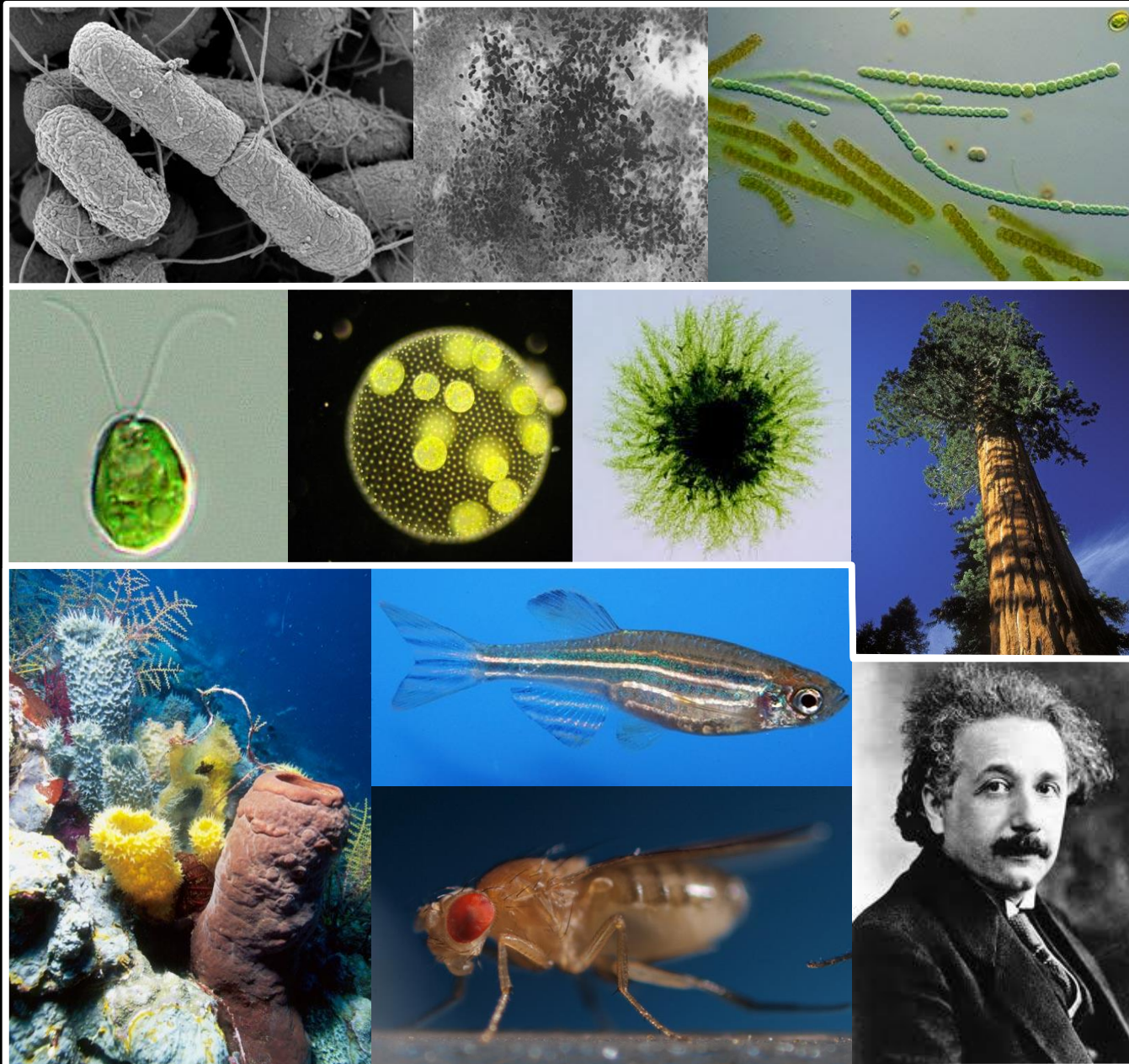


Phylotranscriptomic hourglasses of plant and animal embryogenesis

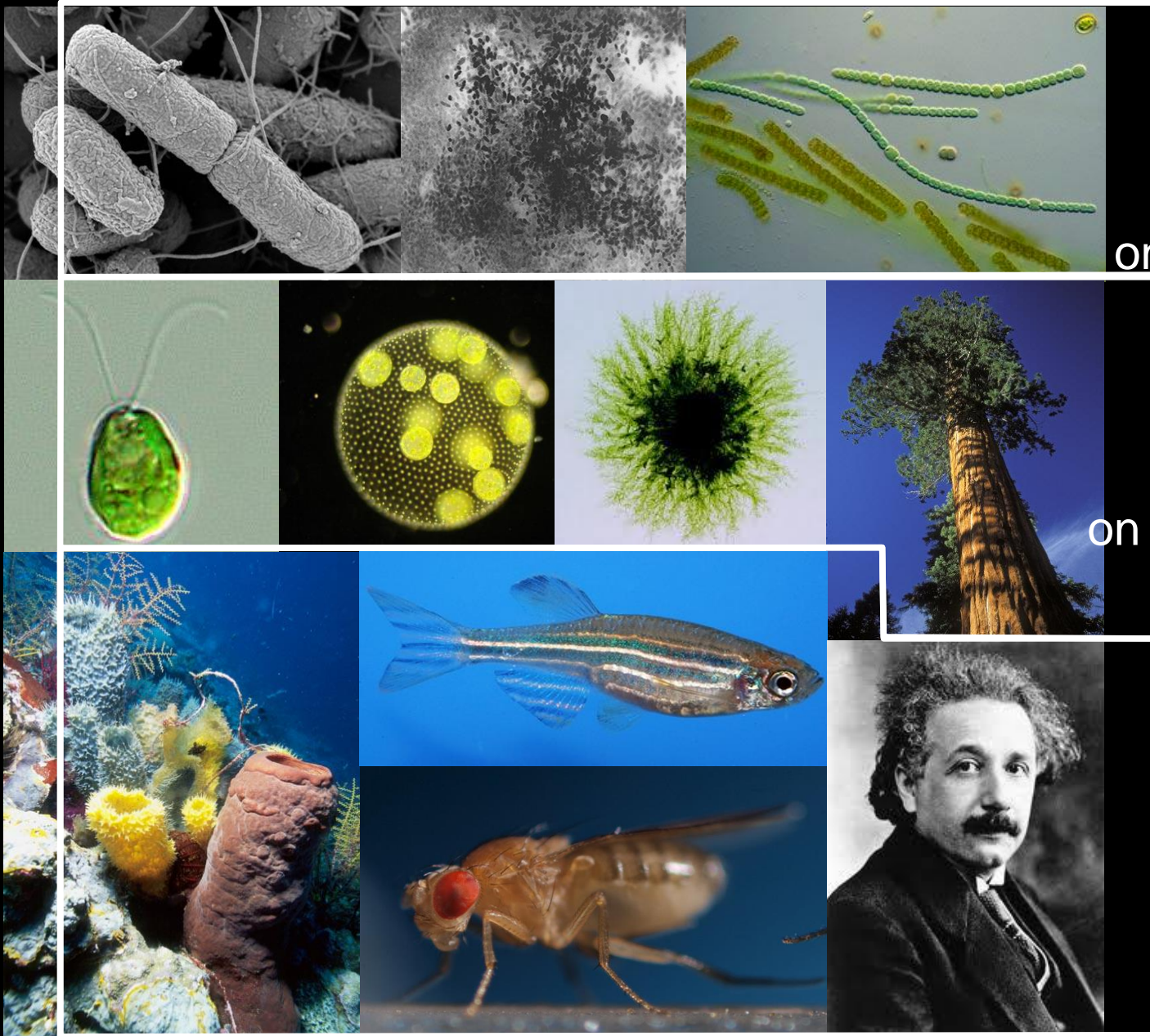
Marcel Quint, Hajk-Georg Drost, Alexander Gabel, Kristian Karsten Ullrich, Markus Boenn, and Ivo Grosse

*Leibniz Institute of Plant Biochemistry
Institute of Computer Science at Halle University
German Center of Integrative Biodiversity Research (iDiv) Halle-Jena-Leipzig*

How did **complex life** emerge?



How did **complex life** emerge?

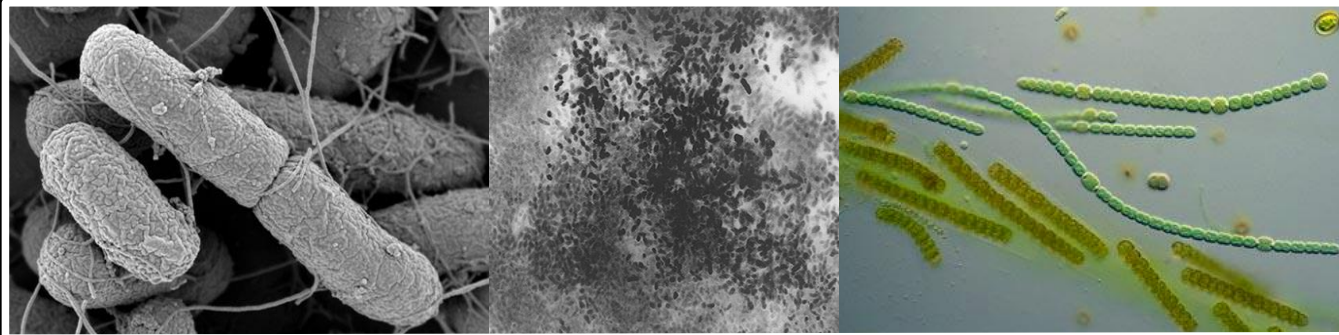


on the **evolutionary** level

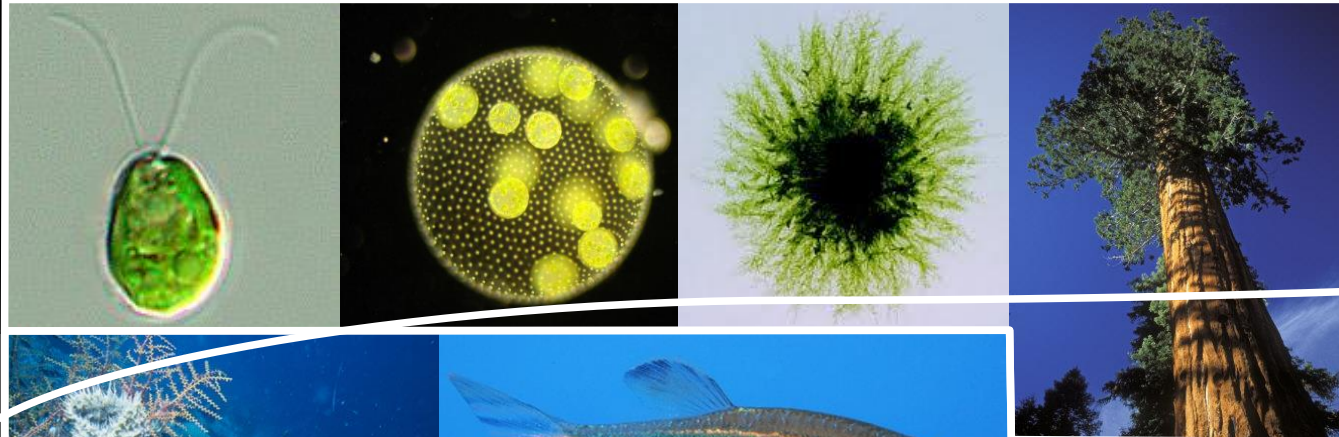
on the **developmental** level

Gradual evolution of complex life and biodiversity

Bacteria



Plants



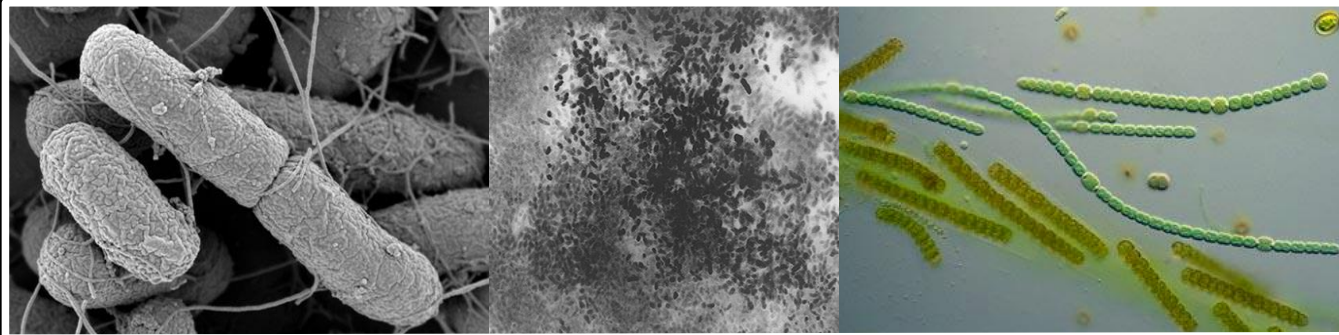
Animals



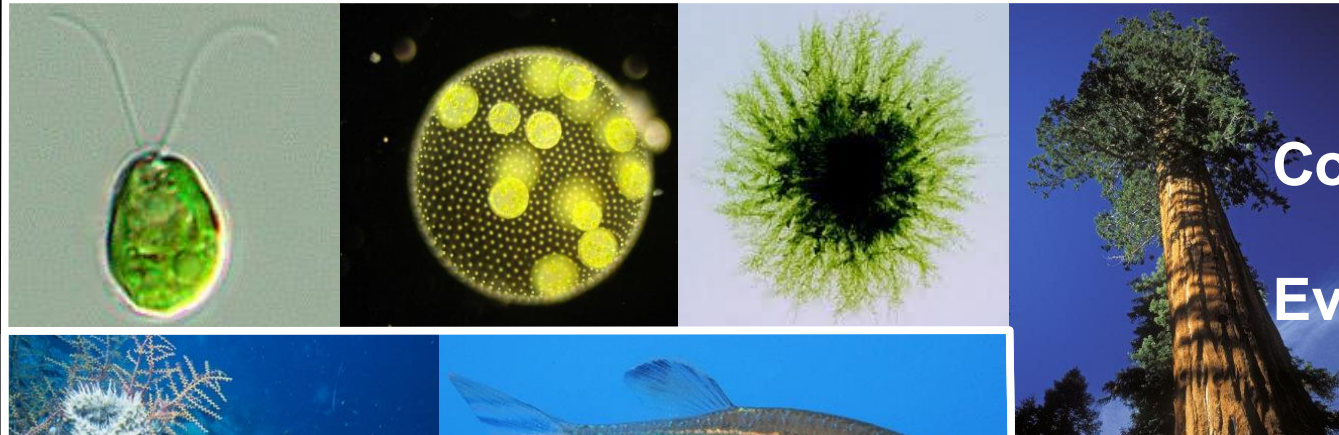
X
Embryo-
genesis

Gradual evolution of complex life and biodiversity

Bacteria



Plants

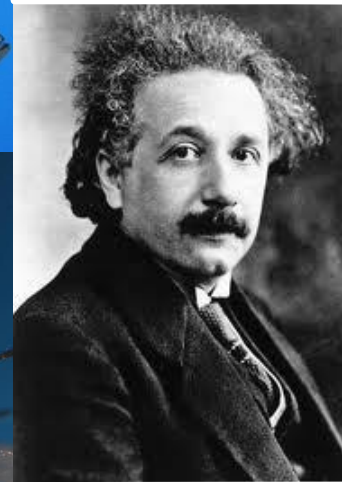


Comparative embryology

Evolutionary embryology

Evo-devo

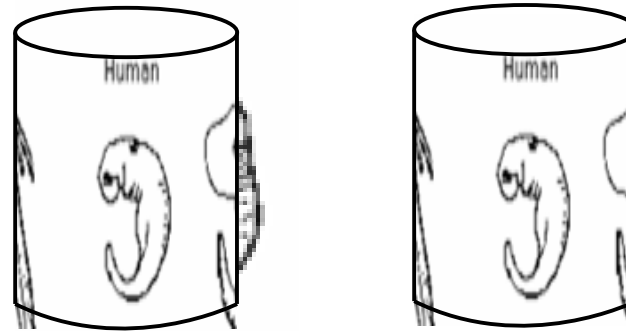
Animals



Comparative embryology - Baer's laws of embryology

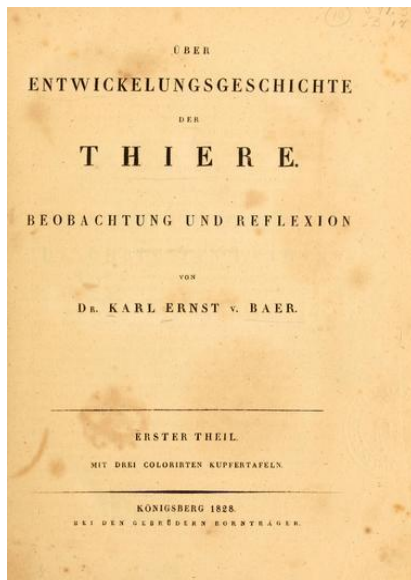


Karl Ernst von Baer
1792-1876



„Ich besitze zwei kleine Embryonen in Weingeist, für die ich versäumt habe die

*- K.E. von Baer, (1828), *Entwicklungsgeschichte der Thiere* (S. 221)*



- (1) General features of the embryo appear earlier than special features
- (2) Special characters develop from general characters
- (3) Embryos of different species progressively diverge from one another during ontogeny**
- (4) Embryos of one animal can never resemble the adult form of another animal, but only its embryo

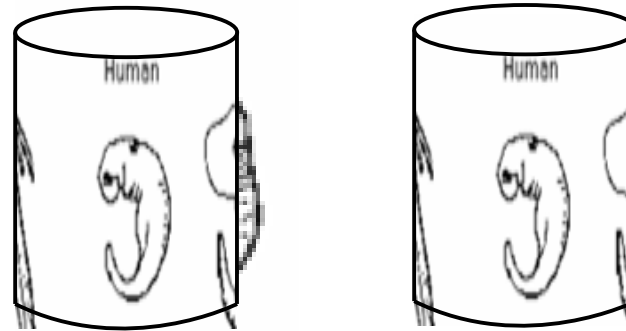
→ Existence of a stage in which embryos of different vertebrate species could not be distinguished

→ One of the milestone discoveries in developmental biology!

Comparative embryology - Baer's laws of embryology

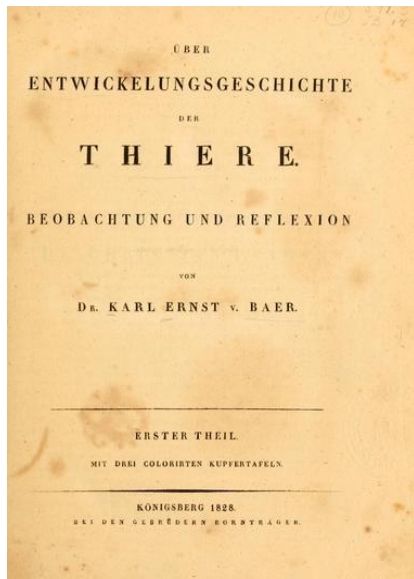


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„Ich besitze zwei kleine Embryonen in Weingeist, für die ich versäumt habe die

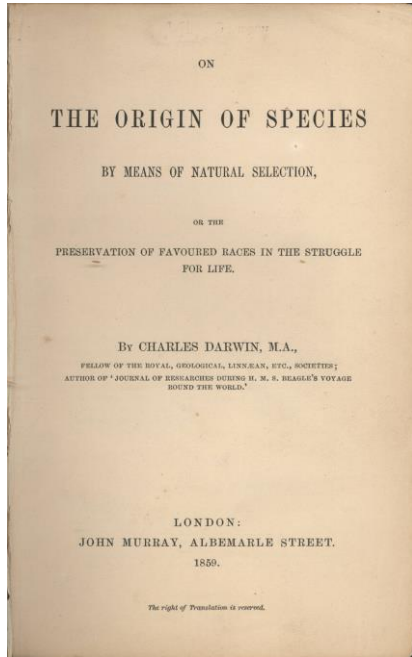
*- K.E. von Baer, (1828), *Entwicklungsgeschichte der Thiere* (S. 221)*



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→ **Pre-Darwin → no connection to Darwinian evolution**

Post-Darwin



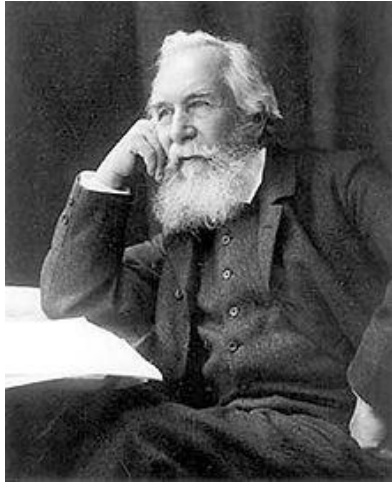
Darwin delivered the connection of:

- Theory of evolution
- Common origin of species

} Mechanism:
Natural Selection

Connection between origin / descent, systematics, and ontogeny?

Post-Darwin - Ernst Haeckel

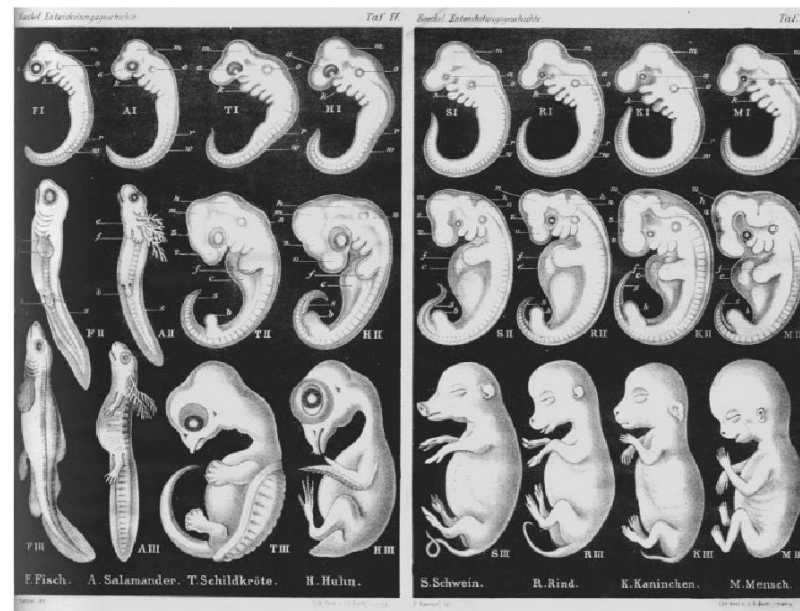


Ernst Haeckel
1834 - 1919

Haeckel's biogenetic law (1866):

“Die Ontogenese ist die kurze und schnelle Recapitulation der Phylogenese”

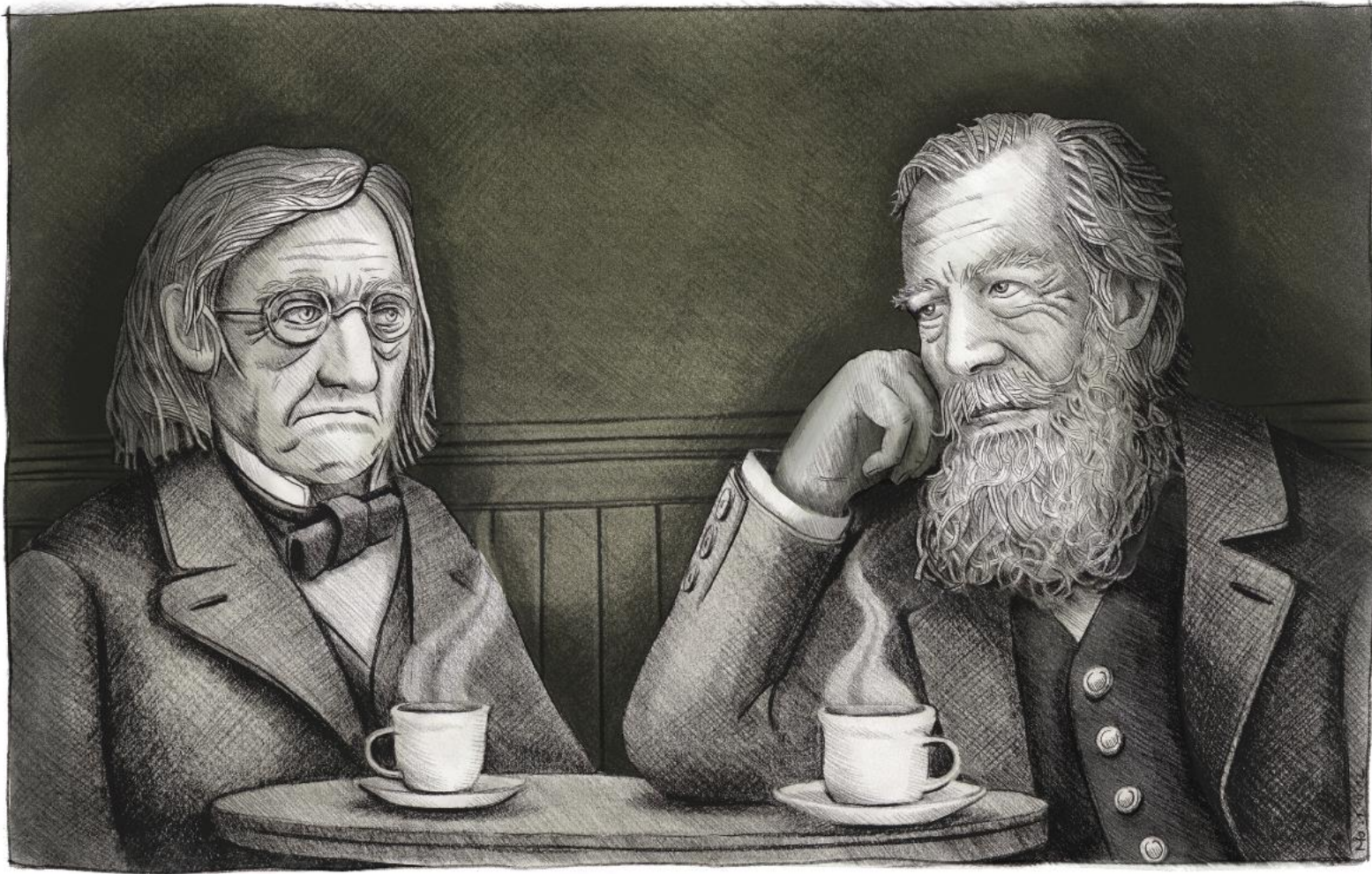
- Mostly wrong and based on purposely falsified data:



- Groundwork for social darwinism and nazi racial ideology

BUT: for the first time approaching developmental biology from an evolutionary perspective

Comparative embryology – 19th century



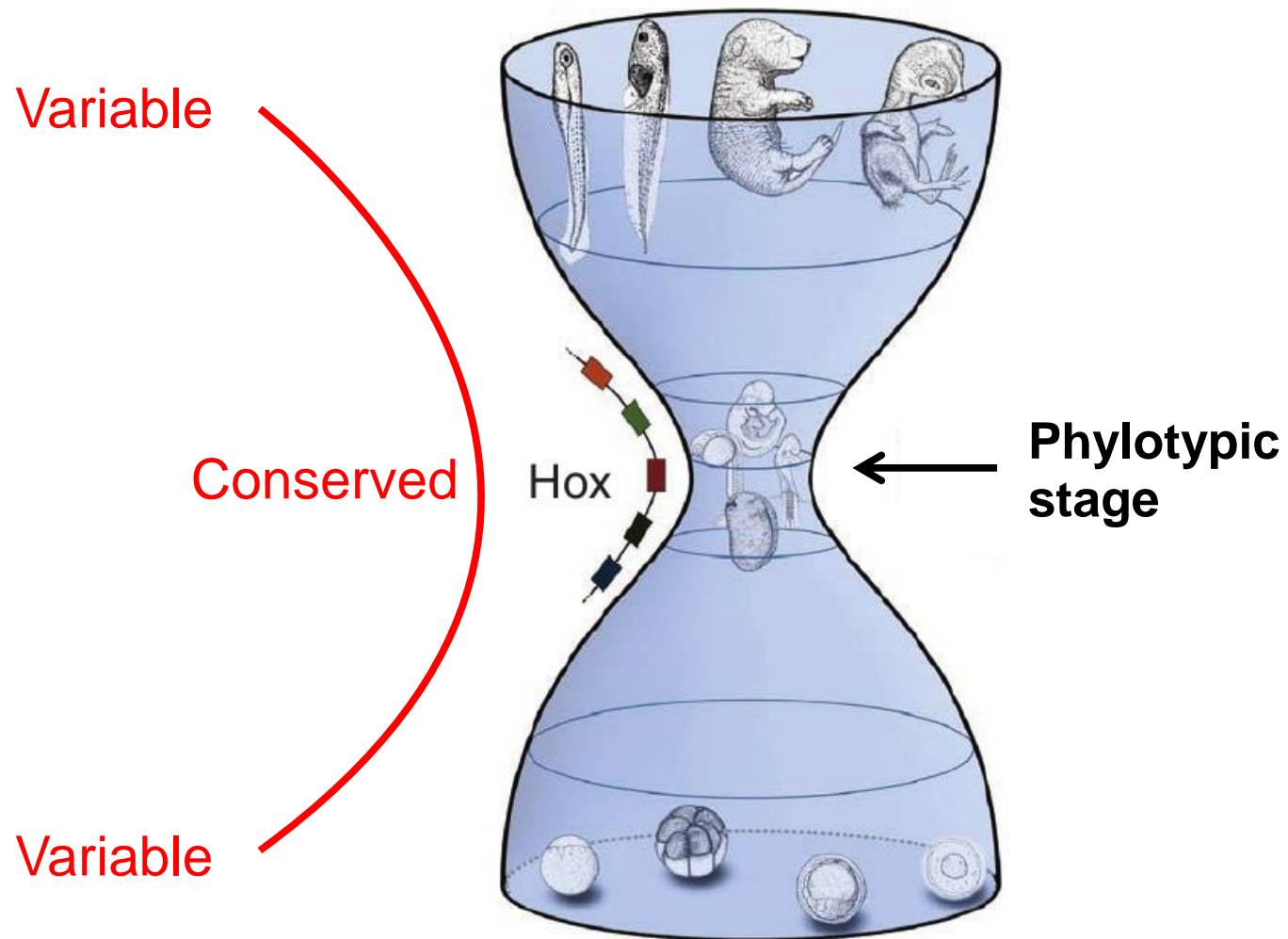
von Baer's contribution

- Existence of a stage of max. morphological conservation during embryogenesis of different vert

Haeckel's contribution

- Use of such data for phylogeny reconstruction

Comparative embryology – The hourglass model



.21st century – the genomic level

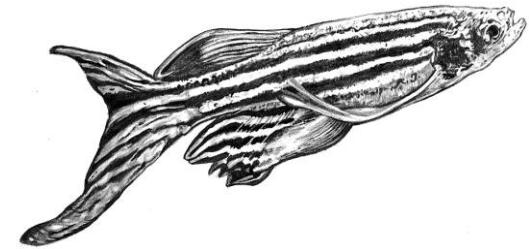
LETTER

doi:10.1038/nature09632

A phylogenetically based transcriptome age index mirrors ontogenetic divergence patterns

Tomislav Domazet-Lošo^{1,2} & Diethard Tautz¹

Nature (2010)



LETTER

doi:10.1038/nature09634

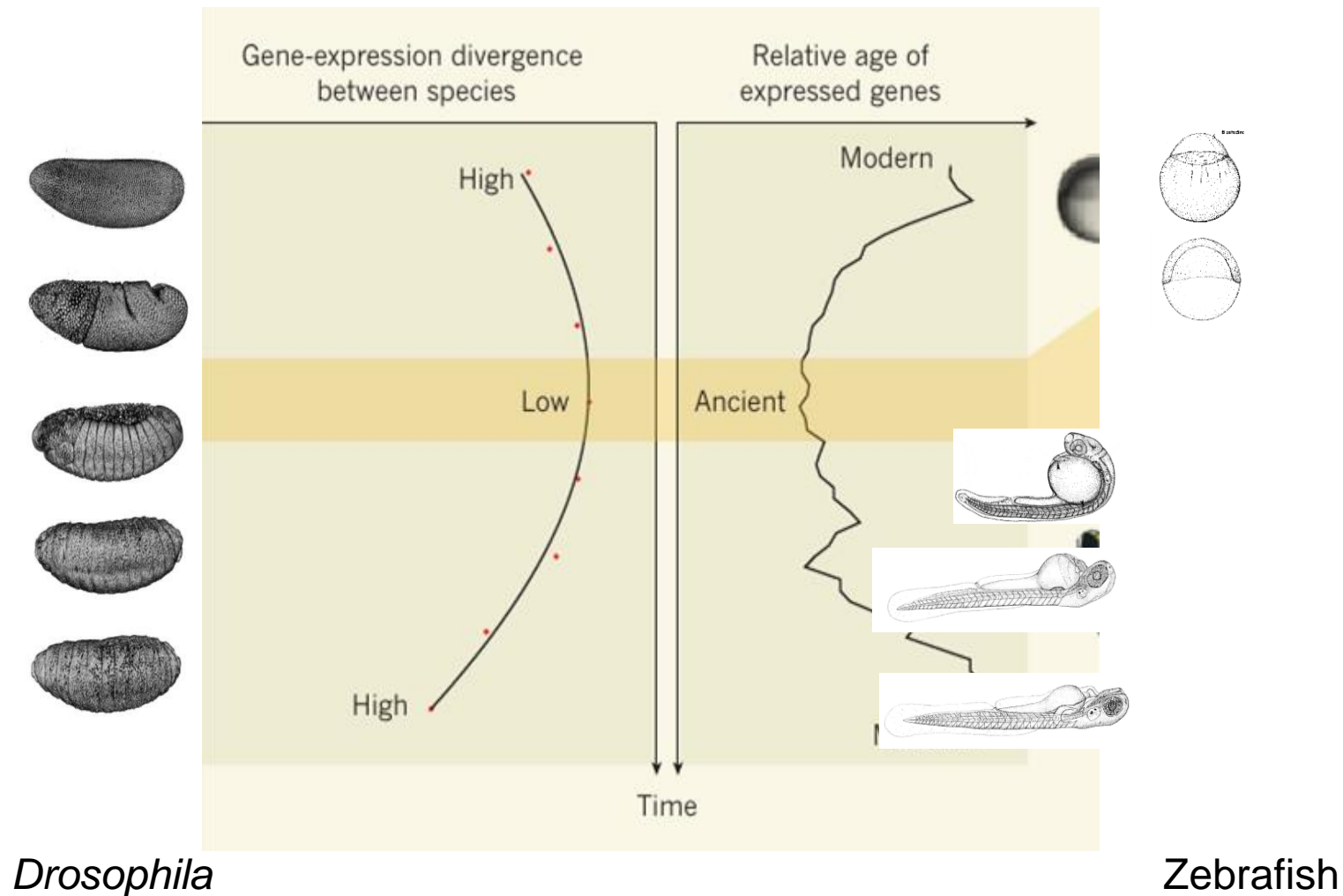
Gene expression divergence recapitulates the developmental hourglass model

Alex T. Kalinka^{1*}, Karolina M. Varga^{1*†}, Dave T. Gerrard², Stephan Preibisch¹, David L. Corcoran³, Julia Jarrells¹, Uwe Ohler³, Casey M. Bergman² & Pavel Tomancak¹

Nature (2010)

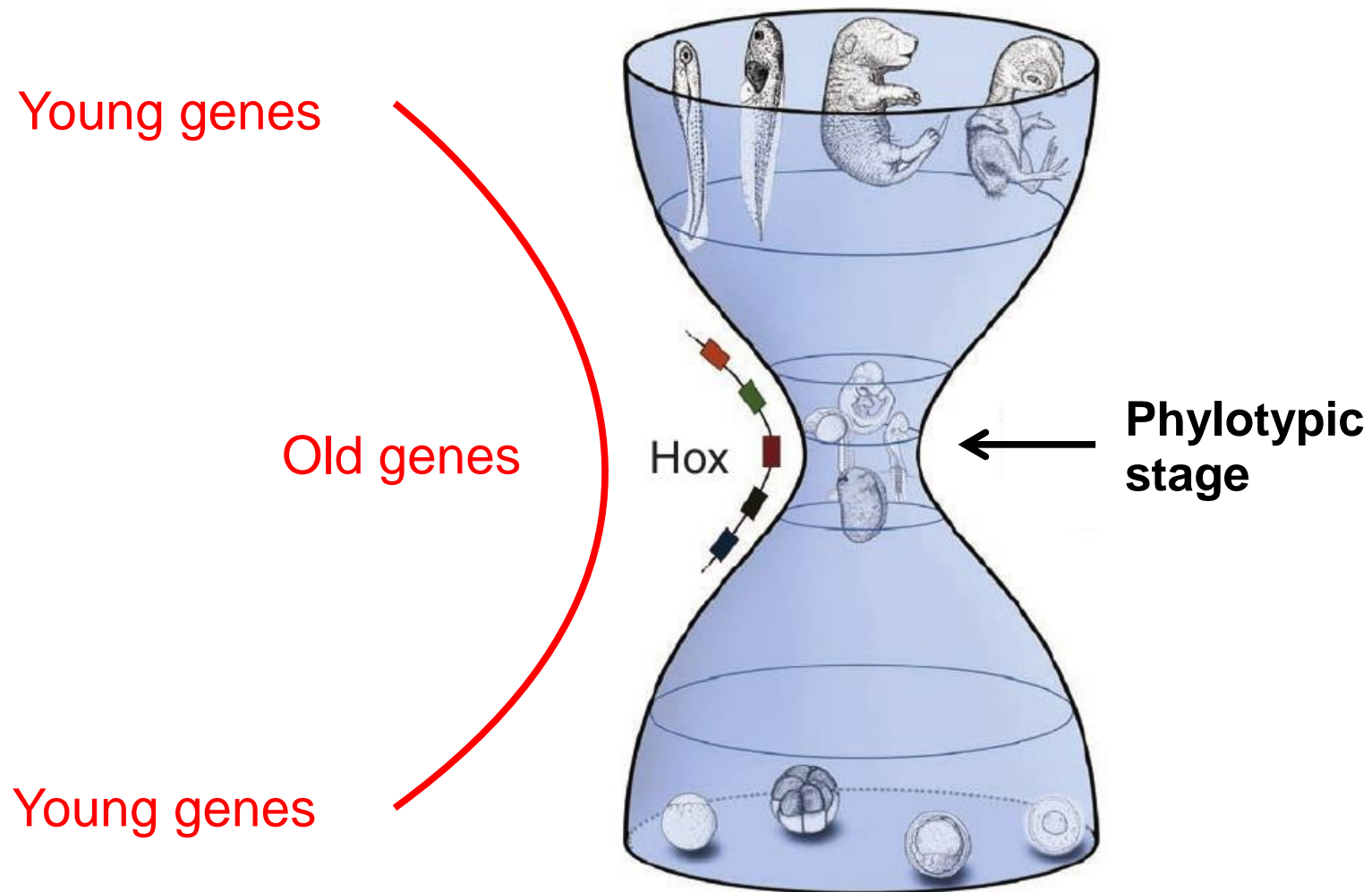


21st century – the genomic level



1. Morphology AND transcriptomes follow an hourglass pattern!
2. Max. conservation of the transcriptomes observed at the morphological phylotypic stage!

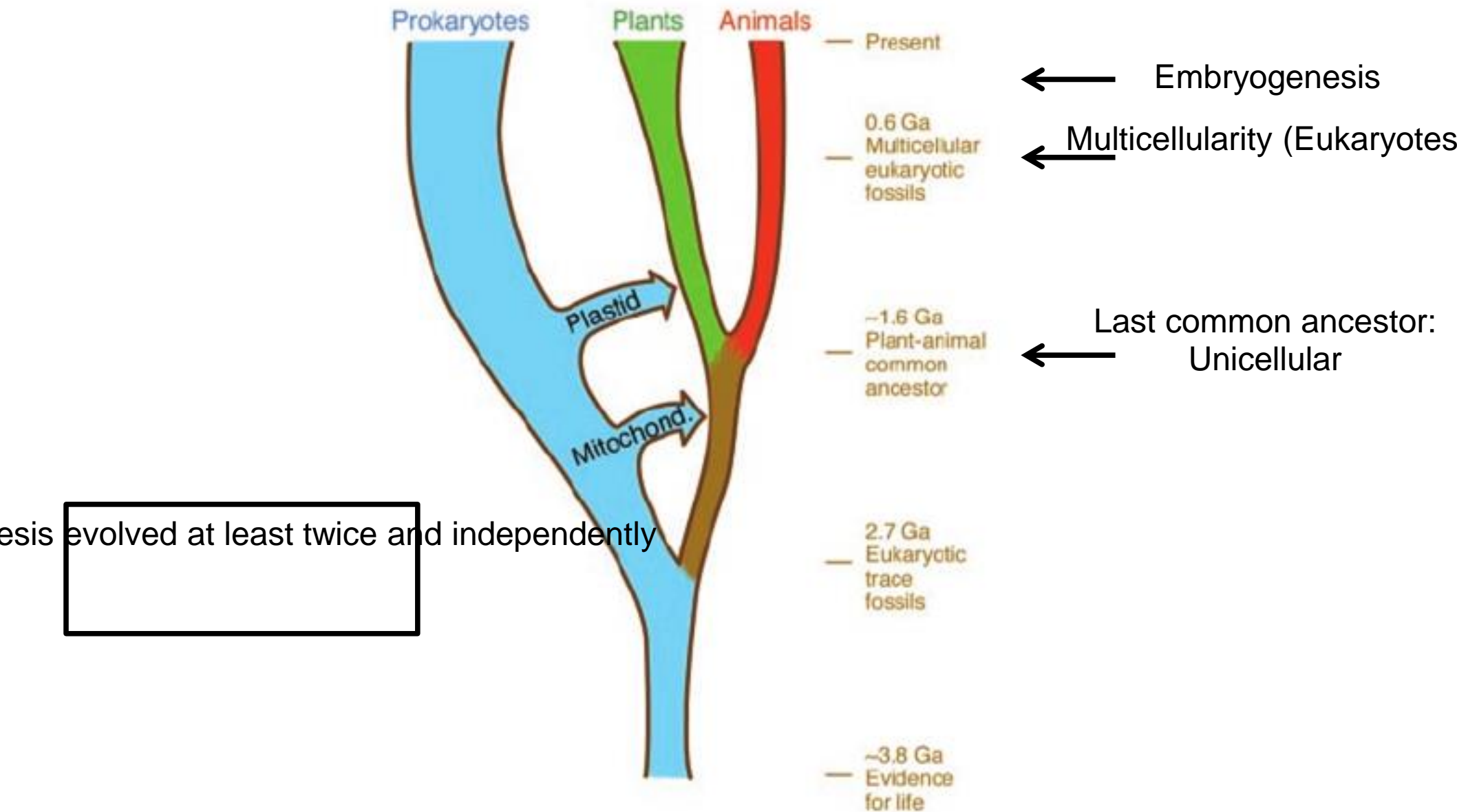
21st century – the genomic level





Evolution of embryogenesis

Unicellular life → Multicellular life → Embryogenesis → Complex life



What about plants?

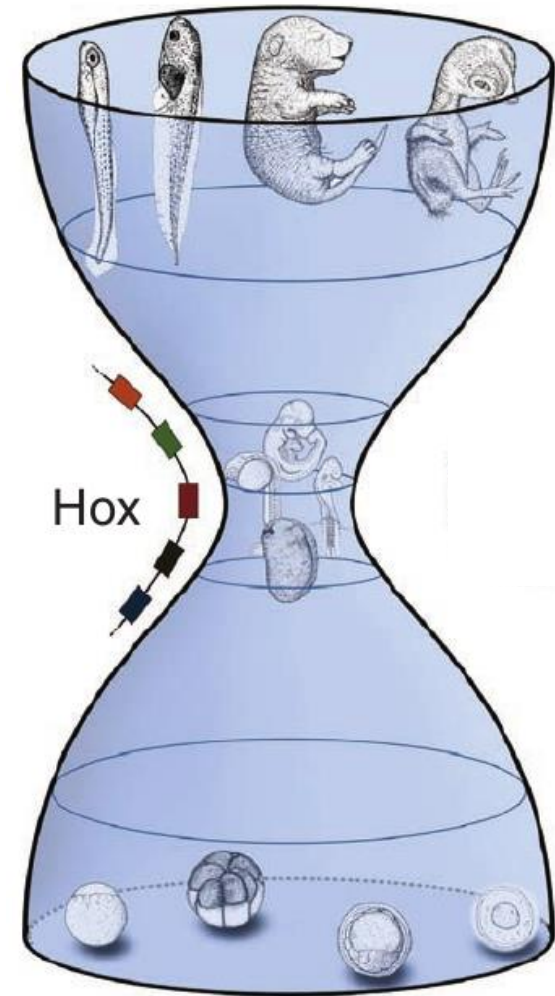
Young genes

.Is there a
transcriptional
hourglass in plants?

.Or any other pattern?

Old genes

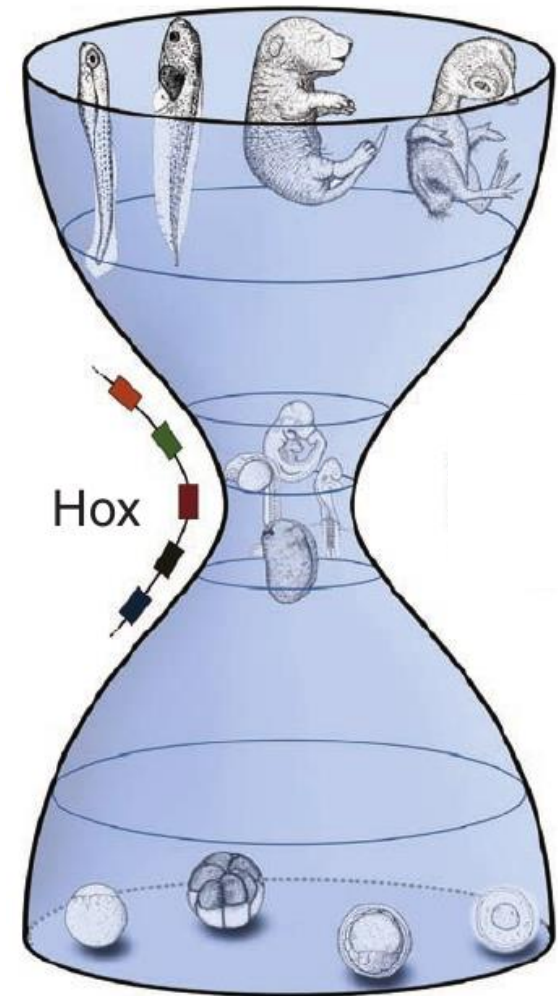
Young genes



What about plants?

Two pieces of information:

1. Evolutionary age for each gene
2. Transcriptomes of various embryo stages



Step 1: Phylogenetic information for each gene - Phylostratigraphy

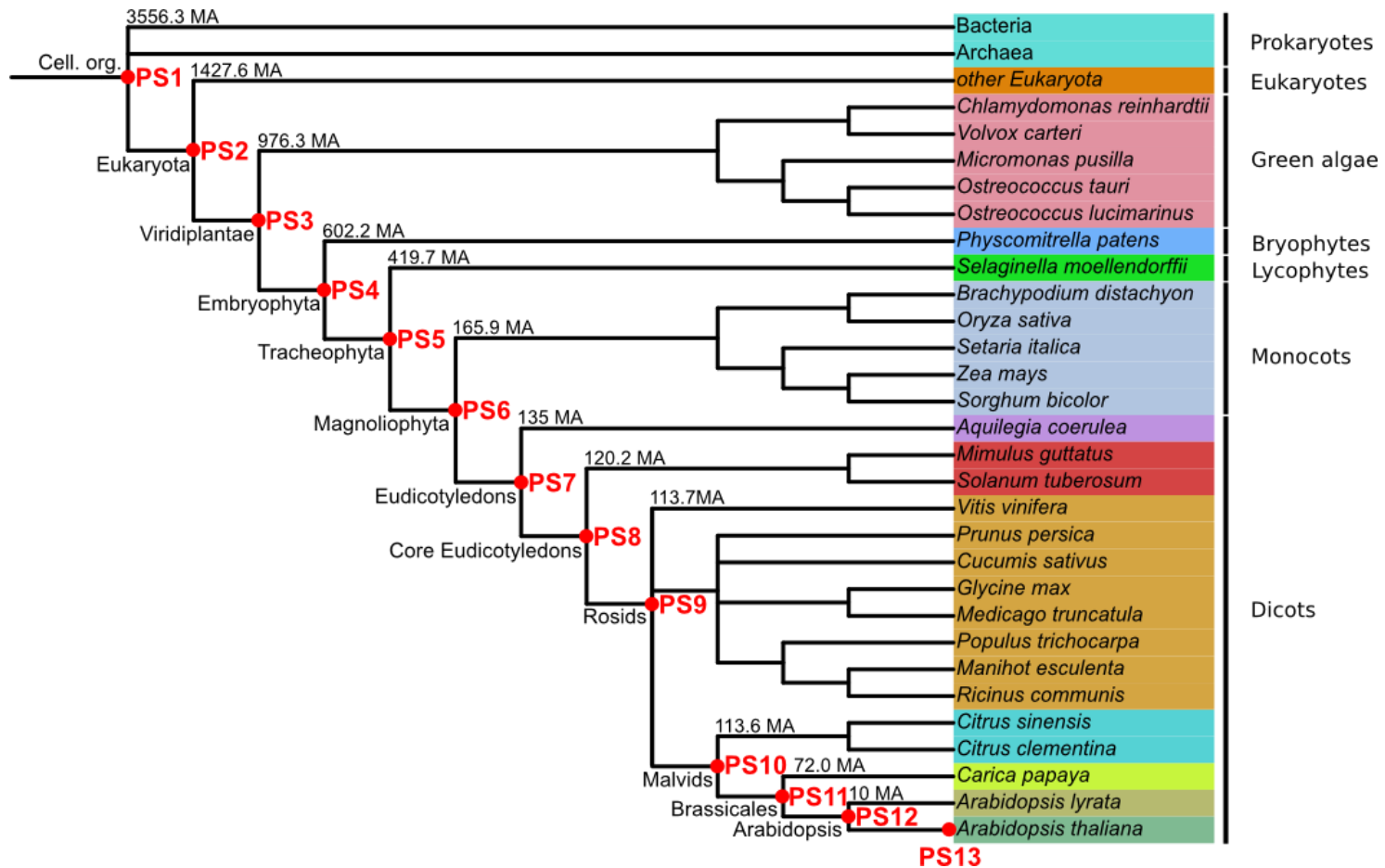
Domazet-Lošo et al., 2007, *Trends Genet*

Reconstruct the phylogeny of *the species of interest* along the tree of life

→ restricted to completely sequenced organisms

Categorize each new node/branch point in the phylogeny to a **phylostratum**

→ from young to old



Step 1: Phylogenetic information for each gene - Phylostratigraphy

Domazet-Lošo et al., 2007, *Trends Genet*

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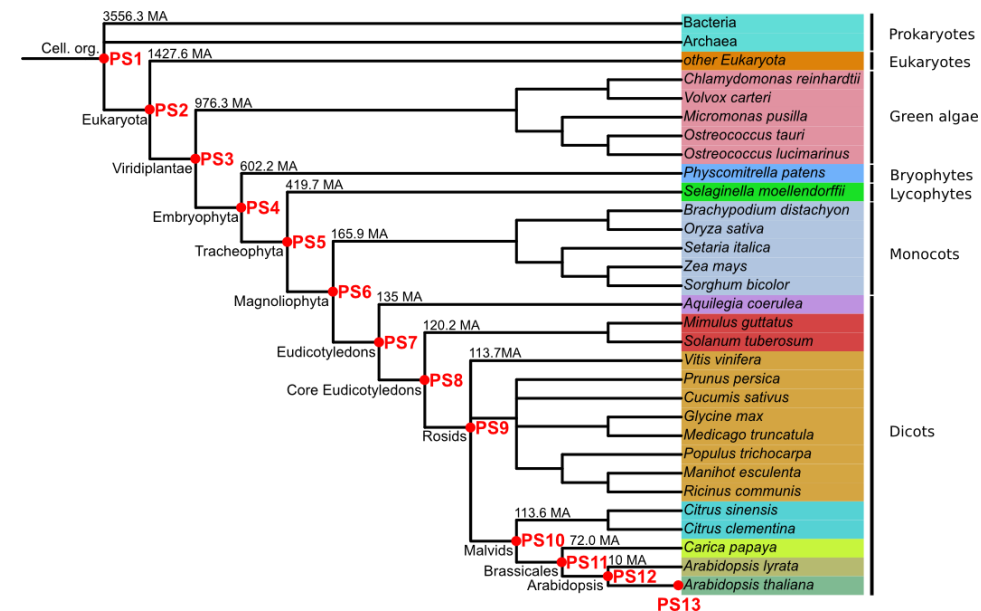
Categorize each new node/branch point in the phylogeny to a **phylostratum**

→ from young to old

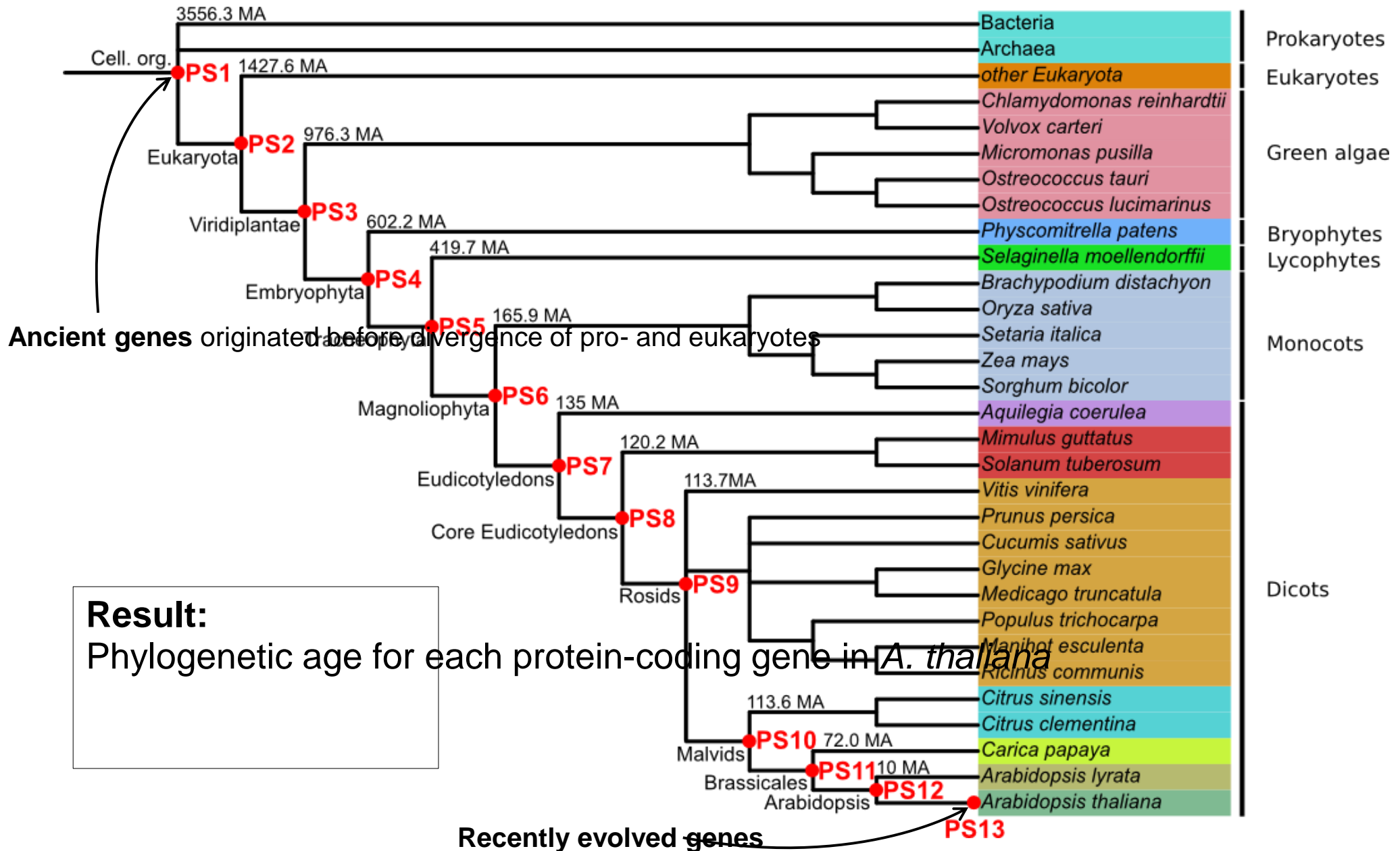
Construct a database with all sequenced genomes in the phylogeny

Blast each gene/protein of *the species of interest* to this database

Sort genes into phylostrata (PS1 through PSn)



Reconstructed phylogeny of *A. thaliana*



Step 2: How can we apply phylostratigraphy to transcriptional data?

adapted from Domazet-Lošo and Tautz, 2010, *Nature*



Introduce expression intensity of a gene as a weight for its phylogene



Sum over all genes

=



Transcriptome Age Index

Definition TAI:

The mean evolutionary age of a transcriptome, where the evolutionary age (PS) of each gene is weighted (TAI)

Step 2: How can we apply phylostratigraphy to transcriptional data?

adapted from Domazet-Lošo and Tautz, 2010, *Nature*



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Sum over all genes

=

← **Transcriptome Age Index**
(TAI)

Definition TAI:

The mean evolutionary age of a transcriptome, where the evolutionary age (PS) of each gene is weighted

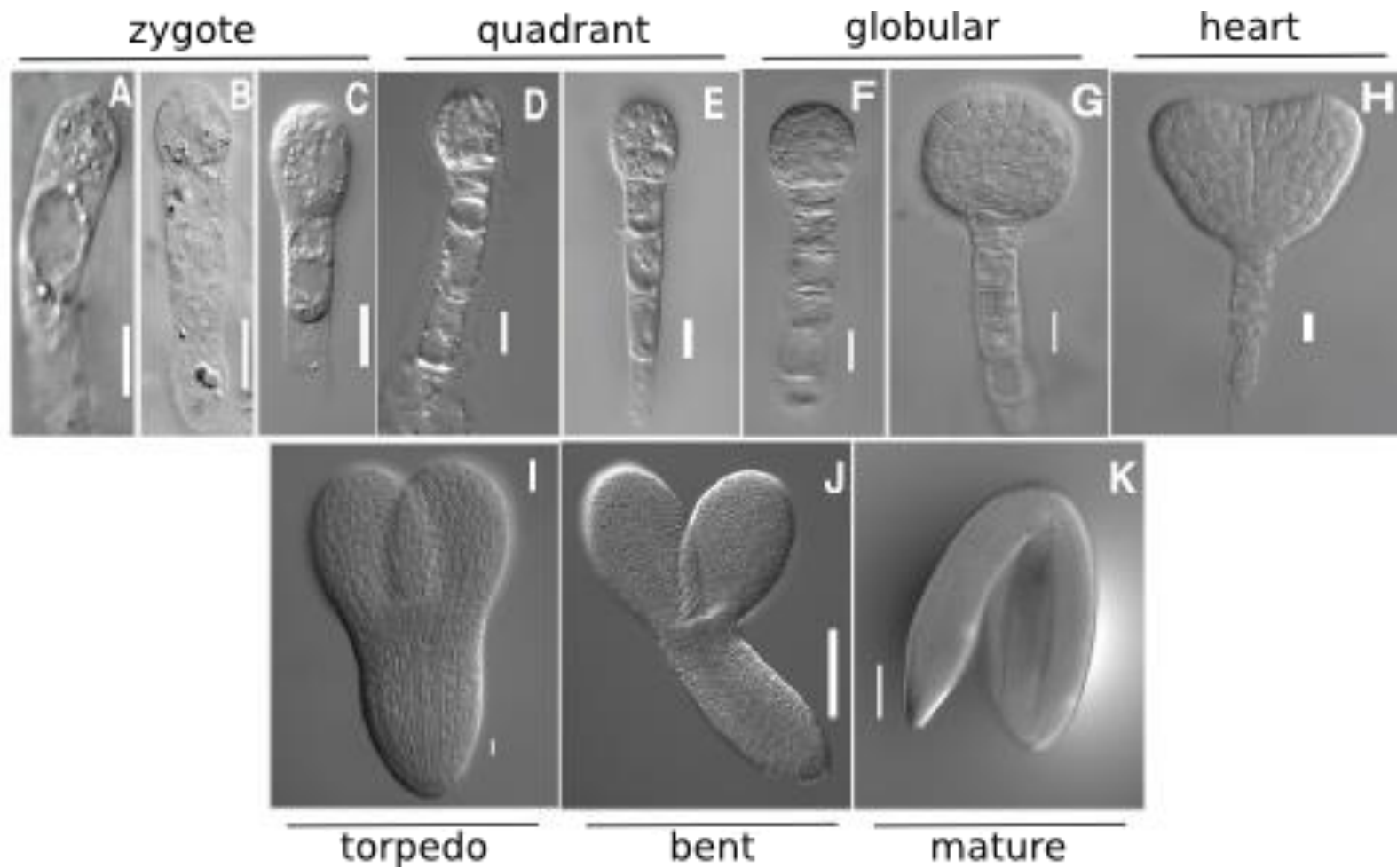


Goal:

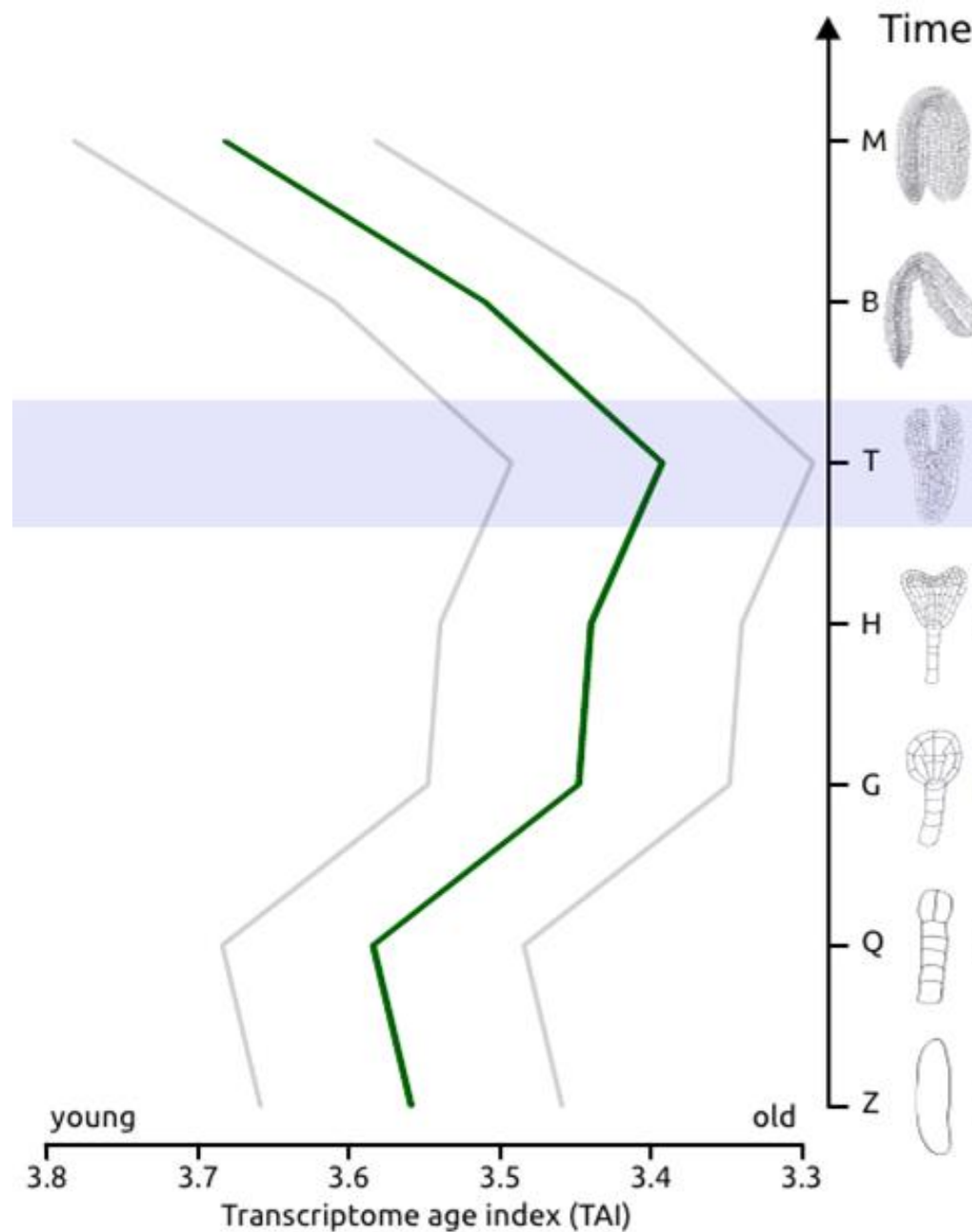
Compute TAI across multiple embryo stages

→ compare TAI profile to hourglass pattern

Embryogenesis in *Arabidopsis thaliana*



Pattern of *A. thaliana* embryogenesis is reminiscent of the developmental hour

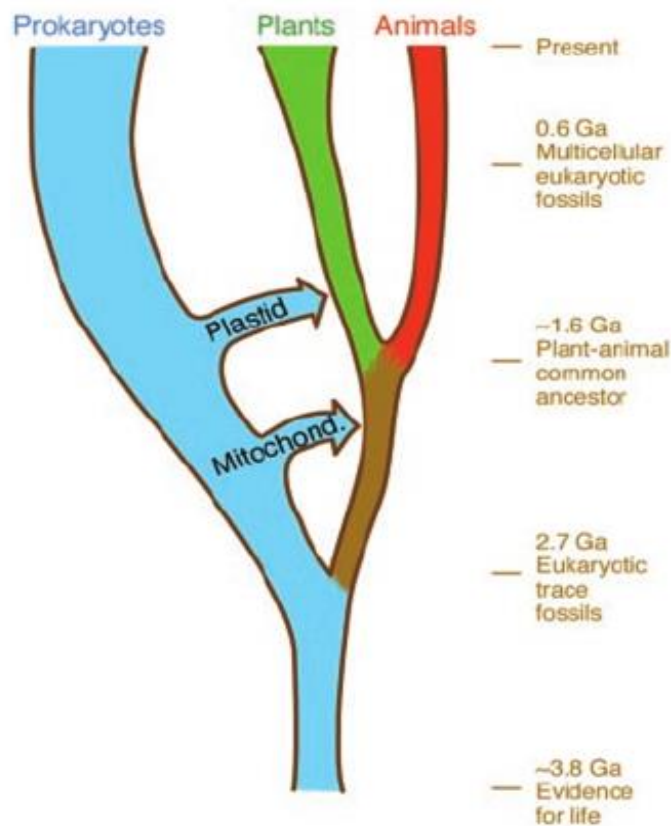


Intermediate conclusions

Embryogenesis evolved twice and independently in animals and plants

Embryogenesis morphologically and genetically different in animals and plants

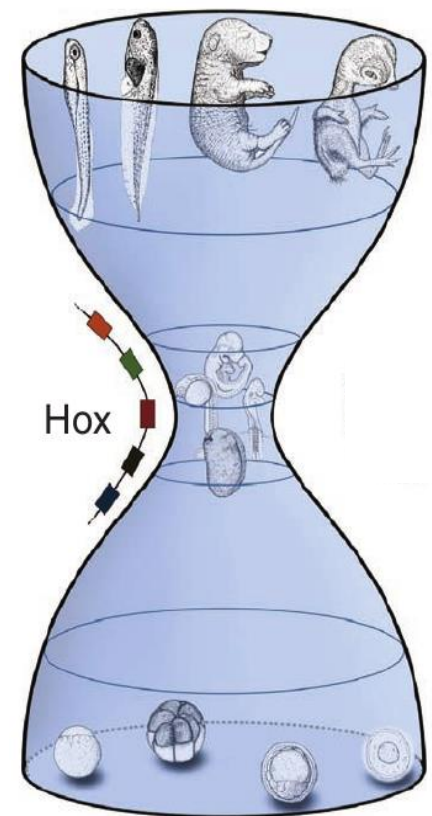
But transcriptomes of animals and plants show the same hourglass pattern



Young genes

Old genes

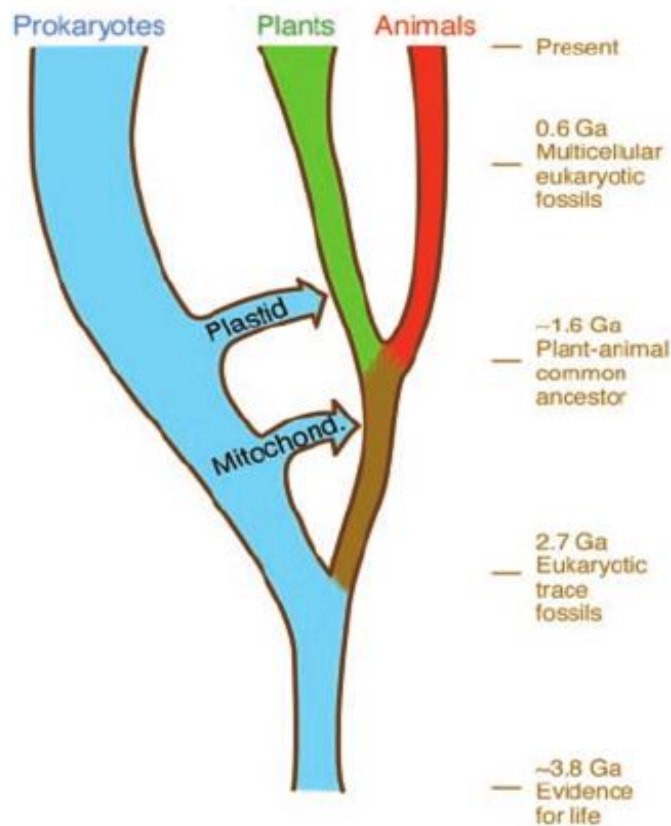
Young genes



Fundamental question

Is the hourglass pattern

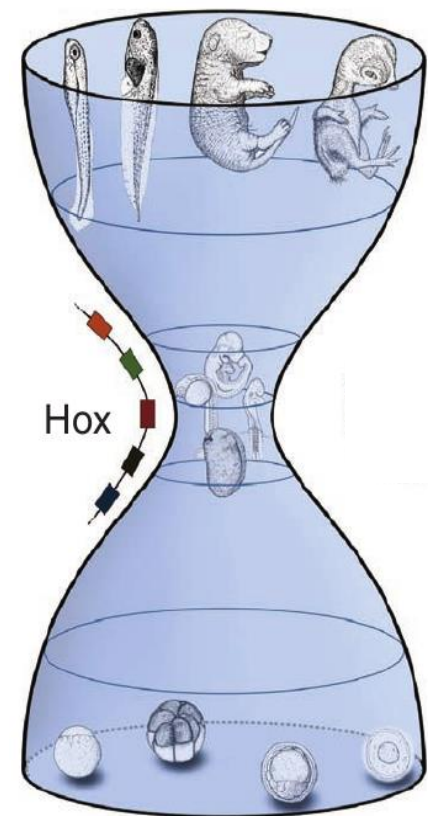
- **an evolutionary relic** and possibly not functional anymore?
- **or actively maintained** and possibly still functional?



Young genes

Old genes

Young genes



Transcriptome divergence index - TDI

TDI based on sequence divergence



Sequence comparison of orthologous genes in related species



Sequence divergence = Ka/Ks

Ka = nonsyn. mutations
 Ks = syn. mutations

Evolutionary conserved genes
→ sequence divergence small

Evolutionary variable genes
→ sequence divergence large



Genome-wide → Ka/Ks ratios for all orthologous gene pairs

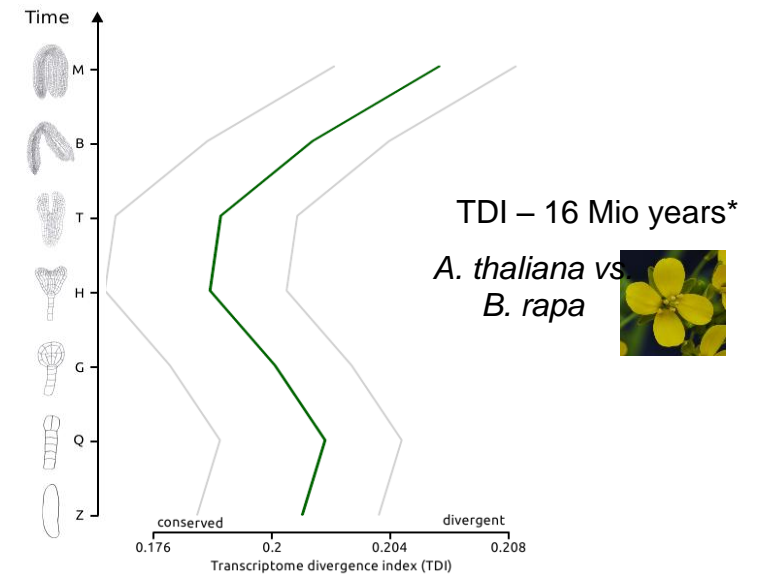
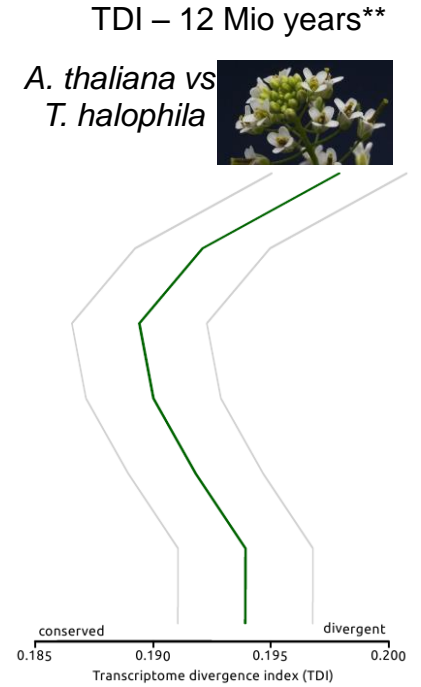
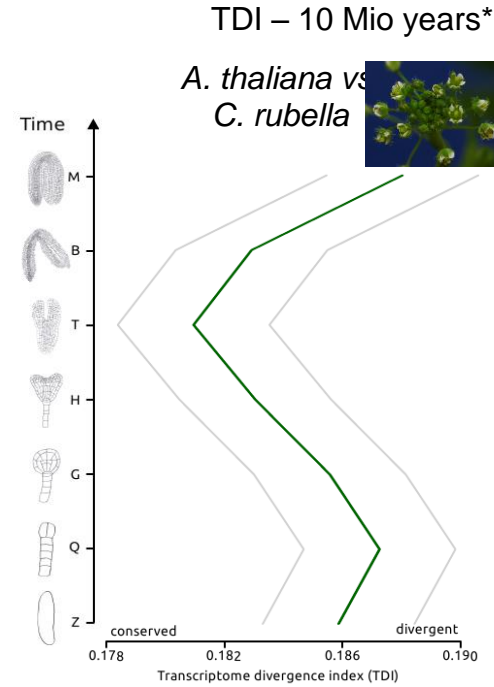
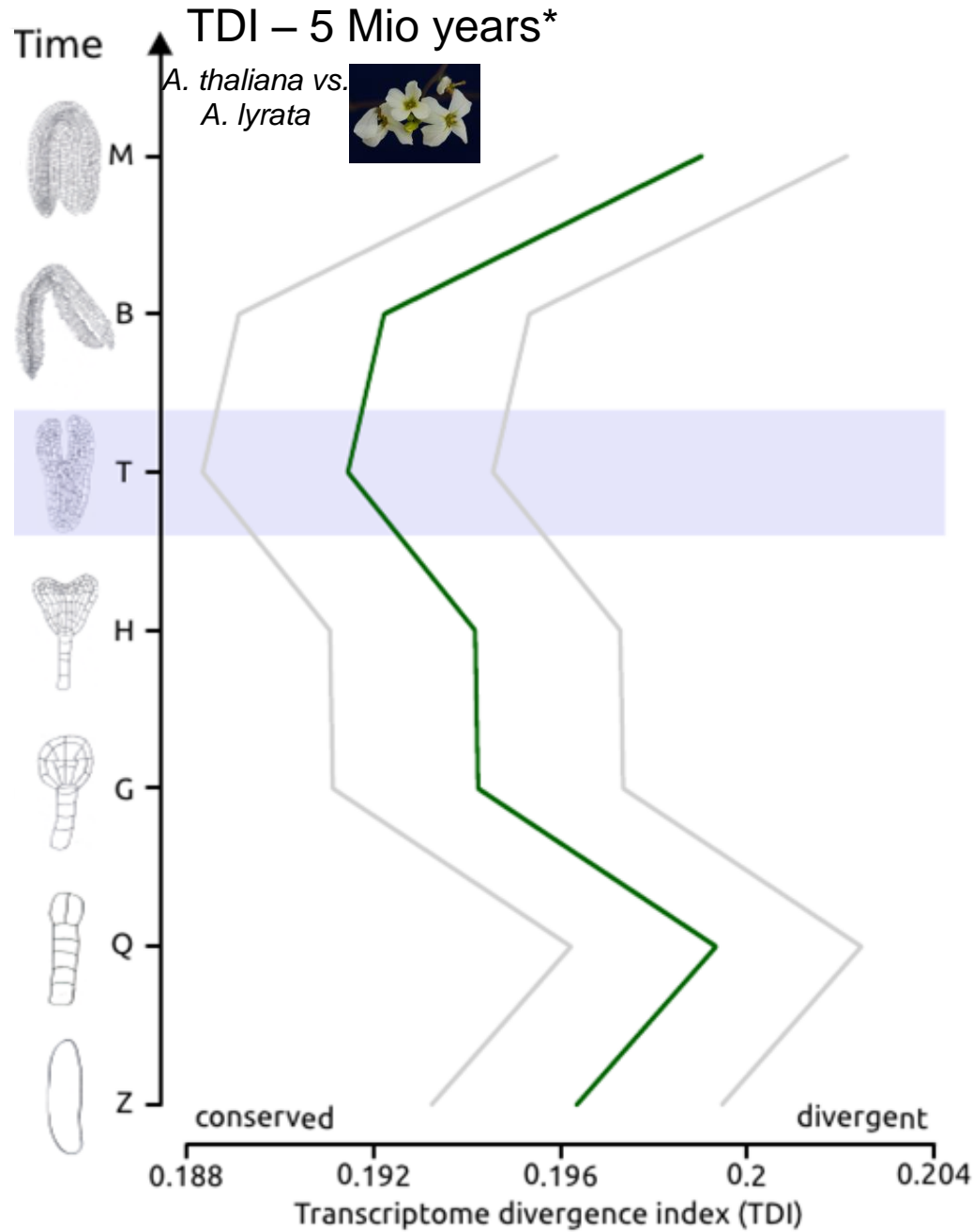
A. thaliana vs. *A. lyrata*



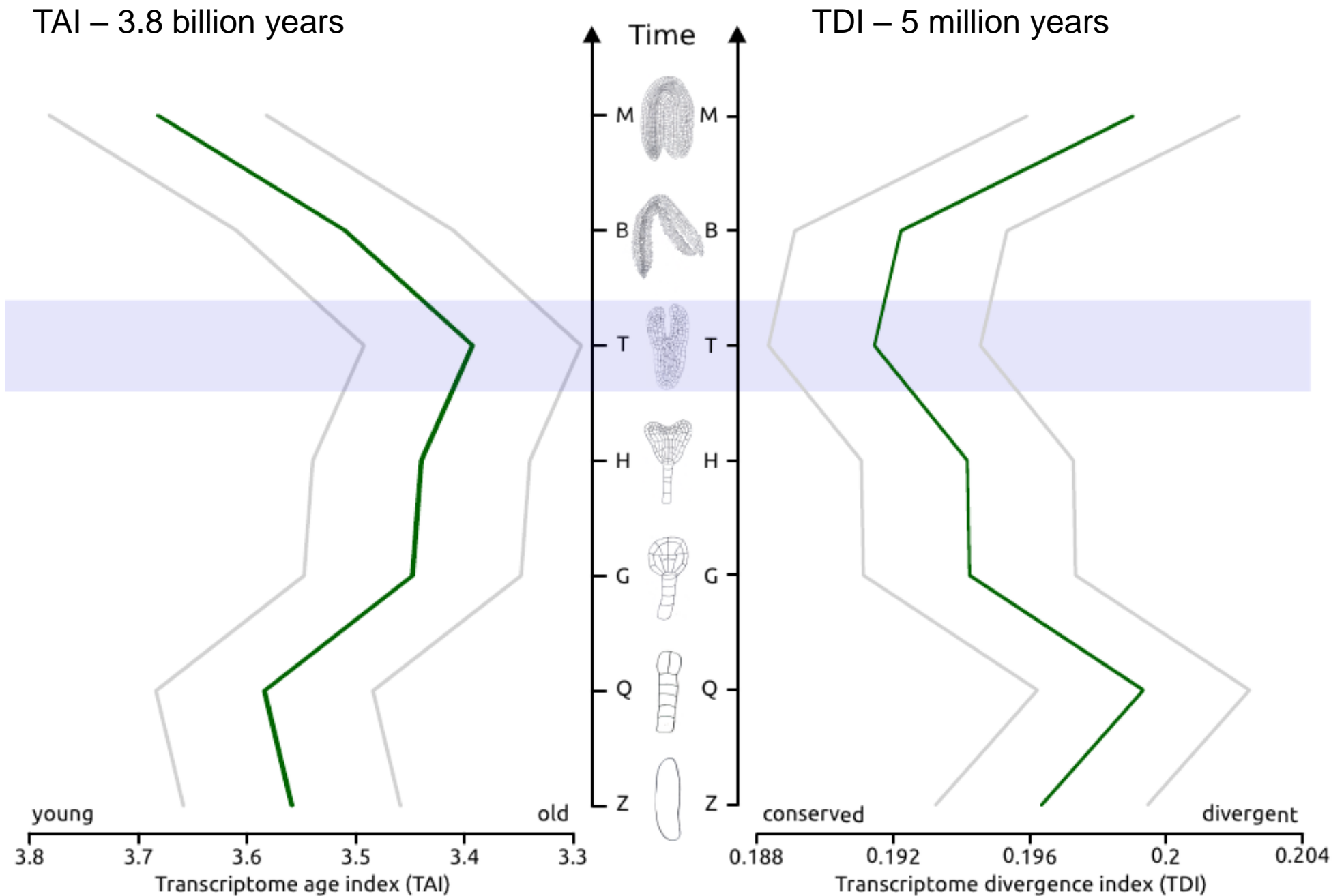
**Ka/Ks
instead of PS**



TDI patterns



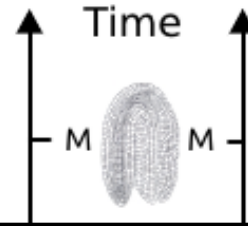
Developmental hourglass for *A. thaliana* embryogenesis



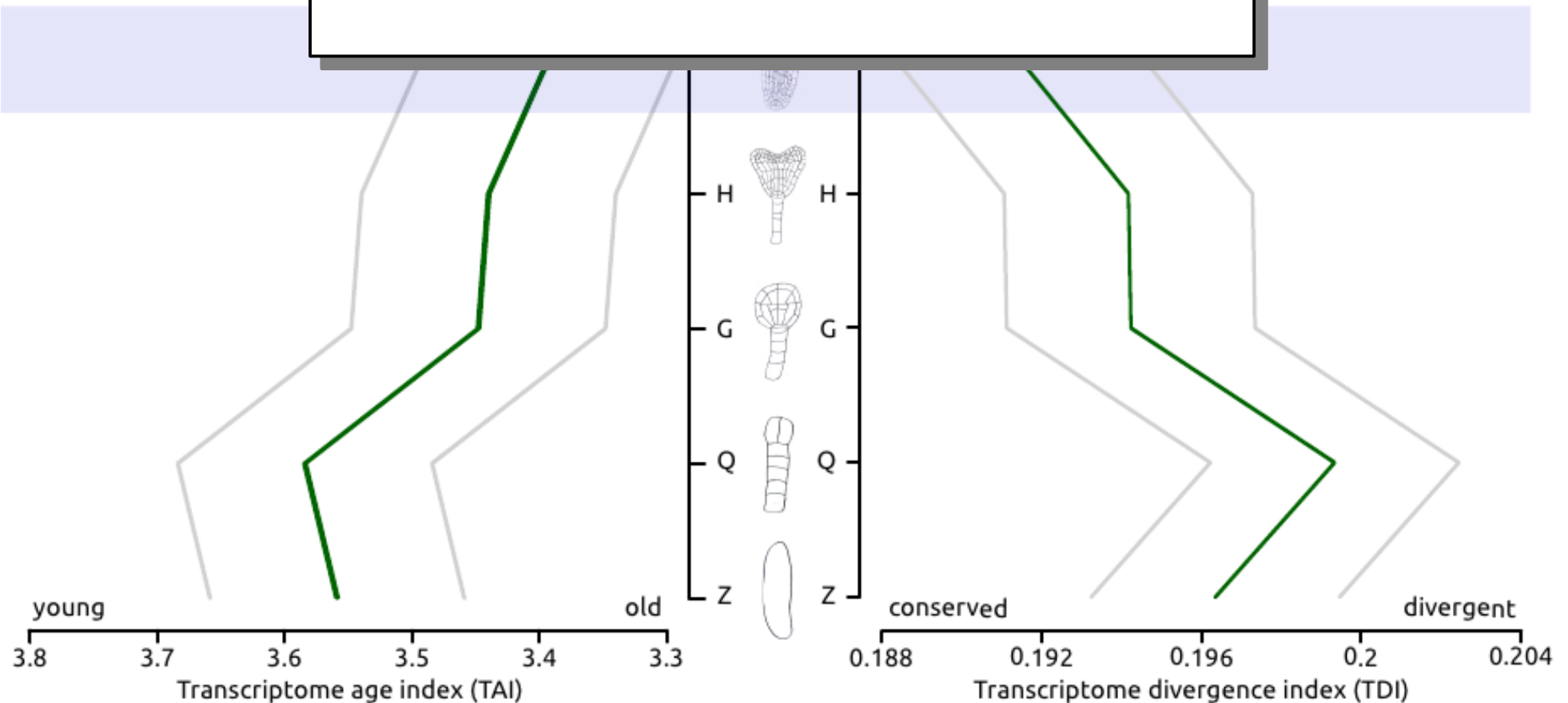
Developmental hourglass for *A. thaliana* embryogenesis

TAI – 3.8 billion years

TDI – 5 million years



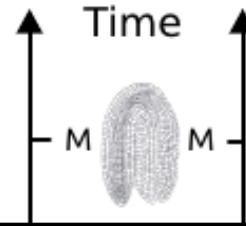
DI hourglass pattern suggests active maintenance and possibly functionality



Developmental hourglass for *A. thaliana* embryogenesis

TAI – 3.8 billion years

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DI hourglass pattern suggests active maintenance and possibly functionality

What about animals?

young

old

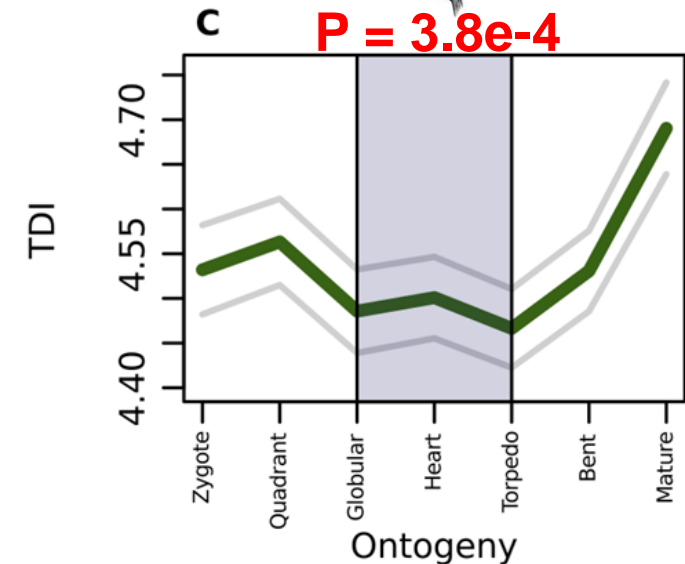
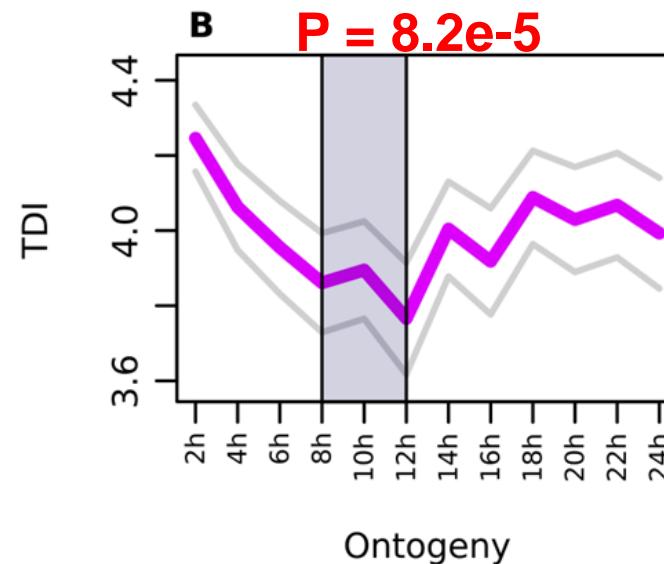
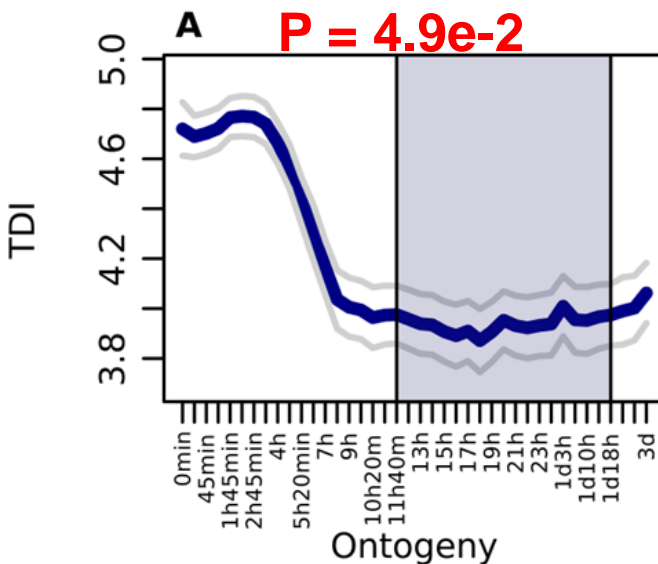
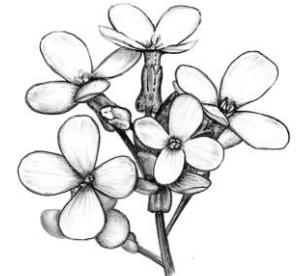
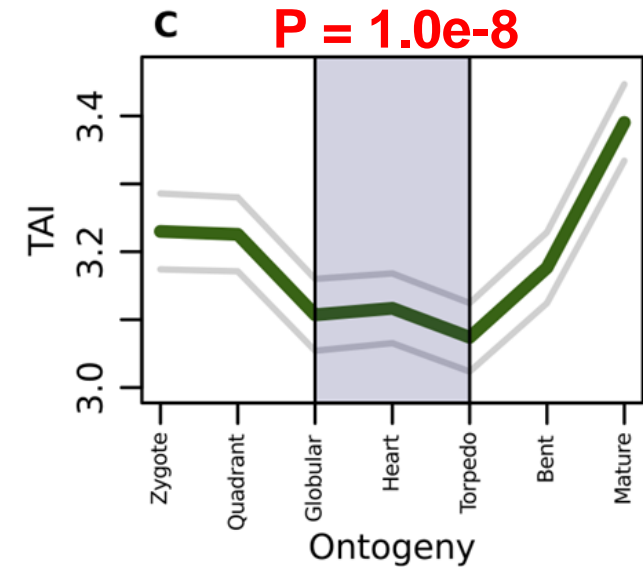
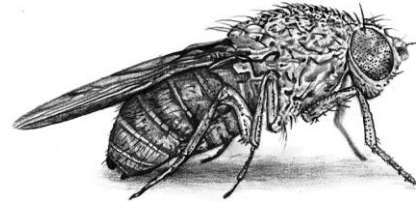
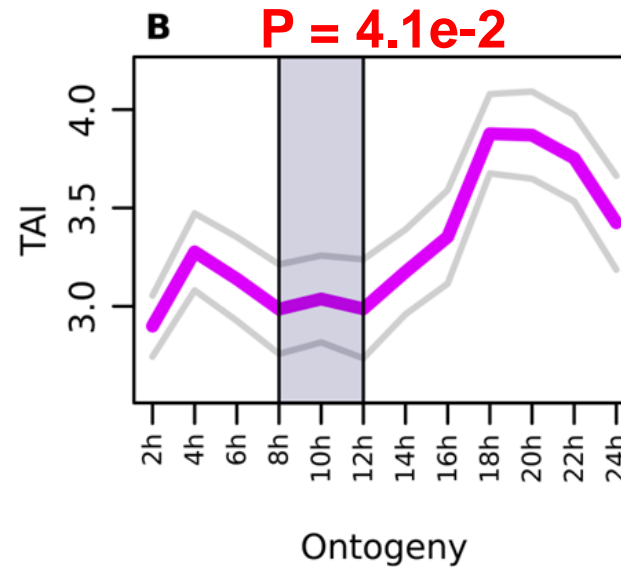
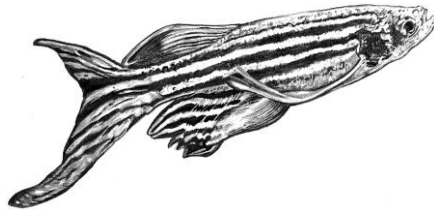
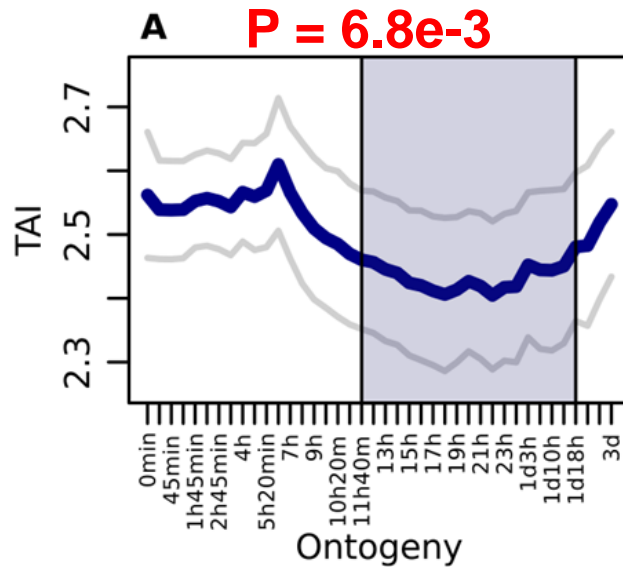
conserved

divergent

3.8 3.7 3.6 3.5 3.4 3.3
Transcriptome age index (TAI)

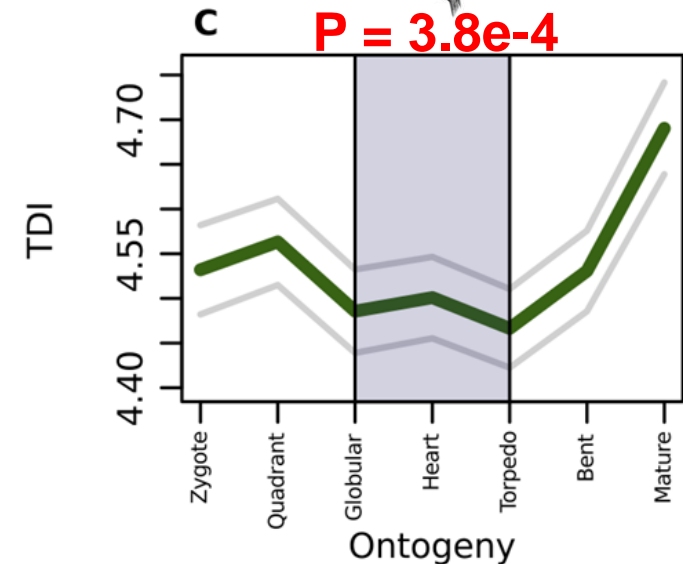
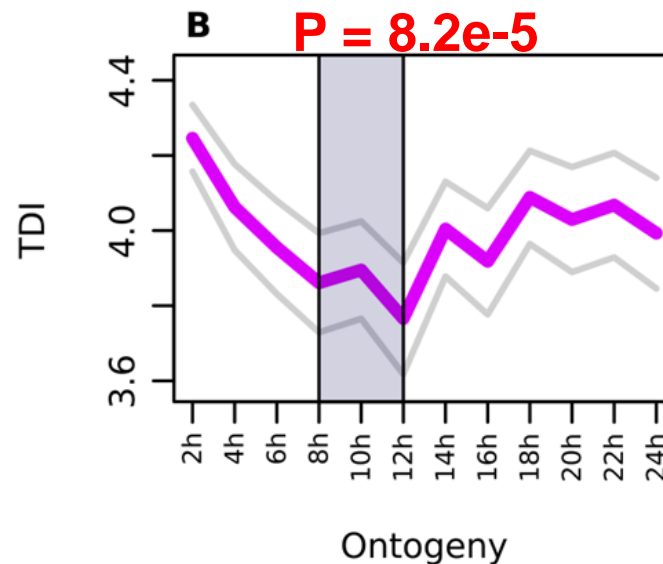
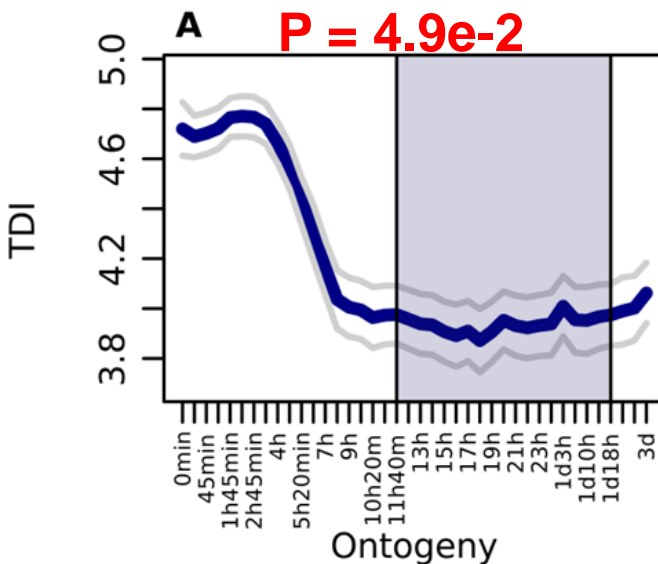
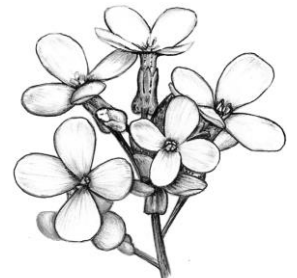
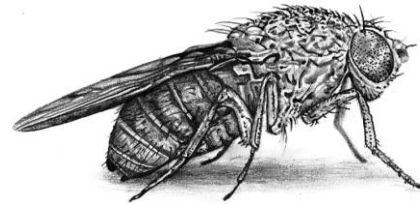
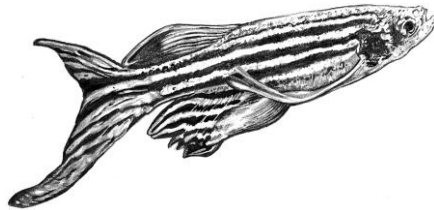
0.188 0.192 0.196 0.2 0.204
Transcriptome divergence index (TDI)

TAI and TDI for embryonic transcriptomes across kingdoms



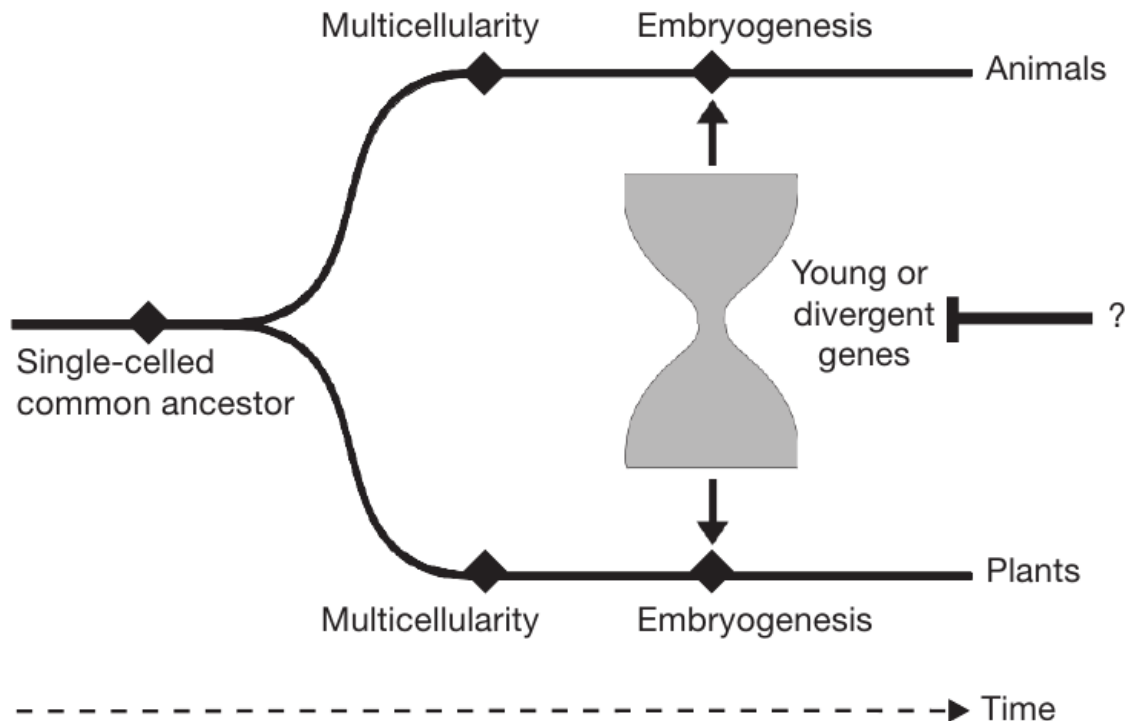
TAI and TDI for embryonic transcriptomes across kingdoms

glass patterns suggest active maintenance and possibly functionality in both

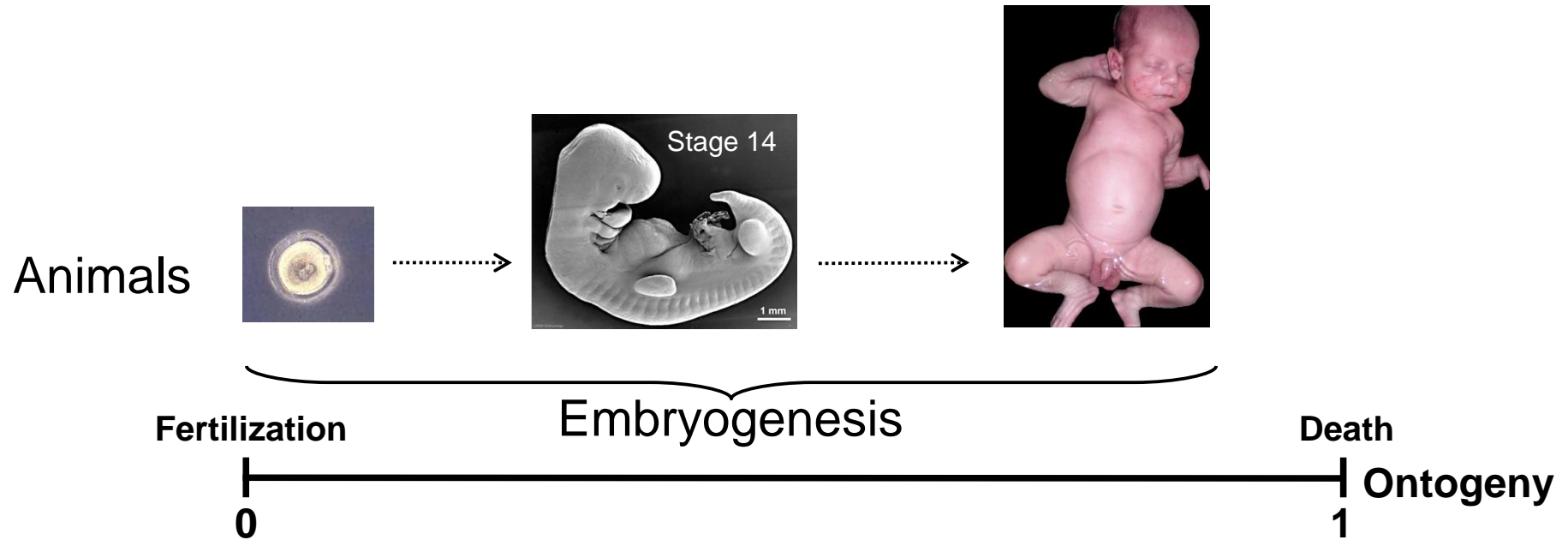


Intermediate conclusions

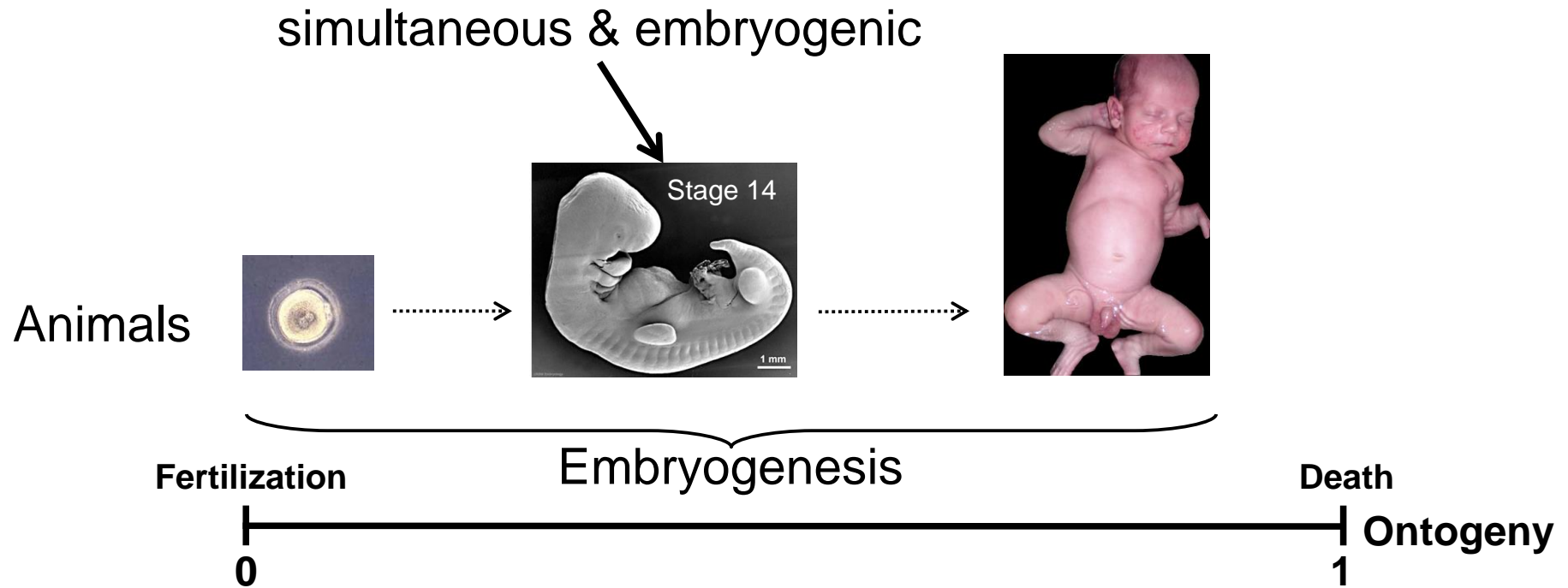
- .Nature invented embryogenesis twice and independently
- .Embryogenesis very different in plants and animals
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- .essential
- .actively maintained
- .possibly still functional?



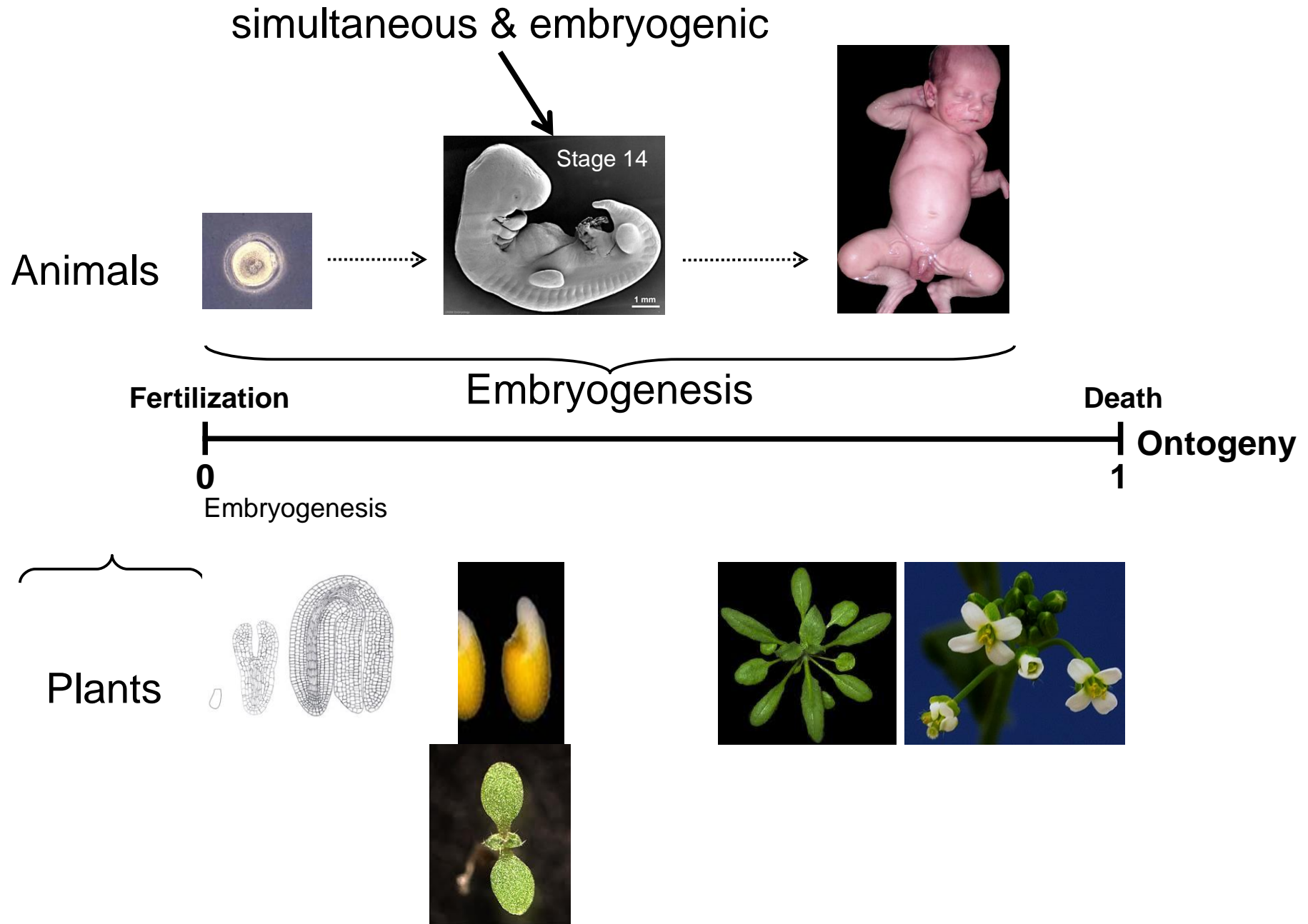
Ontogeny / organ development: animals vs. plants



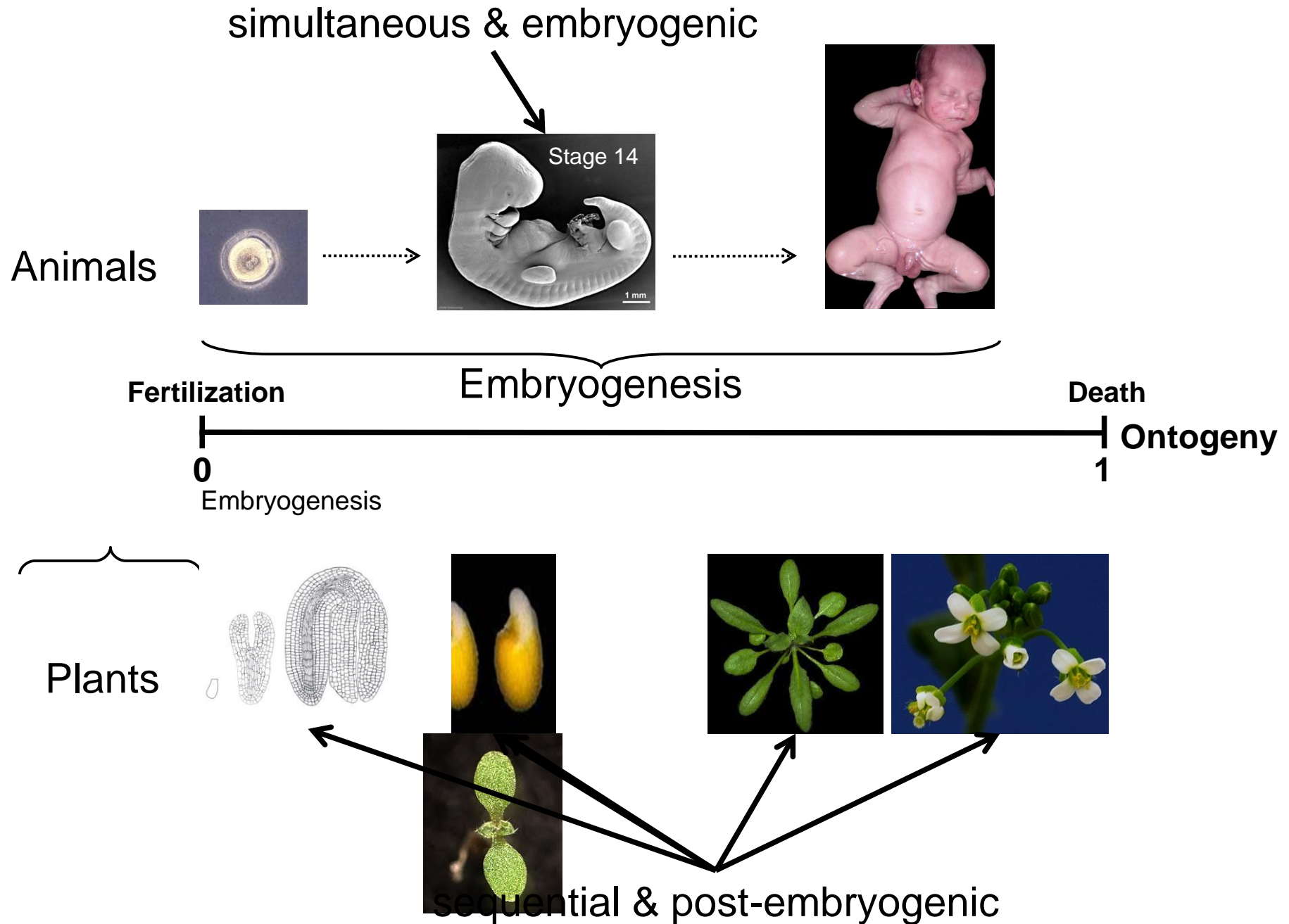
Ontogeny / organ development: animals vs. plants



Ontogeny / organ development: animals vs. plants



Ontogeny / organ development: animals vs. plants



Are transcriptomic hourglasses unique to embryogenesis?

Are they possibly a general feature of developmental transitions?

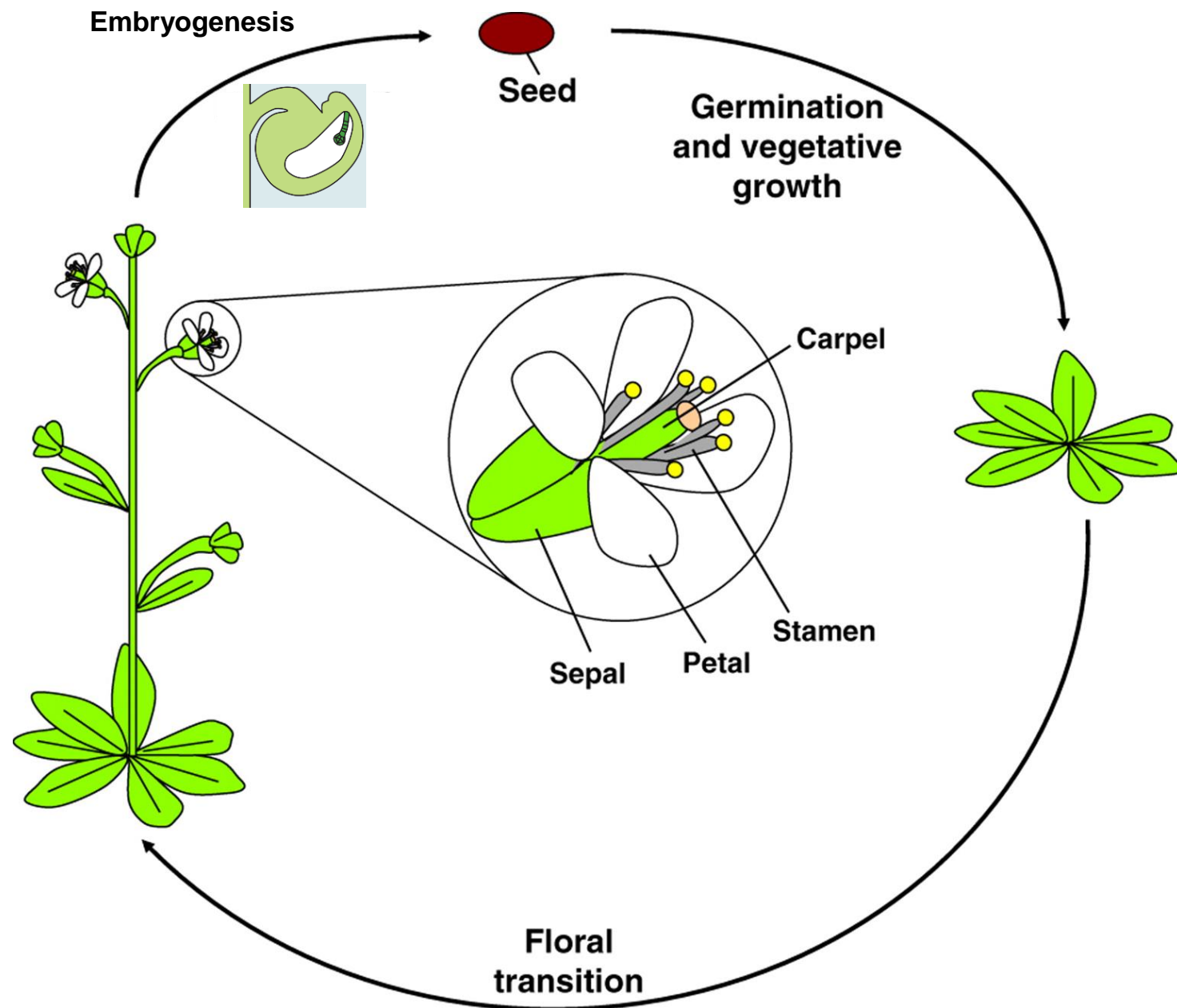


Difficult to address in most animals → development largely embryonic

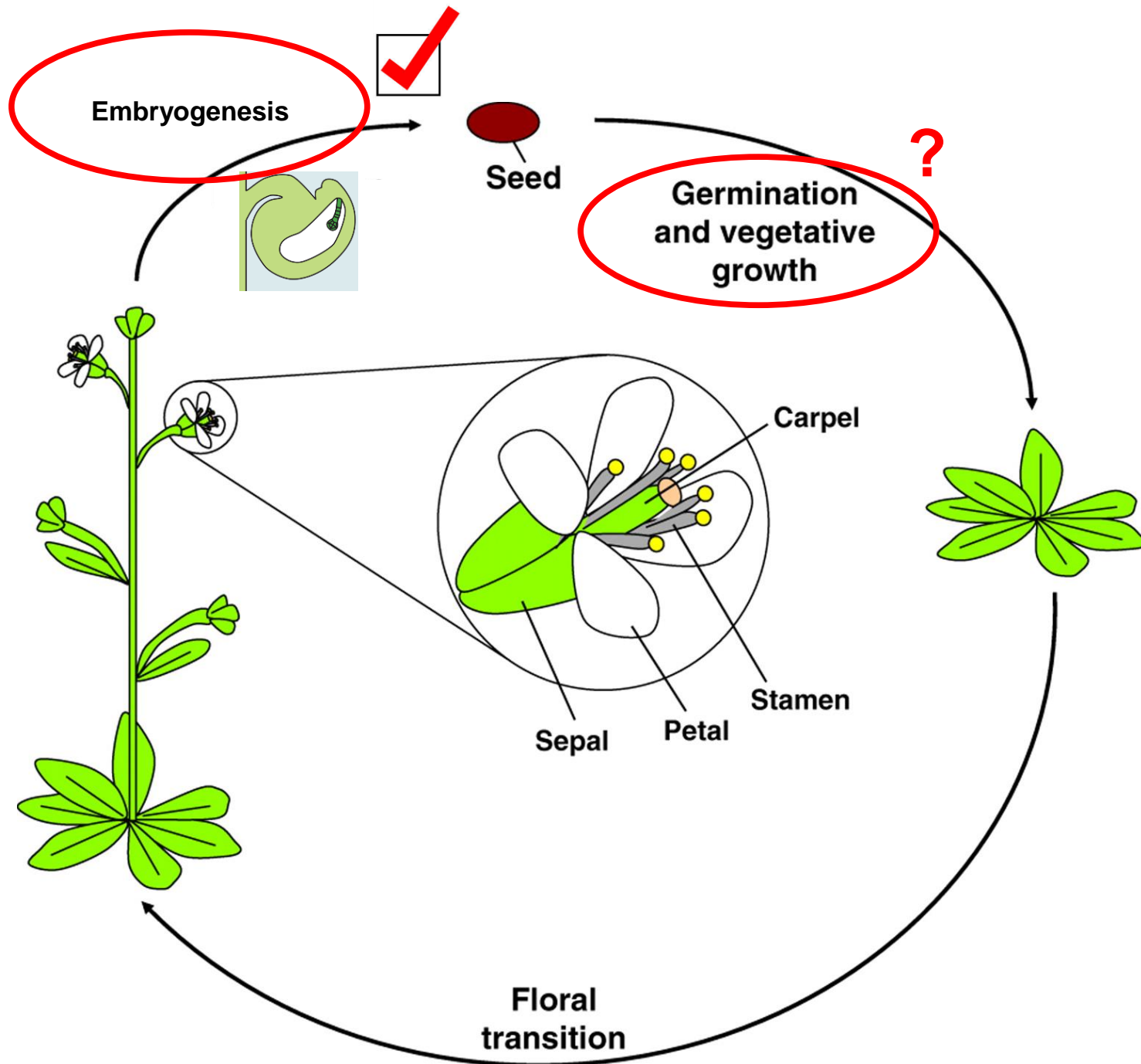


Straight-forward to address in plants
→ development largely post-embryonic

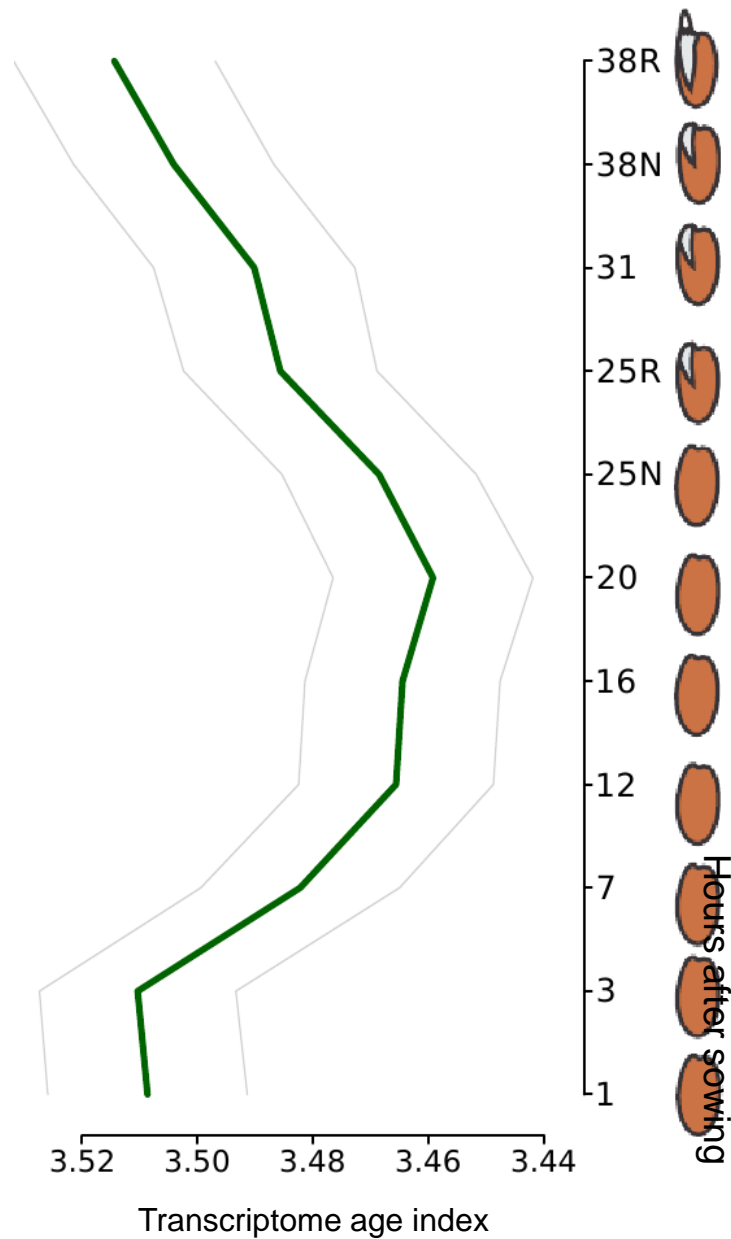
Major developmental transitions in the *A. thaliana* life cycle



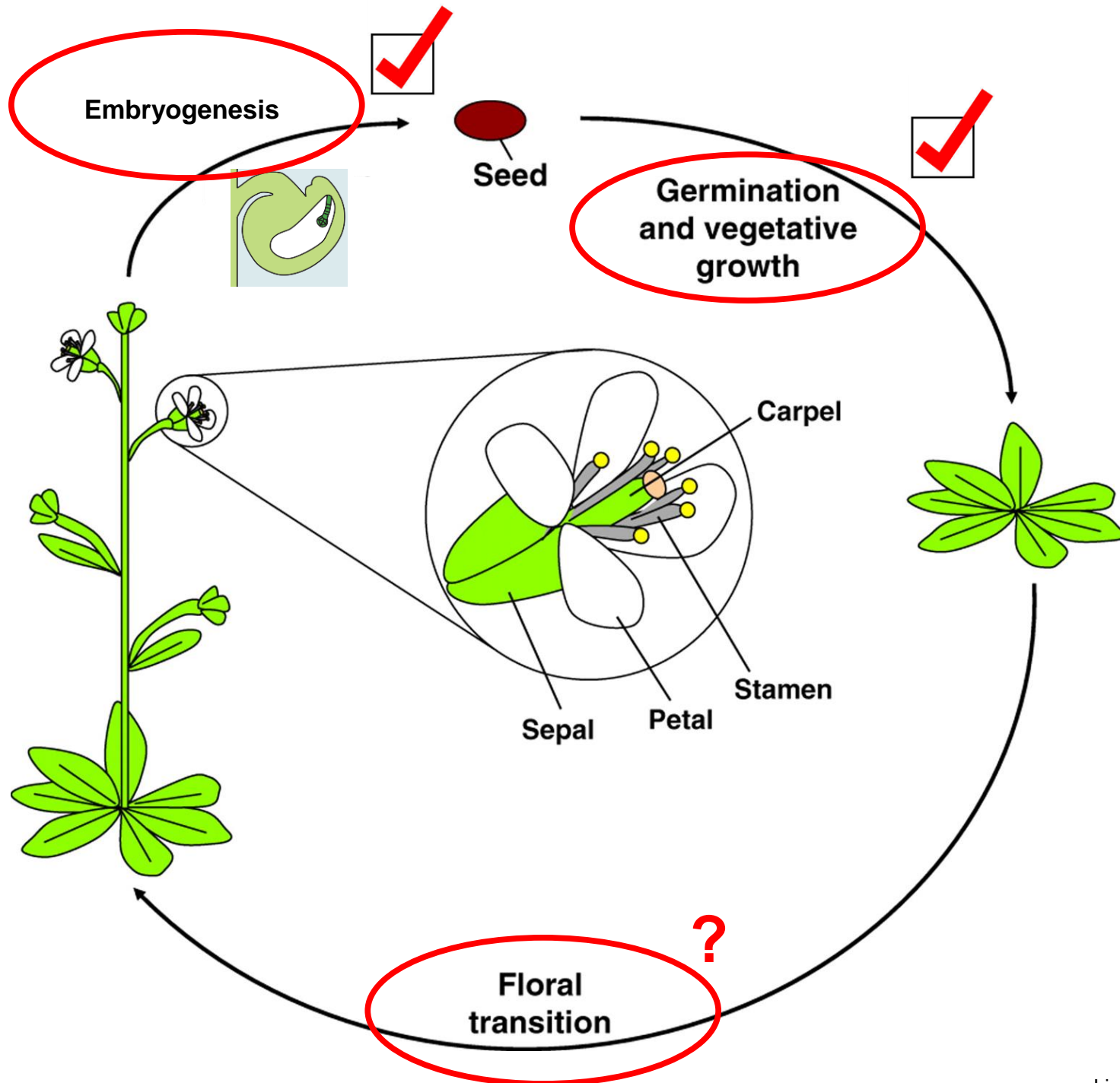
Major developmental transitions in the *A. thaliana* life cycle



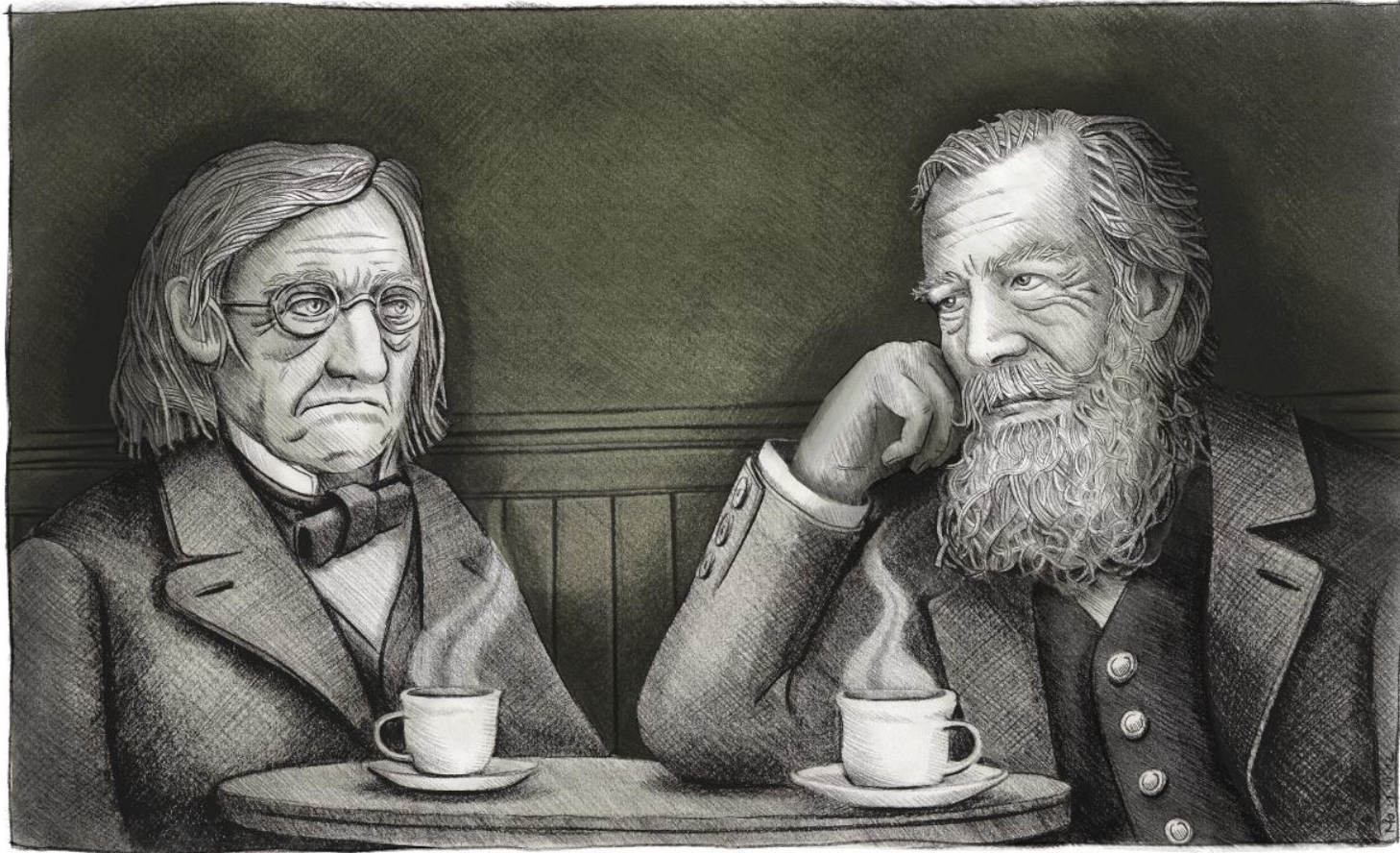
A developmental hourglass for germination



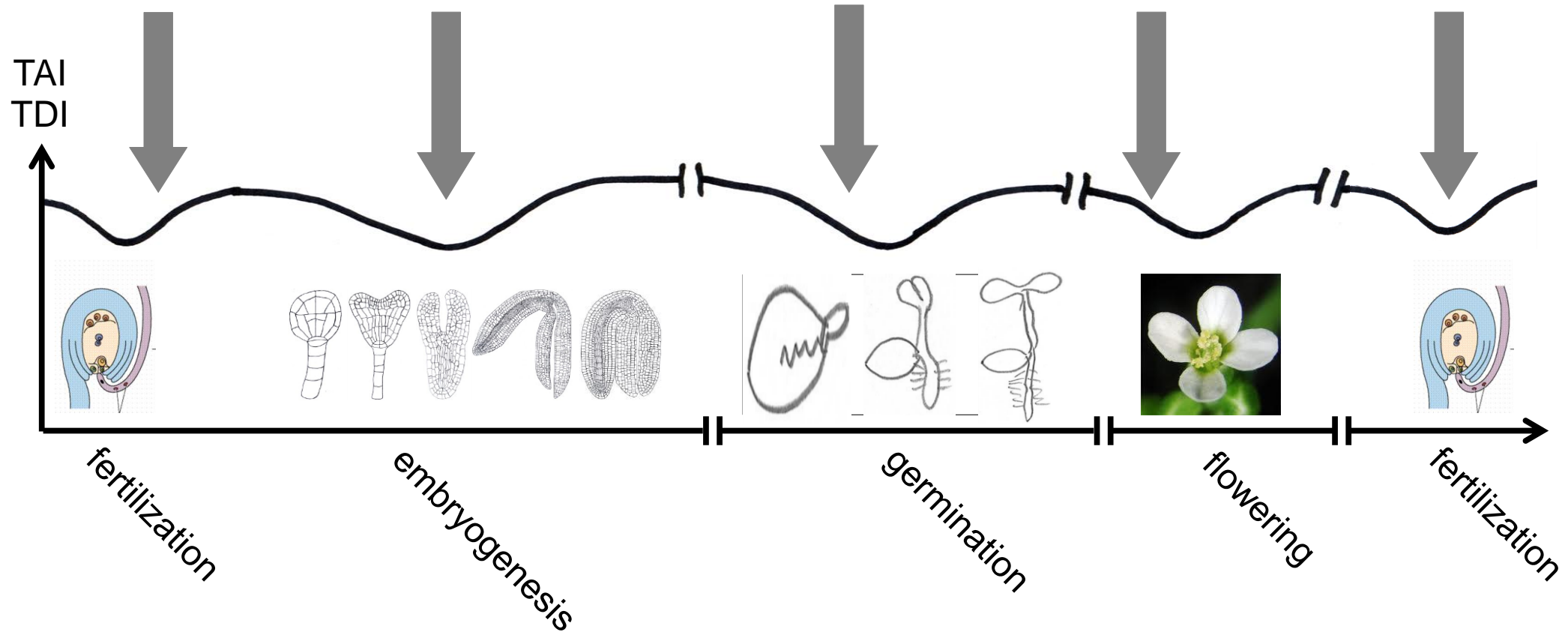
Major developmental transitions in the *A. thaliana* life cycle



Many hourglasses?

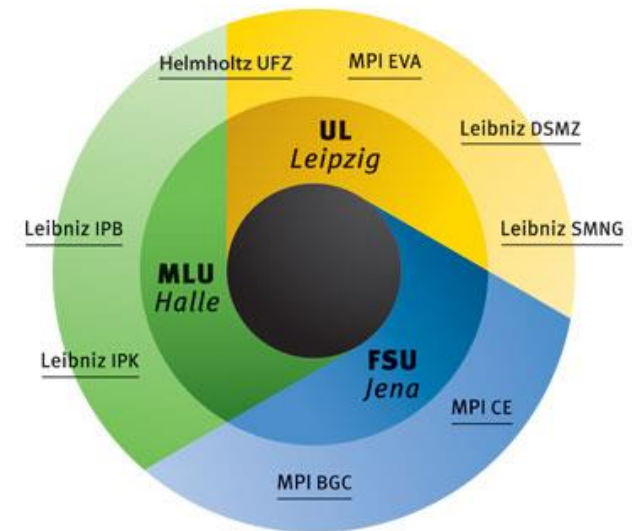
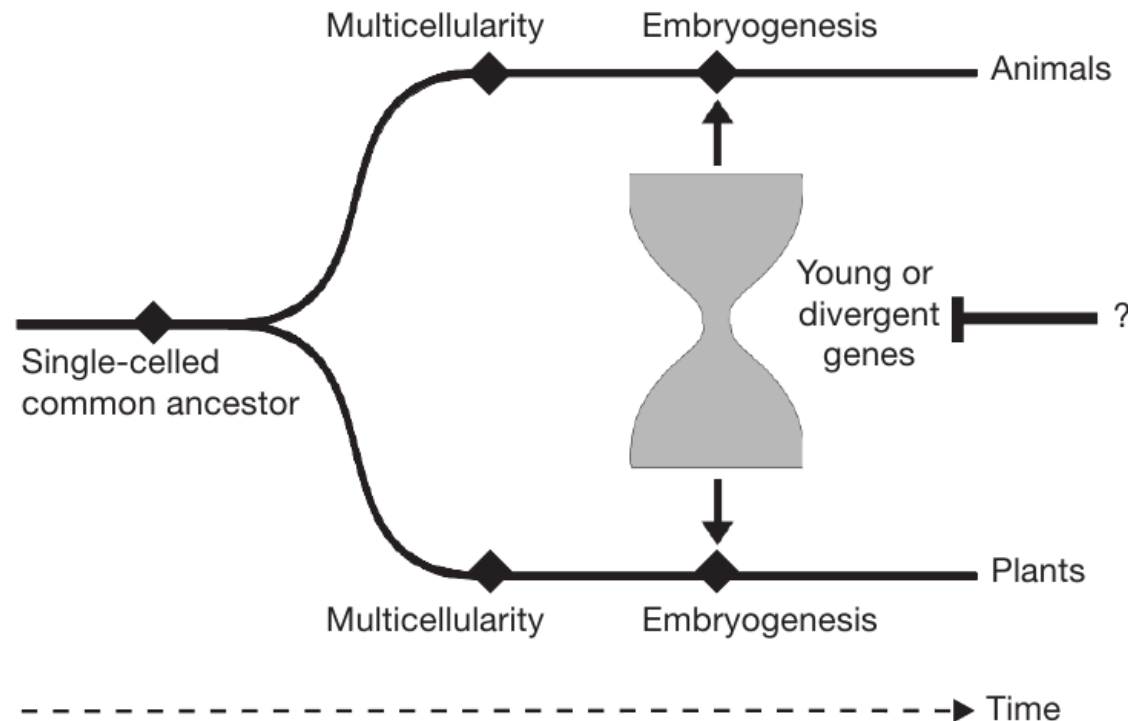


Many hourglasses? Checkpoint hypothesis



Conclusions

- Nature invented embryogenesis twice and independently
- Embryogenesis very different in plants and animals
- Developmental hourglass seems to be
- essential
- actively maintained
- possibly still functional?
- Postembryonic hourglasses in plants in germination and floral transition
- Existence of a universal logic that coordinates (i) embryogenesis in both kingdoms



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