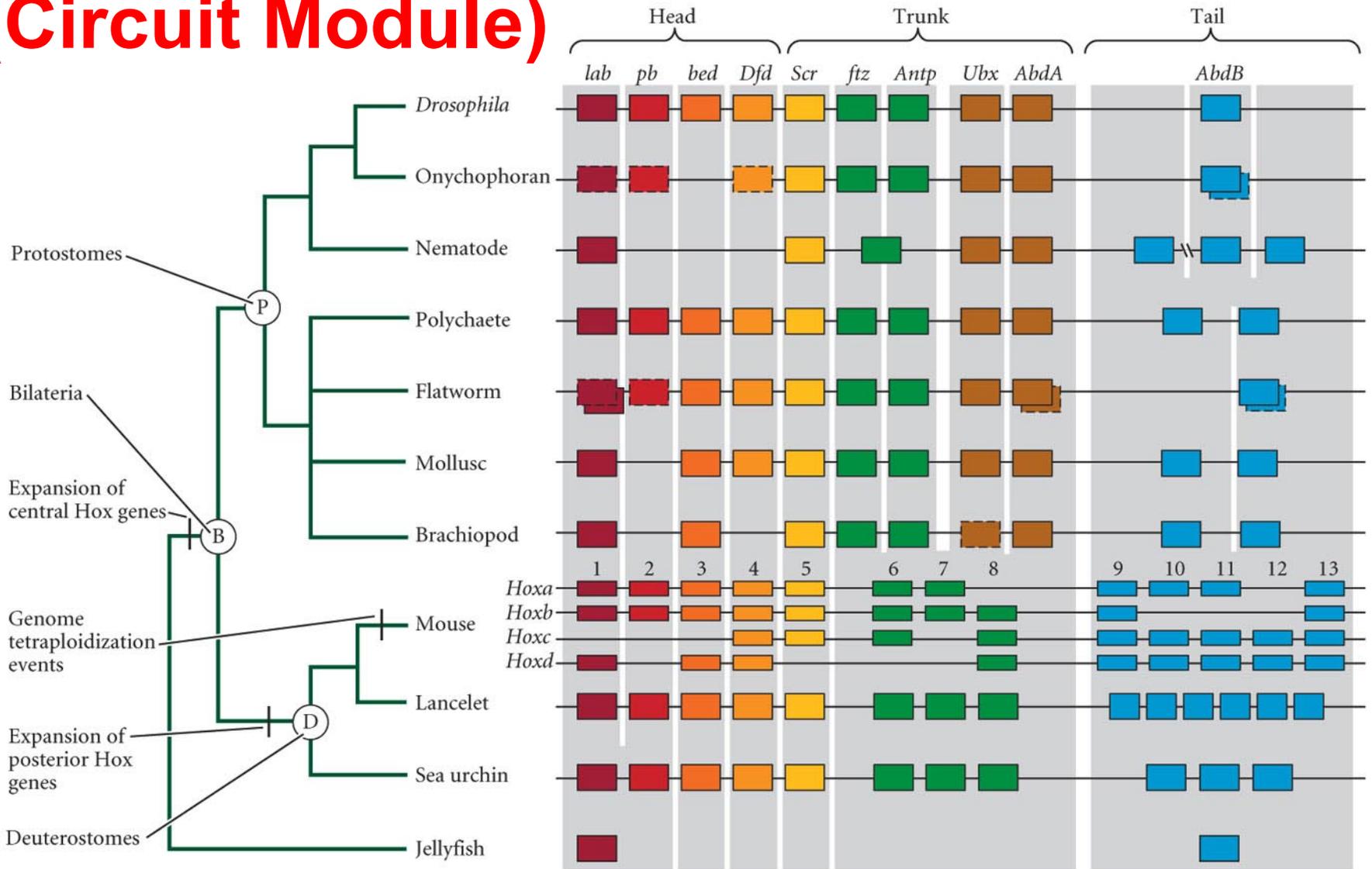
A: Diaphragm!



Genetic Constraints

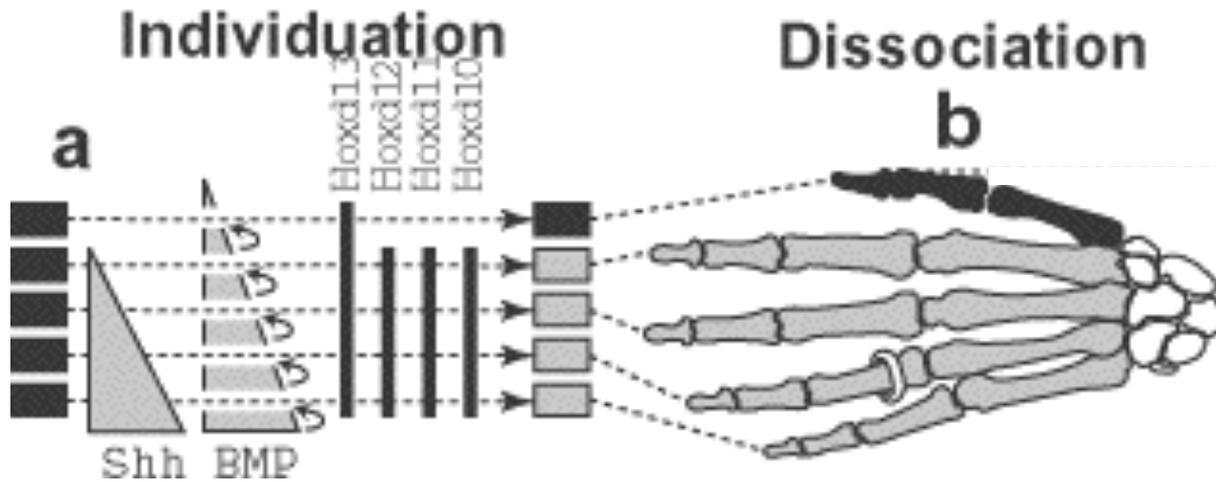


600 Million Yrs. Old! (Circuit Module)



Co-option:

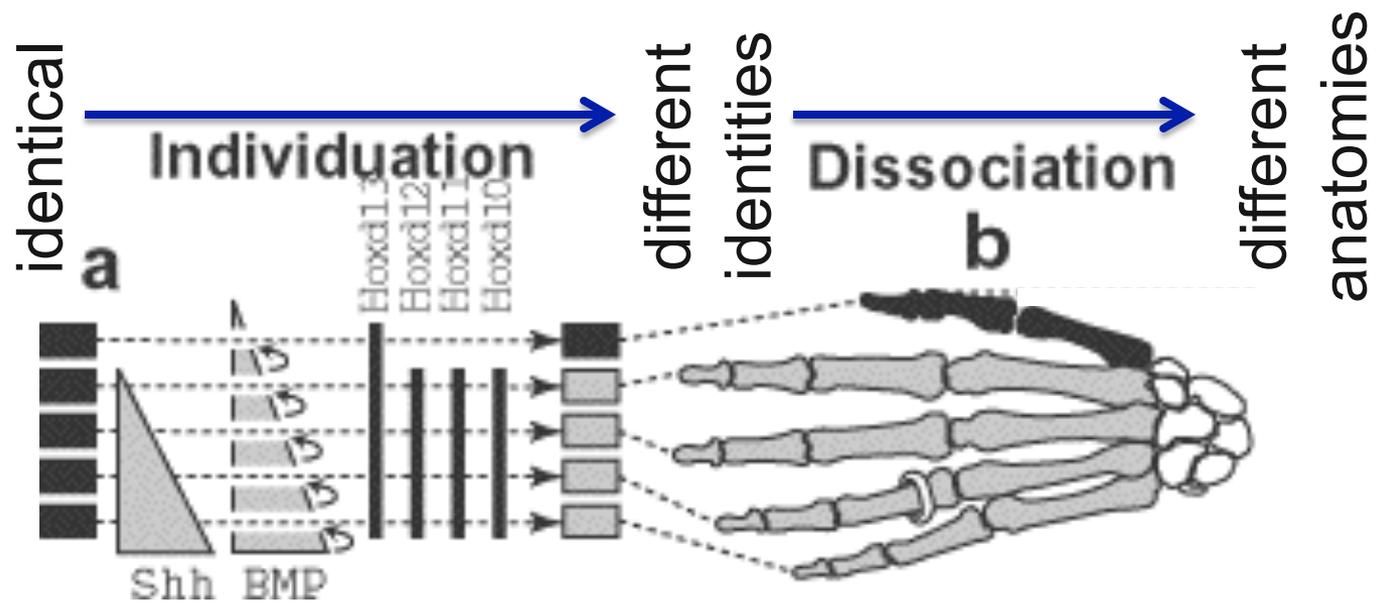
Re-wiring of an old circuit (Hox) for a new function (fingers).



Morphogen (Shh or BMP) = a (diffusible) signaling molecule.

Gradient = assigning cell positions by morphogen concentrations.

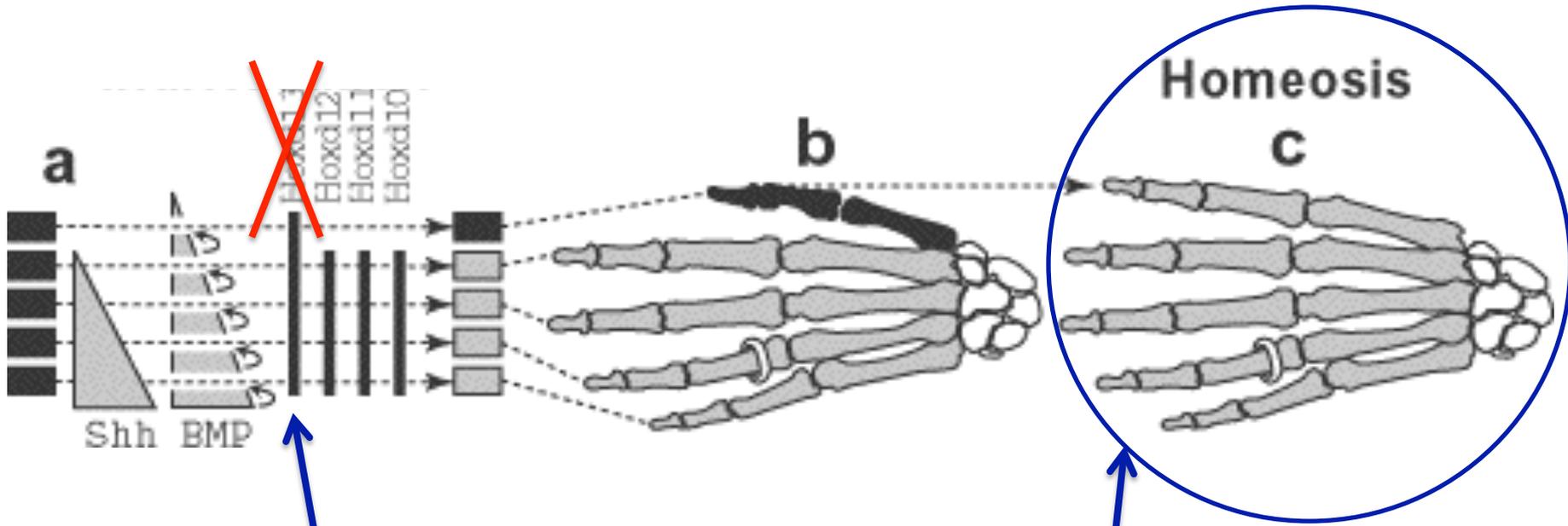
Individuation: Giving different identities to similar structures (via genes).



Dissociation: The independent evolution of ≥ 1 members of a series.

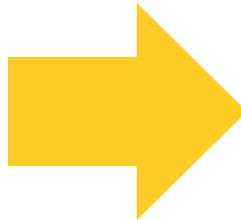
Individuation (in development) allows dissociation (in evolution).

Homeosis: Changing one body part (thumb) to resemble another (finger).



Mutations in Hoxd13 cause a “bear claw” phenotype!

≈



**Q: How does evolution
rewire the genome?**

**A: By co-opting old circuits
for new functions.**

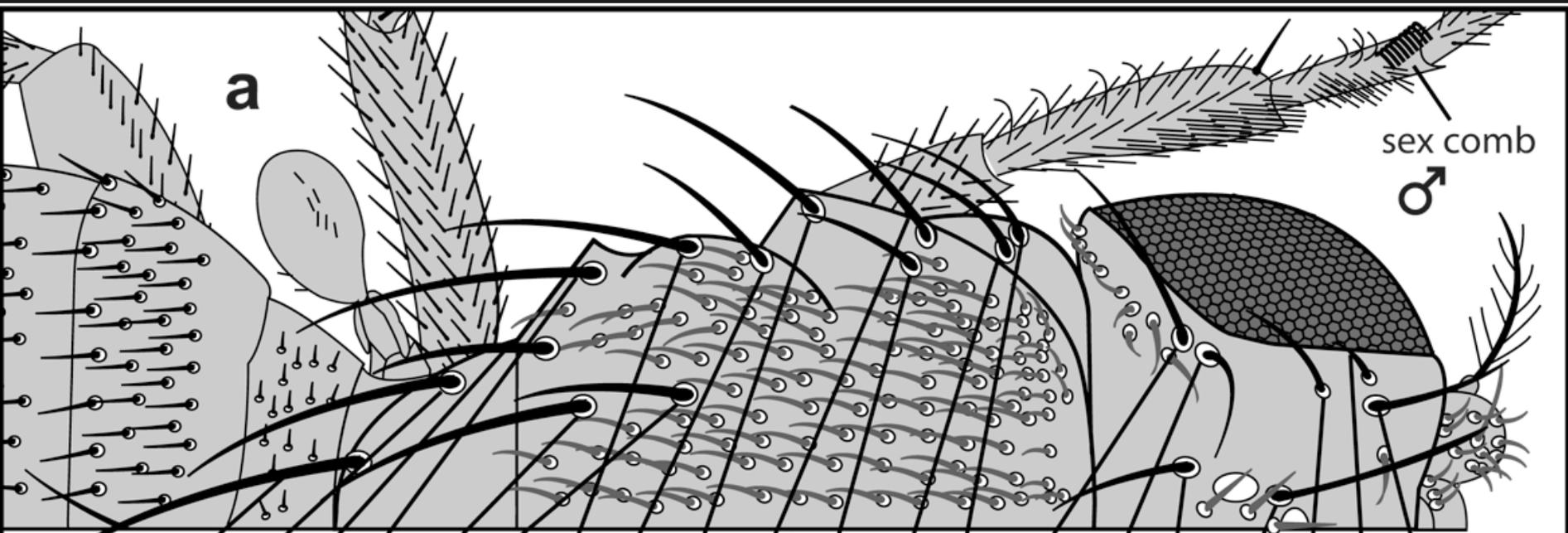
Human Evolution





a

sex comb
♂



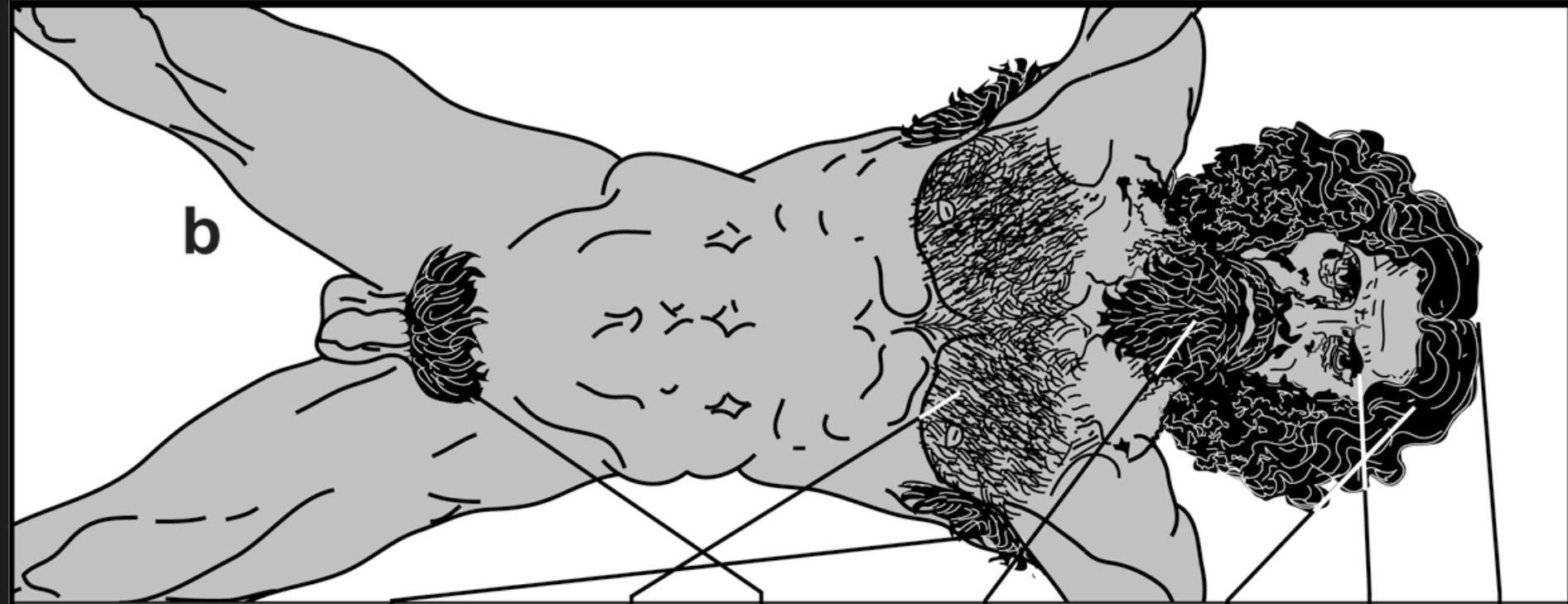
70 kb 60 50 40 30 20 10 0 -10 -20 -30 kb



pigment gene **ac** **sc** *l'sc* *T2* region-specific enhancers #1-8 *ase* *T1*

Bristle Gene Complex

b



imaginary
region-
specific
enhancers
#1-7?

armpit

chest
♂

pubic

→
WNT?

beard
♂

→
?

scalp

eyebrow

crown
♂
(baldness)

Hair Gene Complex?

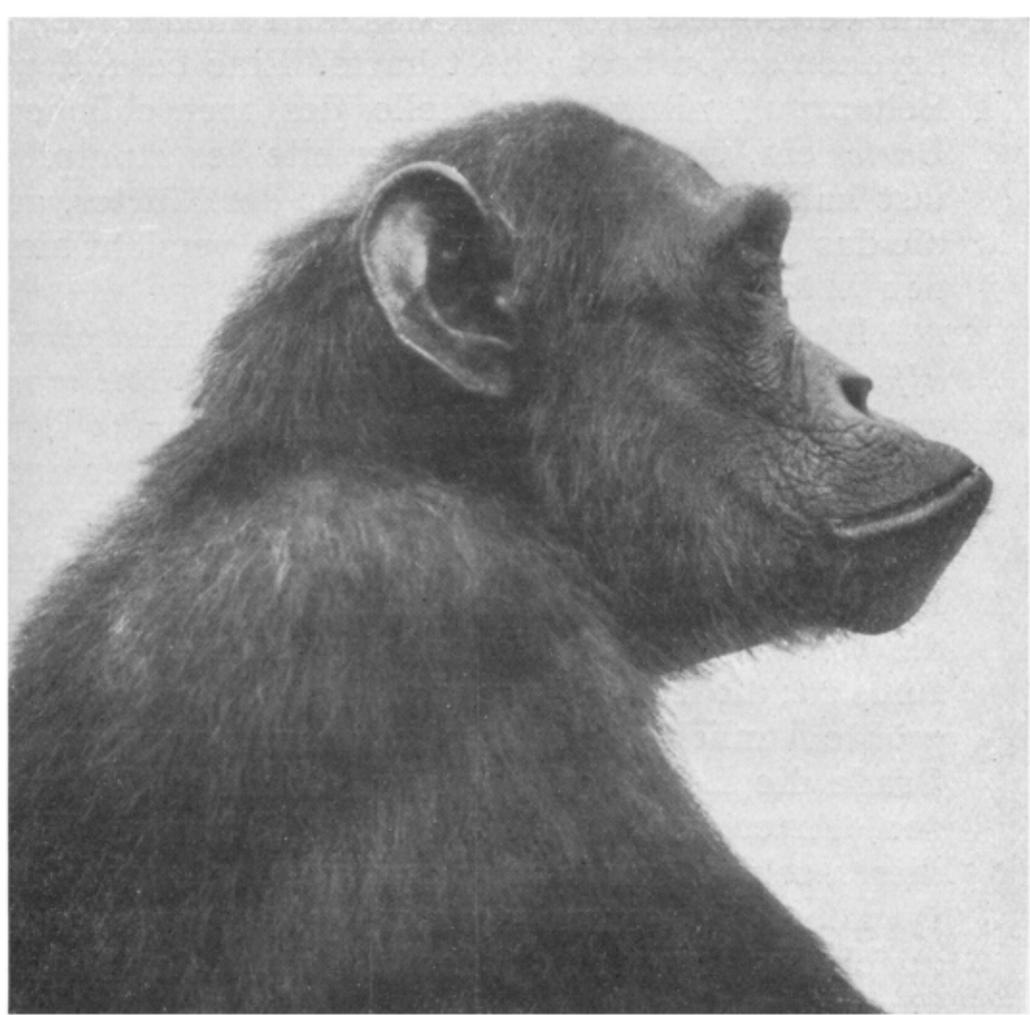
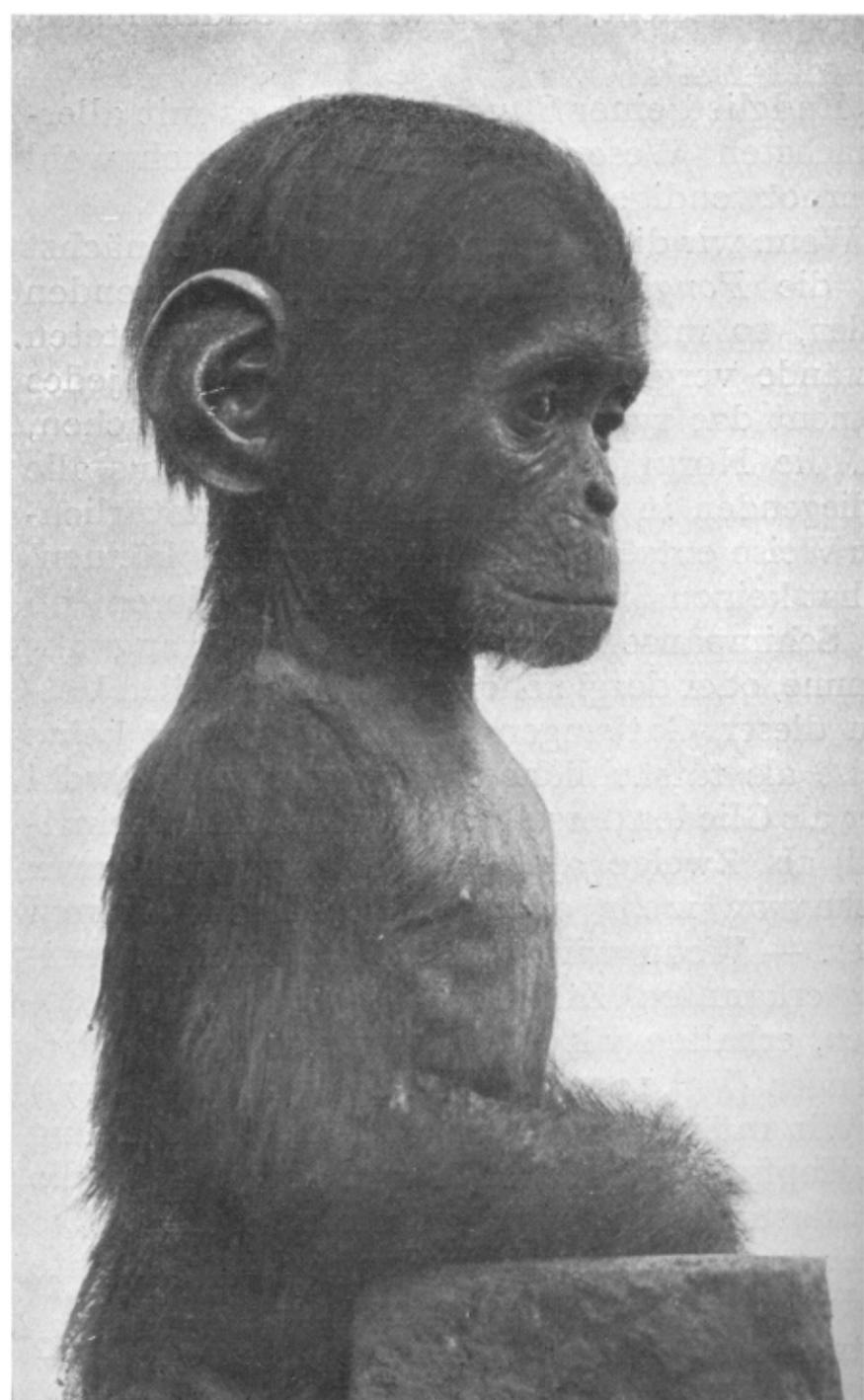
hair gene?

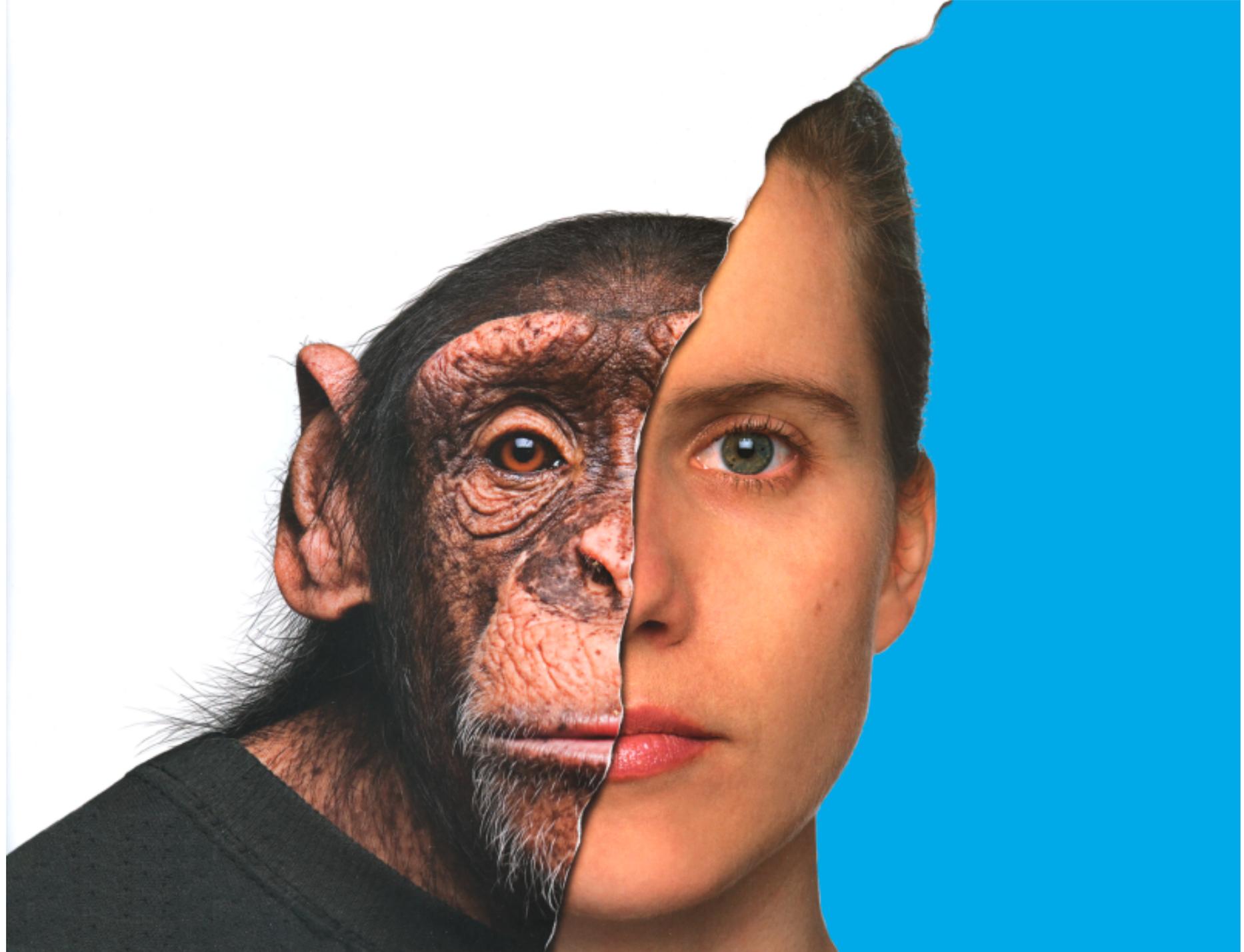
pigment gene?

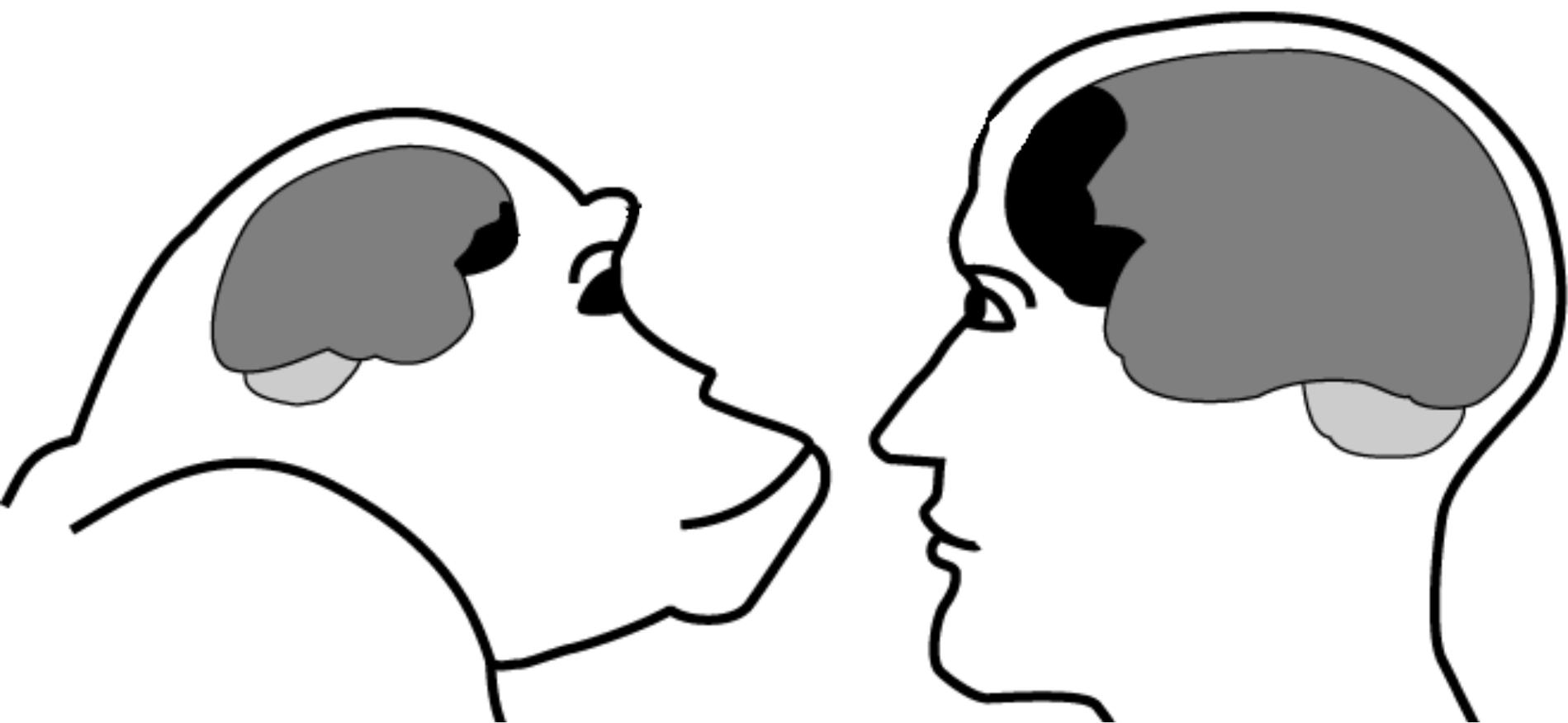
**Q: How does evolution
rewire the genome?**

**A: By tweaking cis-regulatory
elements for area codes.**









3.13

log [brain weight(g)]

0

Age (days post conception)

2023

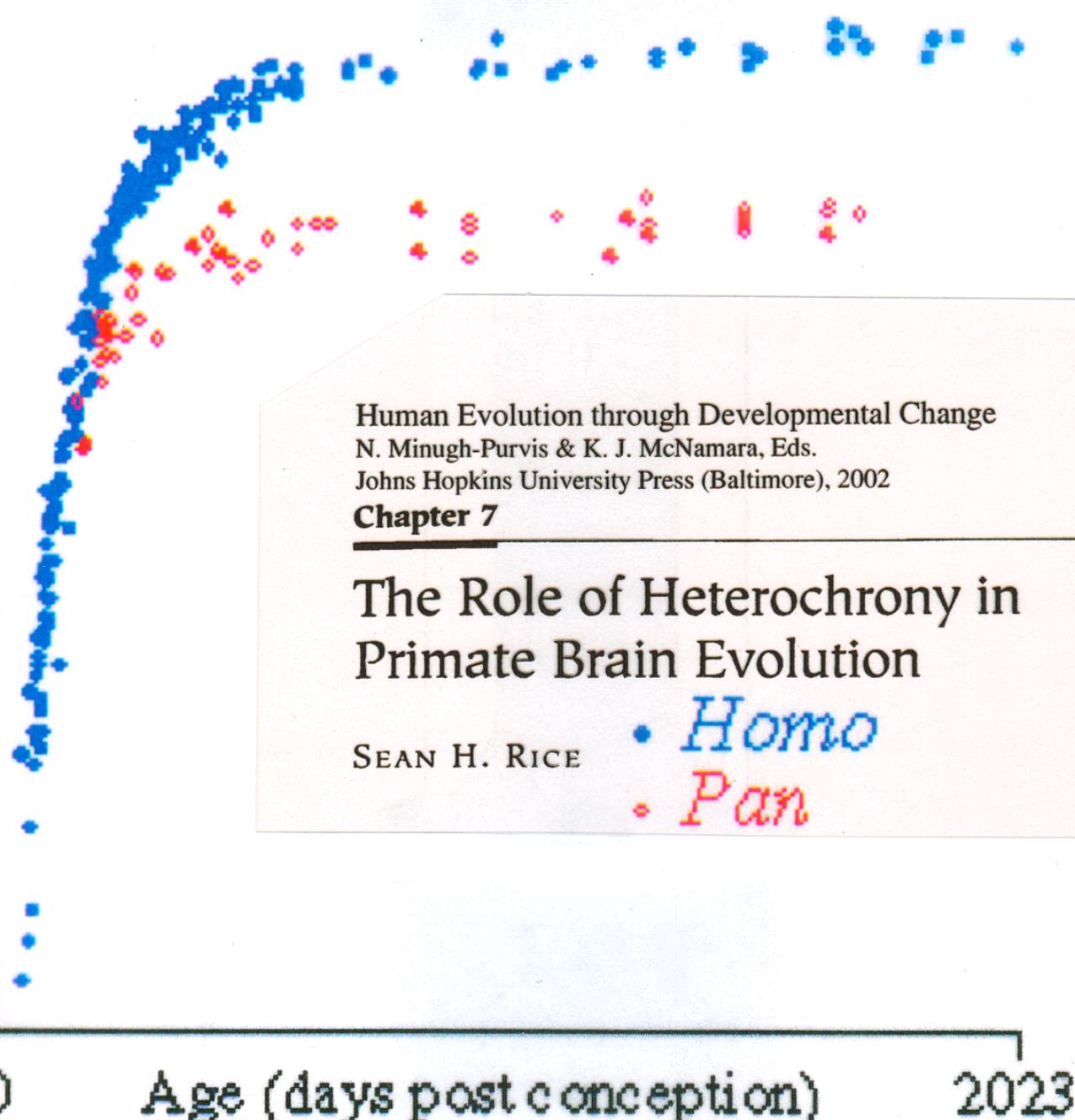
Human Evolution through Developmental Change
N. Minugh-Purvis & K. J. McNamara, Eds.
Johns Hopkins University Press (Baltimore), 2002
Chapter 7

The Role of Heterochrony in Primate Brain Evolution

SEAN H. RICE

• *Homo*

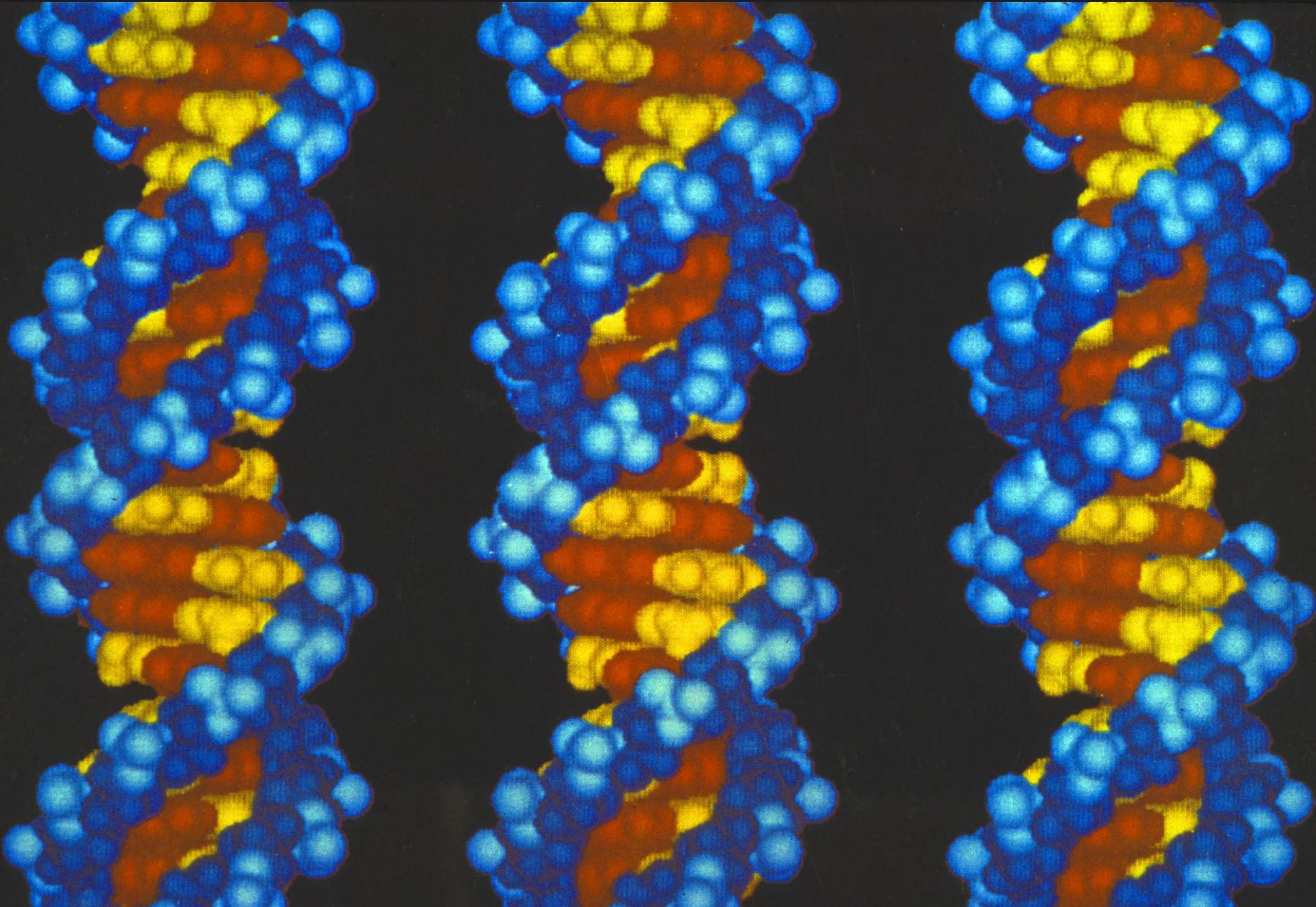
• *Pan*



**Q: How does evolution
rewire the genome?**

**A: By changing the timing
of gene expression
(= Heterochrony).**

The history of evolution is written in our genomes!



Summary

How does evolution re-wire the genome to re-program development to re-configure anatomy?

-
- **By changing the inputs to a conserved program.**
 - **By co-opting old circuits for new functions.**
 - **By tweaking cis-regulatory elements for area codes.**
 - **By changing the timing of gene expression.**

