

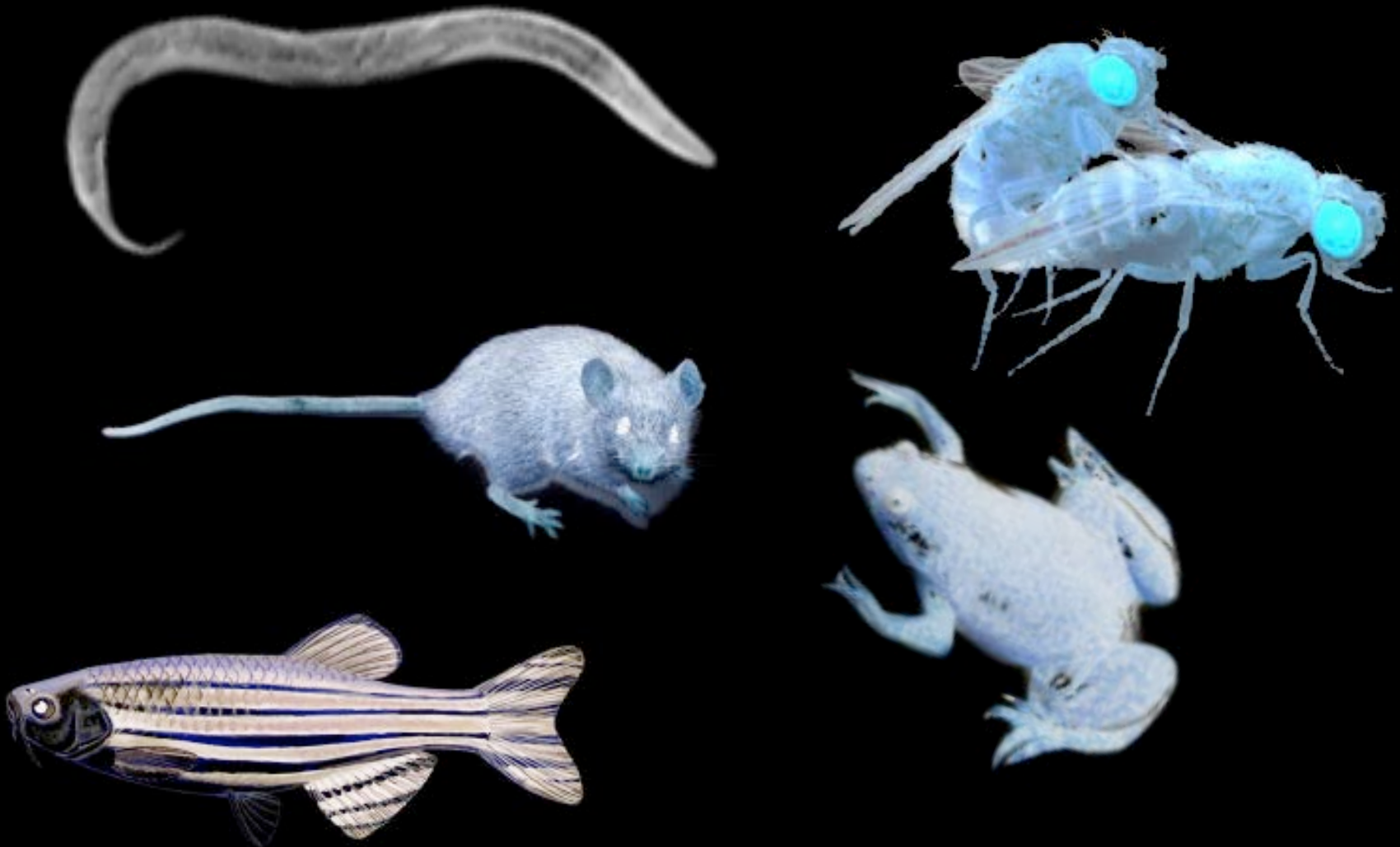
**Alternative model organisms:**  
New tools, new insights  
into the evolution of development



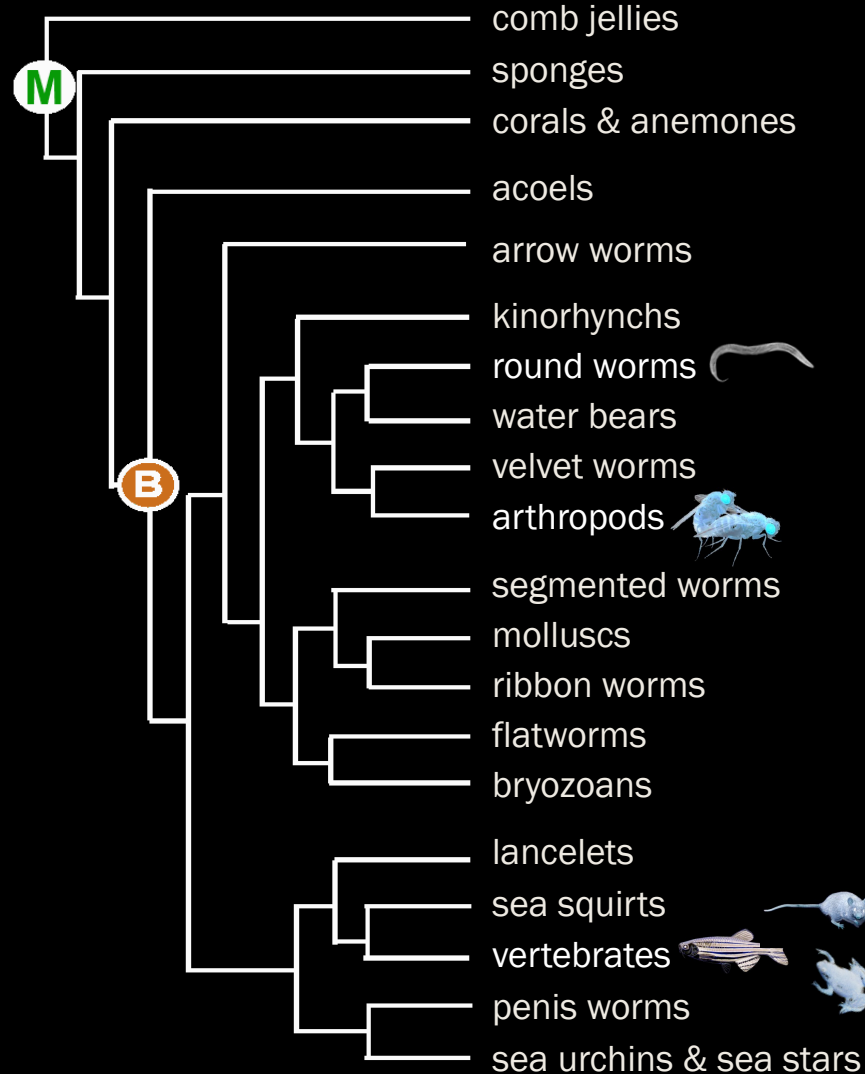
**Cassandra Extavour**  
Harvard University

*Department of Molecular and Cellular Biology*  
*Department of Organismic and Evolutionary Biology*

**Model organisms have provided invaluable case studies of developmental processes but...**

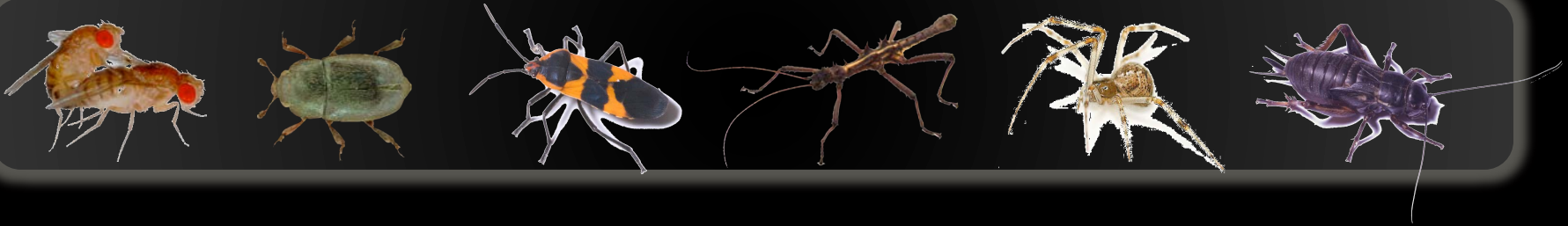


# ...traditional model organisms represent very little of animal diversity





1. How do we choose organisms for study?
2. Case study: a basally branching model offers insight into the evolution of germ line specification.
3. Tool building: quantitative analysis of putative “ancestral” insect development.



# 1. How do we choose organisms for study?

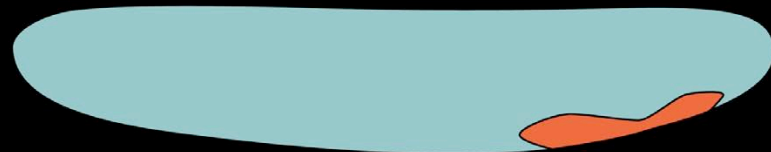
2. Case study: a basally branching model offers insight into the evolution of germ line specification.
3. Tool building: quantitative analysis of putative “ancestral” insect development.

# Traditional model organisms are derived relative to basally branching members of their clades



most body plan elements  
established maternally  
(earlier in development)

**Lab models tend to be selected because of their:**  
*fast development*  
*rapid sexual maturation*  
*large population sizes*  
*high tolerance for environmental variability*



most body plan elements  
established zygotically  
(later in development)

## *Drosophila melanogaster*



- Easy culture
- Community
- Complete genome
- Embryos all year
- 24 hour embryogenesis
- 10 day life cycle
- Powerful genetics

## *Gryllus bimaculatus*



- Easy culture
- Transcriptome
- Draft genome
- Embryos all year
- 10 day embryogenesis
- 6 week life cycle
- RNA interference, CRISPR/Cas9



*cricket embryo dissection*

## *Drosophila melanogaster*

### DERIVED FEATURES

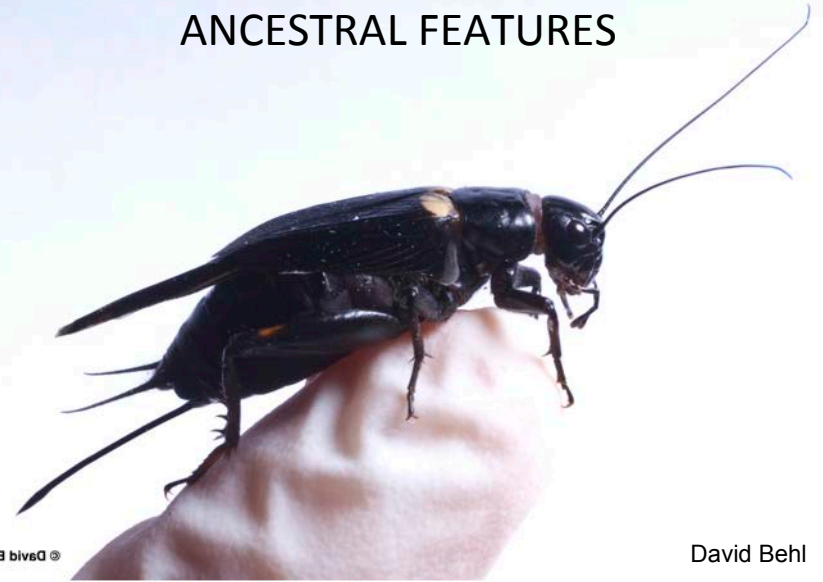


Kuntzei lab

- Holometabola (metamorphosis)
- Long germ development
- Simultaneous segmentation
- Meroistic ovaries (nurse cells)
- Reduced extra embryonic membrane
- Large embryonic rudiment
- No blastokinesis

## *Gryllus bimaculatus*

### ANCESTRAL FEATURES




InfoB bivaG ©

David Behl

- Hemimetabola (direct development)
- Short germ development
- Sequential segmentation
- Panoistic ovaries (no nurse cells)
- 2 Large extra embryonic membranes
- Small embryonic rudiment
- Blastokinesis



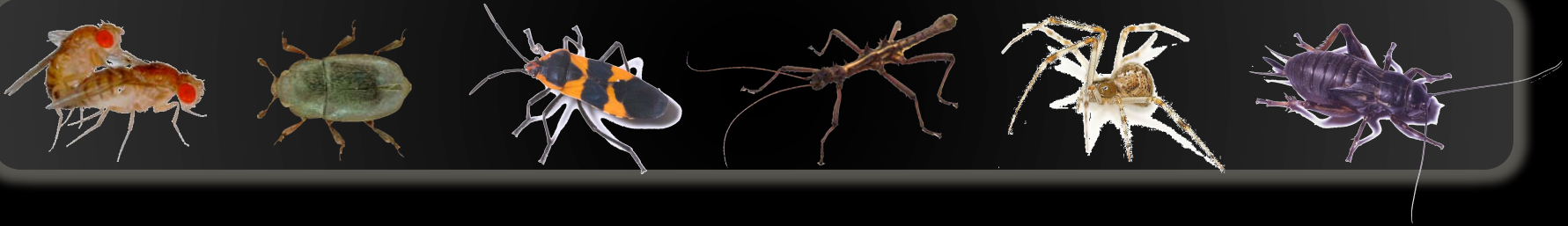
# Embryonic development in *Gryllus bimaculatus*



white light

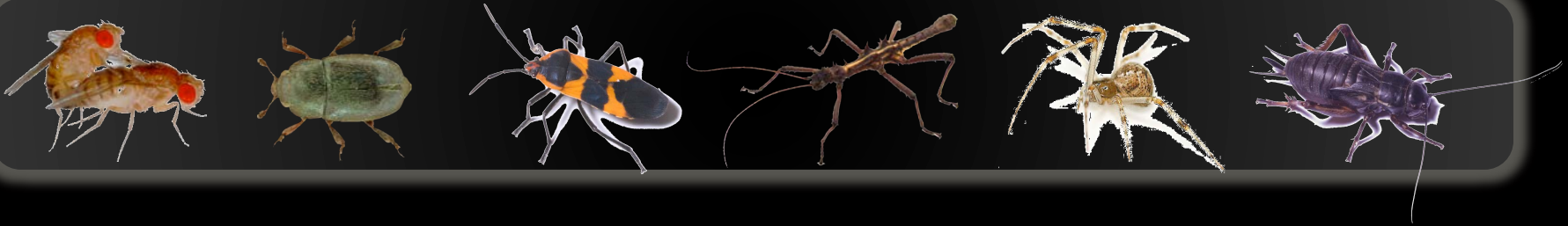


Histone2B-eGFP



## 1. How do we choose organisms for study?

- The most convenient laboratory organisms may provide limited evolutionary inference.
- Phylogenetic position and experimental tools should correspond to the questions being asked.
- Basally branching insect and arthropod models can help place findings from *Drosophila melanogaster* in an appropriate evolutionary context.
- *Gryllus bimaculatus* is a promising model to understand the evolution of arthropod development.

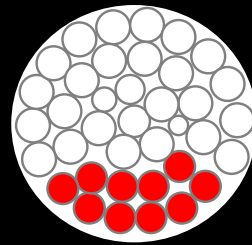
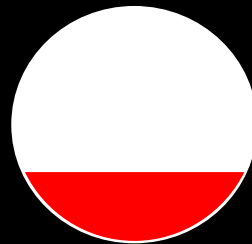
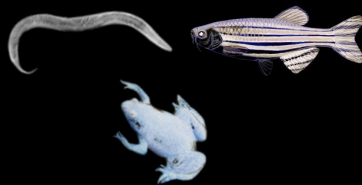


1. How do we choose organisms for study?
2. Case study: a basally branching model offers insight into the evolution of germ line specification.
3. Tool building: quantitative analysis of putative “ancestral” insect development.

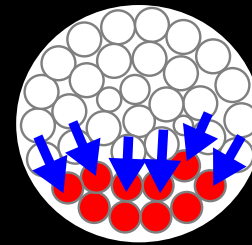
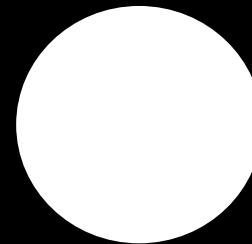
# Animals have two main ways of making primordial germ cells (PGCs)



**Inheritance:**  
cytoplasmic  
determinants  
(germ plasm)



**Inheritance**



**Induction**



**Induction:**  
cell-cell  
signals  
may be  
ancestral in  
animal PGC  
formation.



germ cells



somatic cells

Zygotic mechanisms rather than maternal determinants  
make germ cells in basally branching arthropods



*Oncopeltus fasciatus*



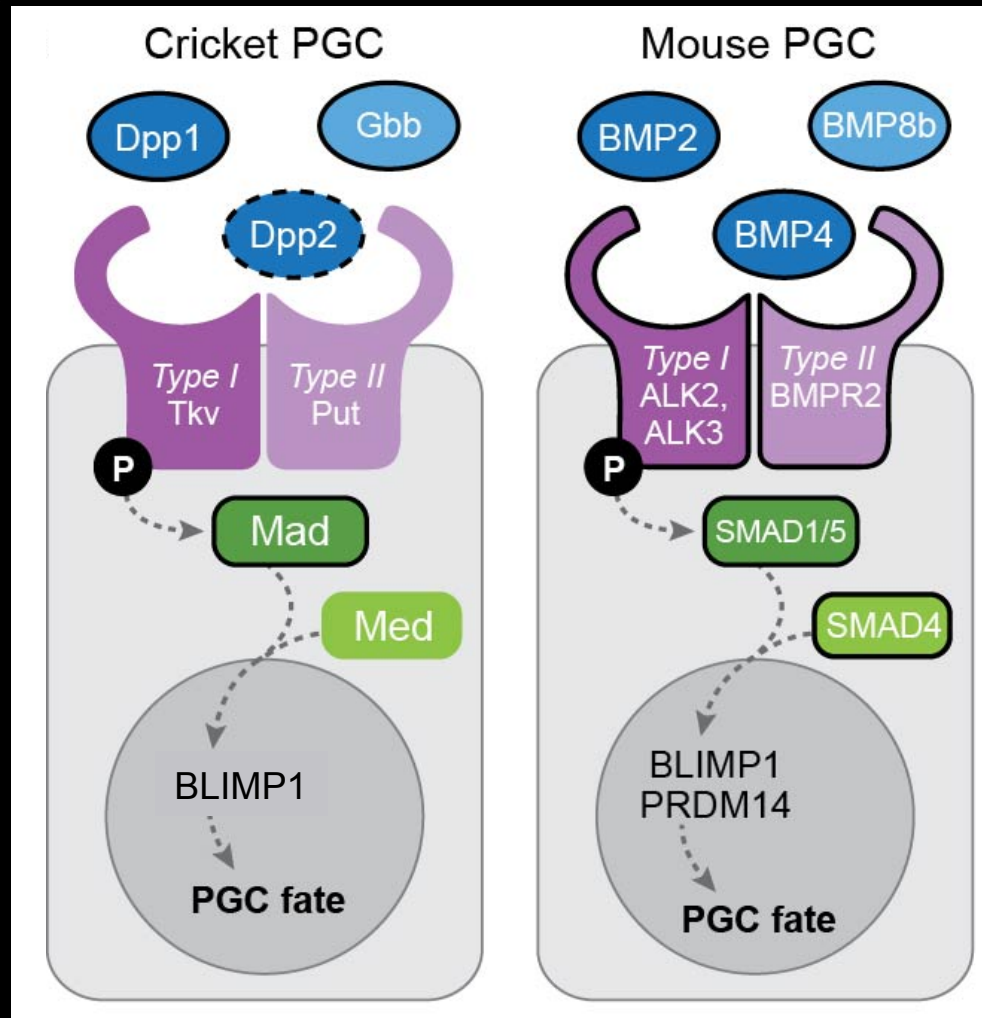
*Gryllus bimaculatus*



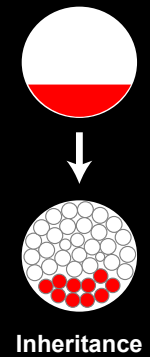
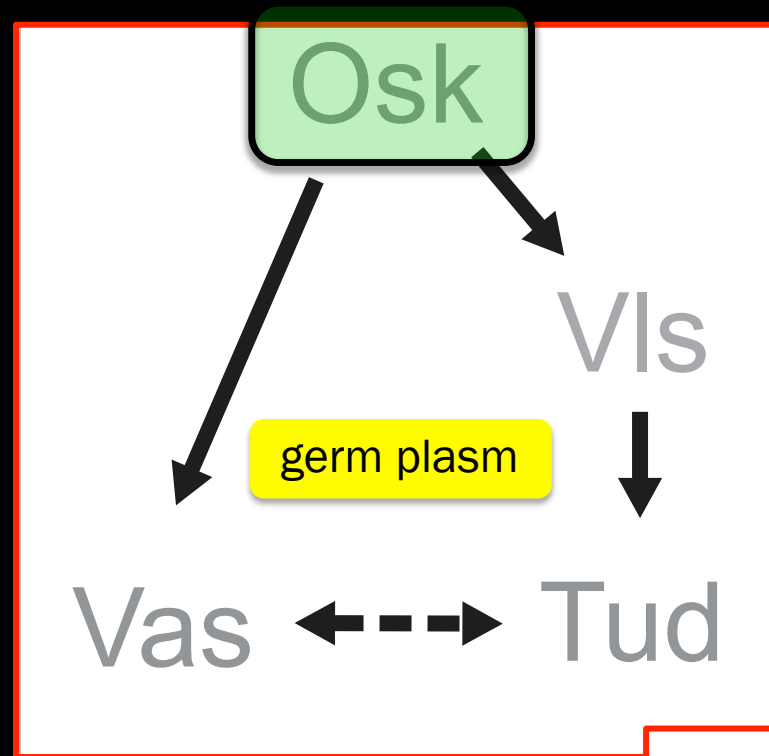
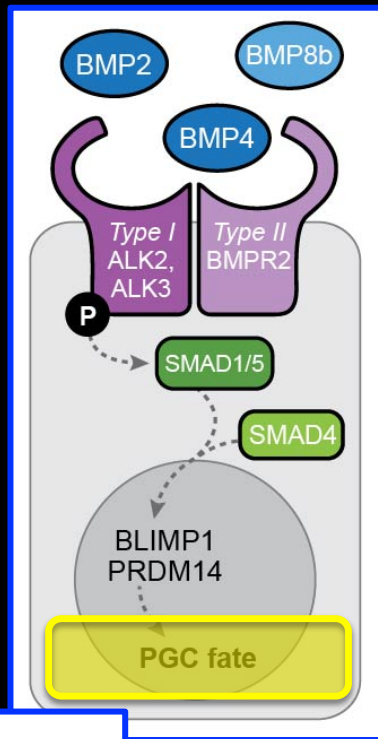
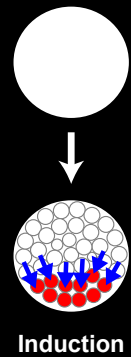
*Parasteatoda tepidariorum*

Ewen-Campen et al. *BMC Genomics* 2011  
Zeng et al. *BMC Genomics* 2011  
Zeng & Extavour *Database* 2012  
Zeng et al. *PLoS ONE* 2013  
Ewen-Campen et al. *Current Biology* 2013  
Ewen-Campen et al. *BiO* 2013  
Posnien et al. *PLoS ONE* 2014  
Schwager et al. *Developmental Biology* 2015

# Mouse and cricket PGCs are generated by BMP signalling upstream of Blimp-1



# How did germ plasm assembly evolve?



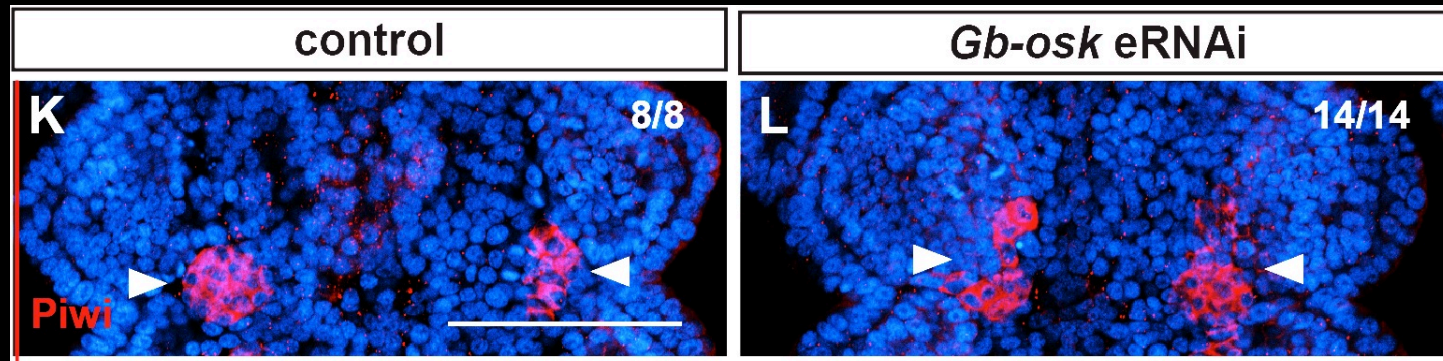
*Mus, Gryllus:*  
**BMP**  
Induction

There is a conserved molecular signature of germ cell fate, independent of the upstream mechanisms

*Drosophila:*  
**Oskar**  
inheritance

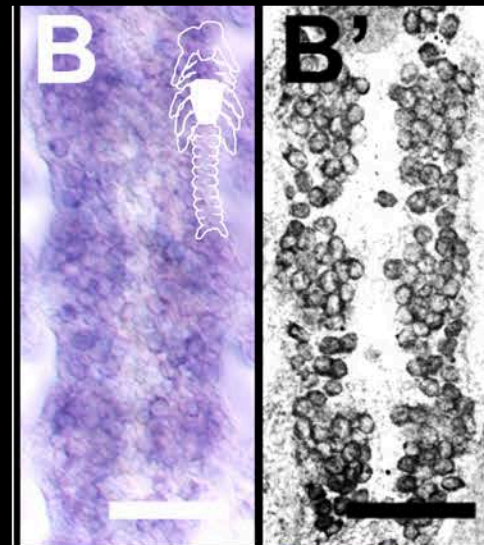
# *Gryllus oskar* does not play “canonical” *oskar* roles

DNA  
Germ  
cells



## *Drosophila oskar*:

- Asymmetric localization in oocyte or early embryo
- PGC formation
- Oogenesis
- Posterior determinant localization or patterning



*Gb-osk*  
**transcript** and  
**protein** are  
expressed in  
cricket  
neuroblasts.

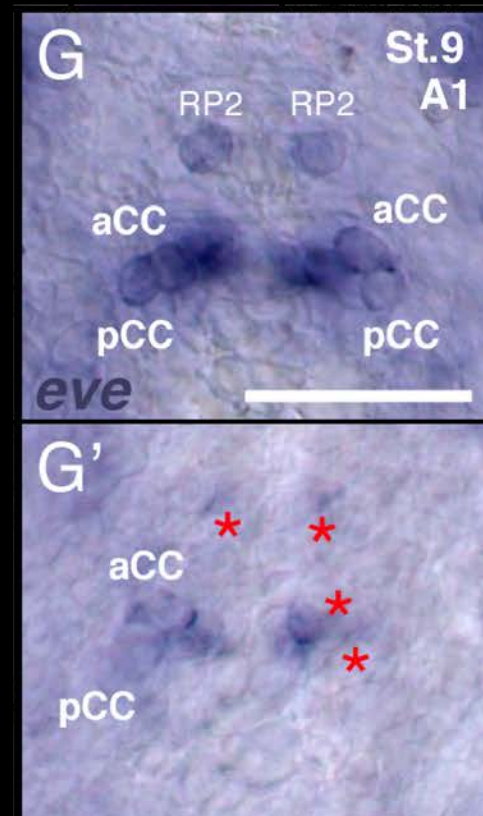
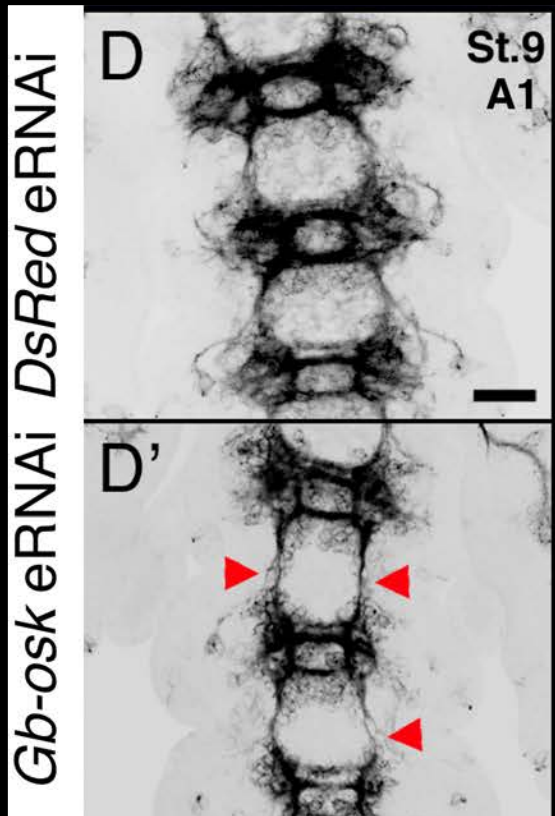


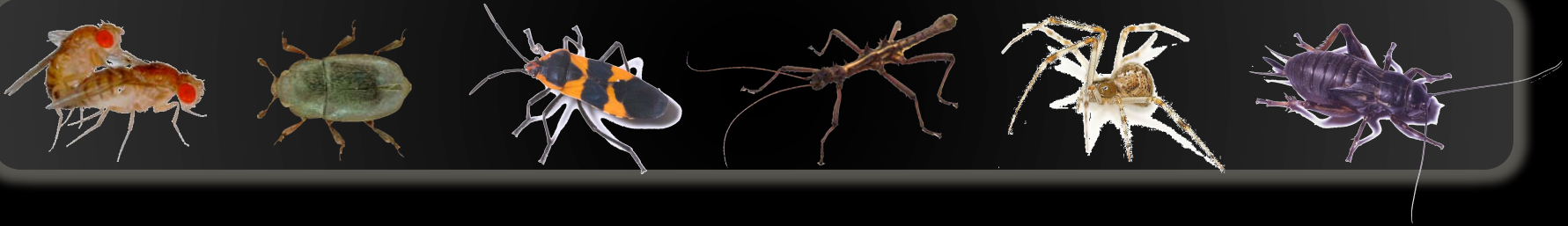
# *Gryllus oskar* knockdown causes defects in embryonic axonal morphogenesis and neuronal patterning

Longitudinal connective defects

Anterior & posterior commissure defects

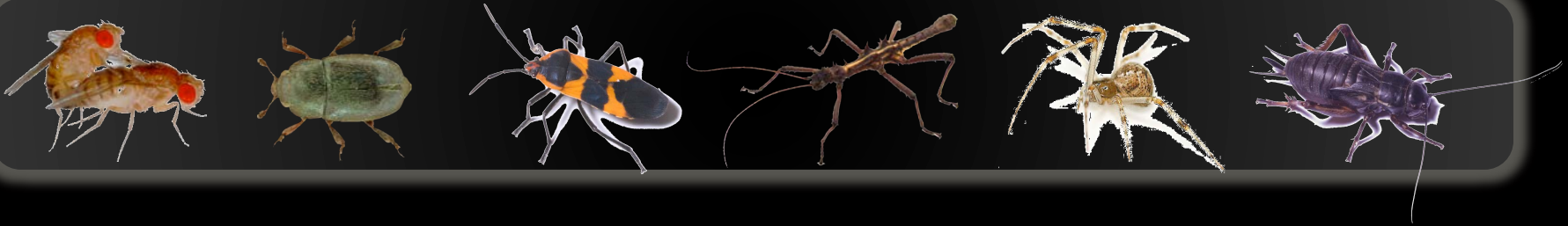
Missing RP2, aCC or pCC neurons





## 2. Case study: a basally branching model offers insight into the evolution of germ line specification.

- Germ plasm is a derived mode of germ line specification in insects (and likely in animals).
- Cell-cell signaling (BMP/Blimp-1 based?) likely ancestral.
- *oskar* may have played an ancestral neural role in Hemimetabola, and then been co-opted to a germ line role in Holometabola, facilitating the evolution of germ plasm.

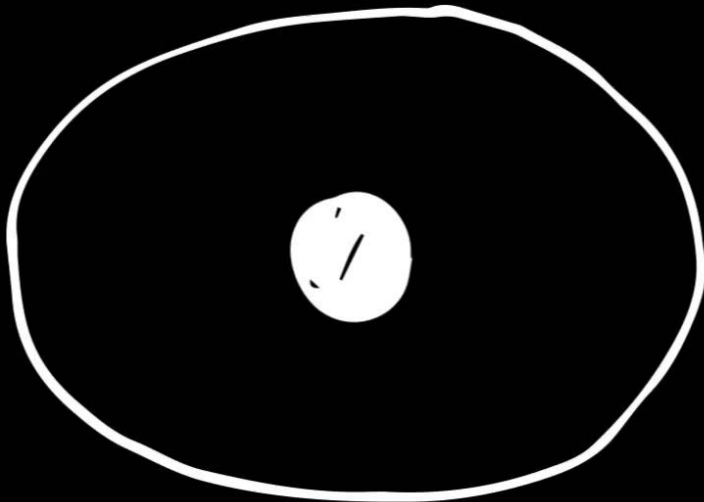


1. How do we choose organisms for study?
2. Case study: a basally branching model offers insight into the evolution of germ line specification.
3. Tool building: quantitative analysis of putative “ancestral” insect development.

# Animals have one of two types of embryonic cleavage

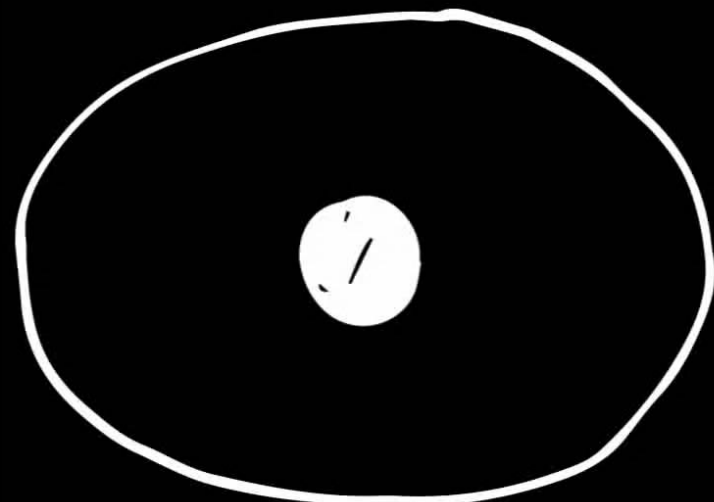
## Holoblastic

Nuclear division  
*with* cell cleavage

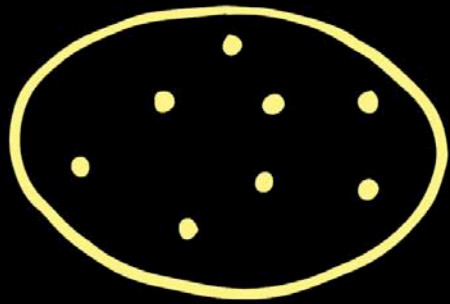
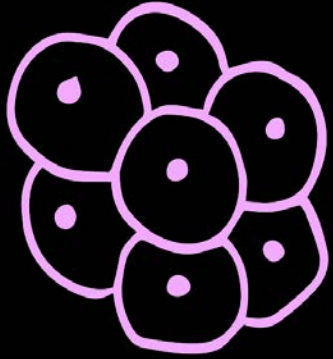


## Syncytial

Nuclear division  
*without* cleavage



Nuclear divisions with cell cleavages



Nuclear divisions without cell cleavages

Anemones

Vertebrates

Nematodes

Crustaceans



springtail



dragonfly



cricket



milkweed bug



beetle



moth

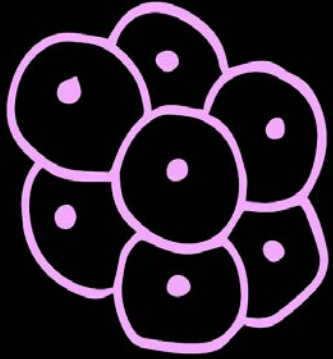


flea



fruit fly

Nuclear divisions with cell cleavages



Anemones

Vertebrates

Nematodes

Crustaceans



springtail



dragonfly



cricket



milkweed bug



beetle



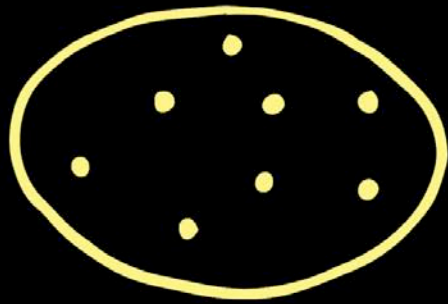
moth



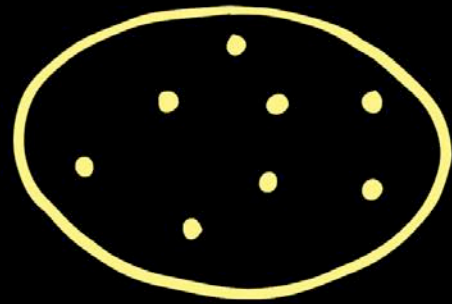
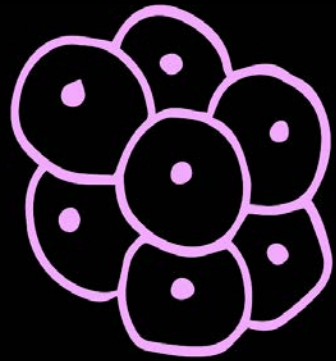
flea



fruit fly



Nuclear divisions without cell cleavages



Millions of years ago: 480 405

Anemones

Vertebrates

Nematodes

Crustaceans



springtail



dragonfly



cricket



milkweed bug



beetle



moth



flea



fruit fly

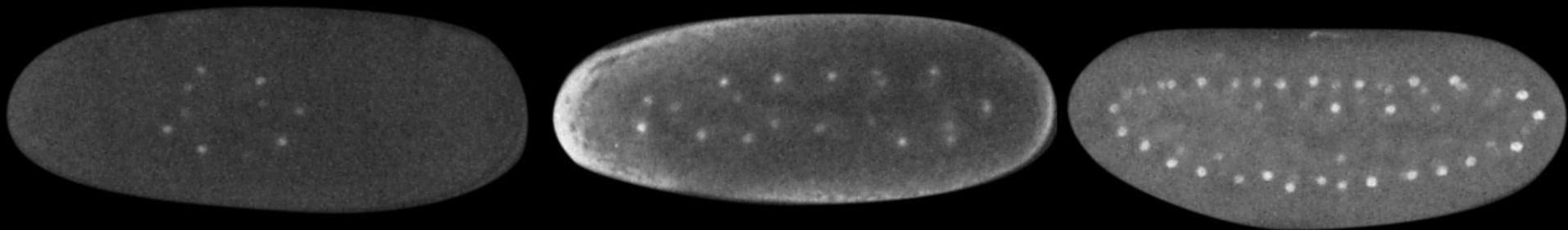
# Syncytial cleavage is not uniform across insects

*Gryllus bimaculatus*



Nakamura et al., 2010

*Drosophila melanogaster*

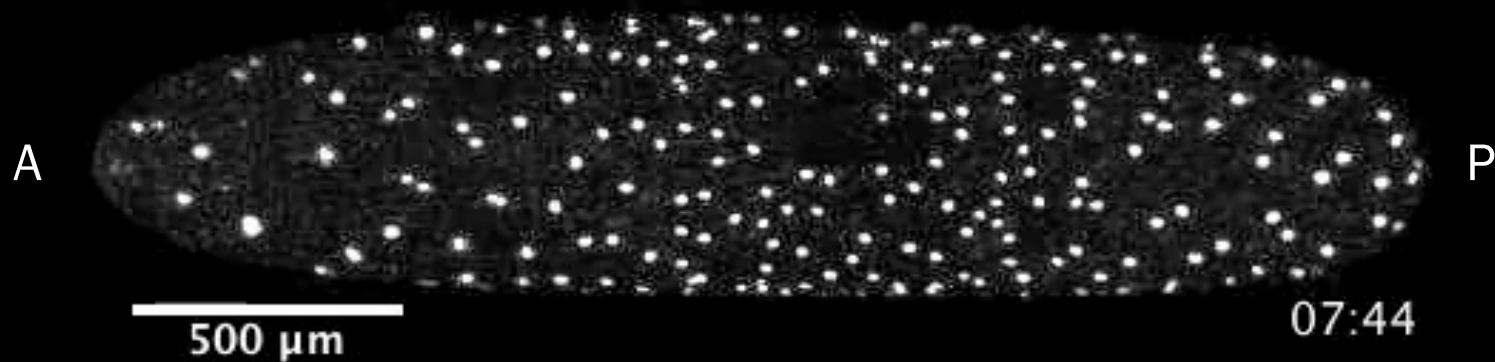


Tomer et al. 2012



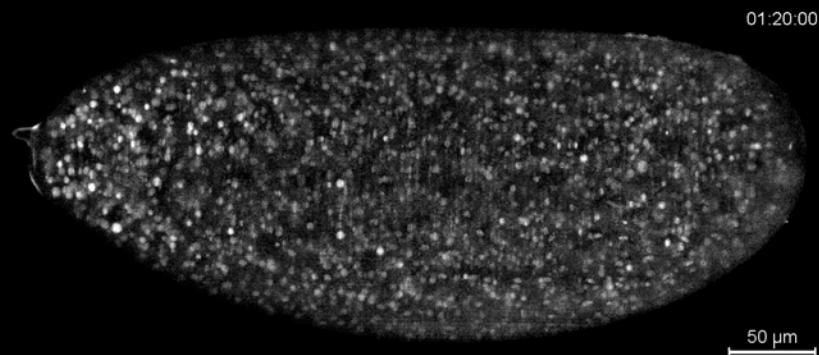
# Blastoderm stages, embryonic and extra-embryonic establishment are not uniform across insects

*Gryllus bimaculatus*

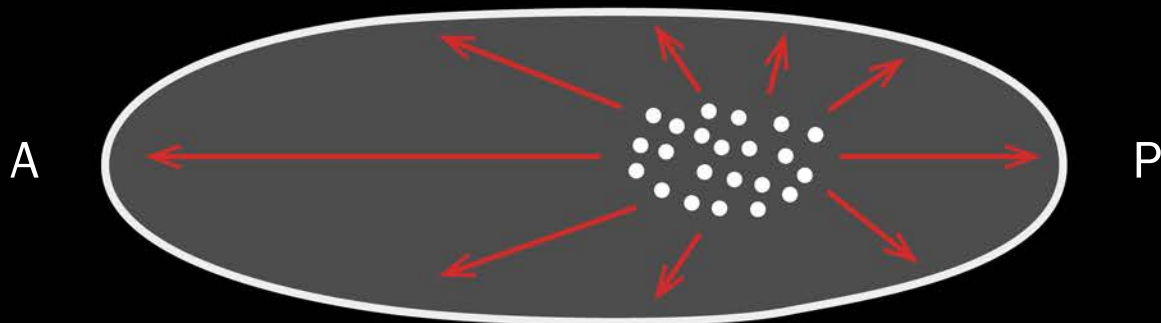


Nakamura et al., 2010

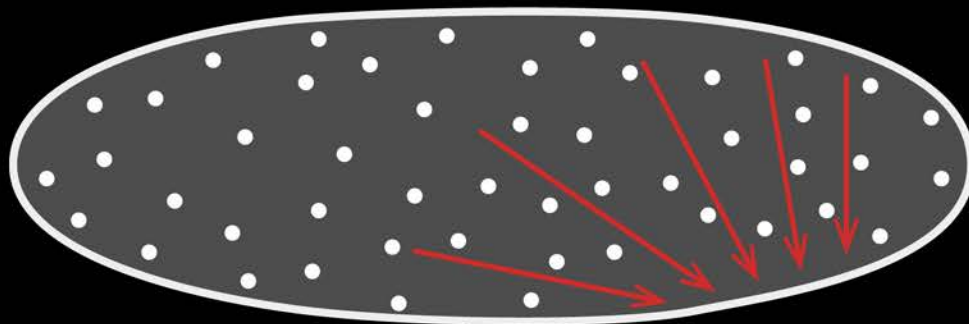
*Drosophila melanogaster*



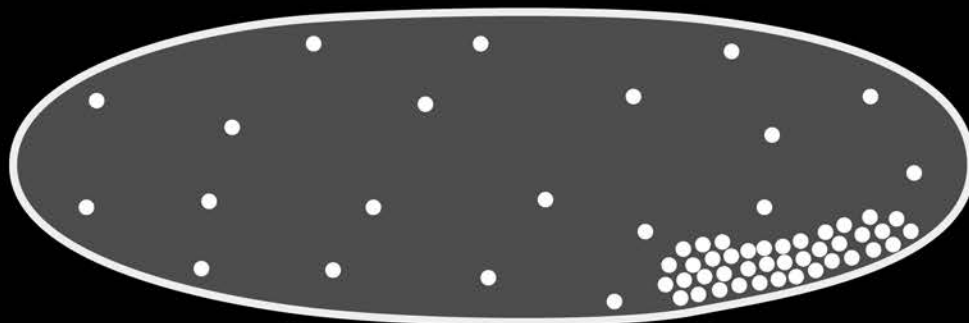
Tomer et al. 2012



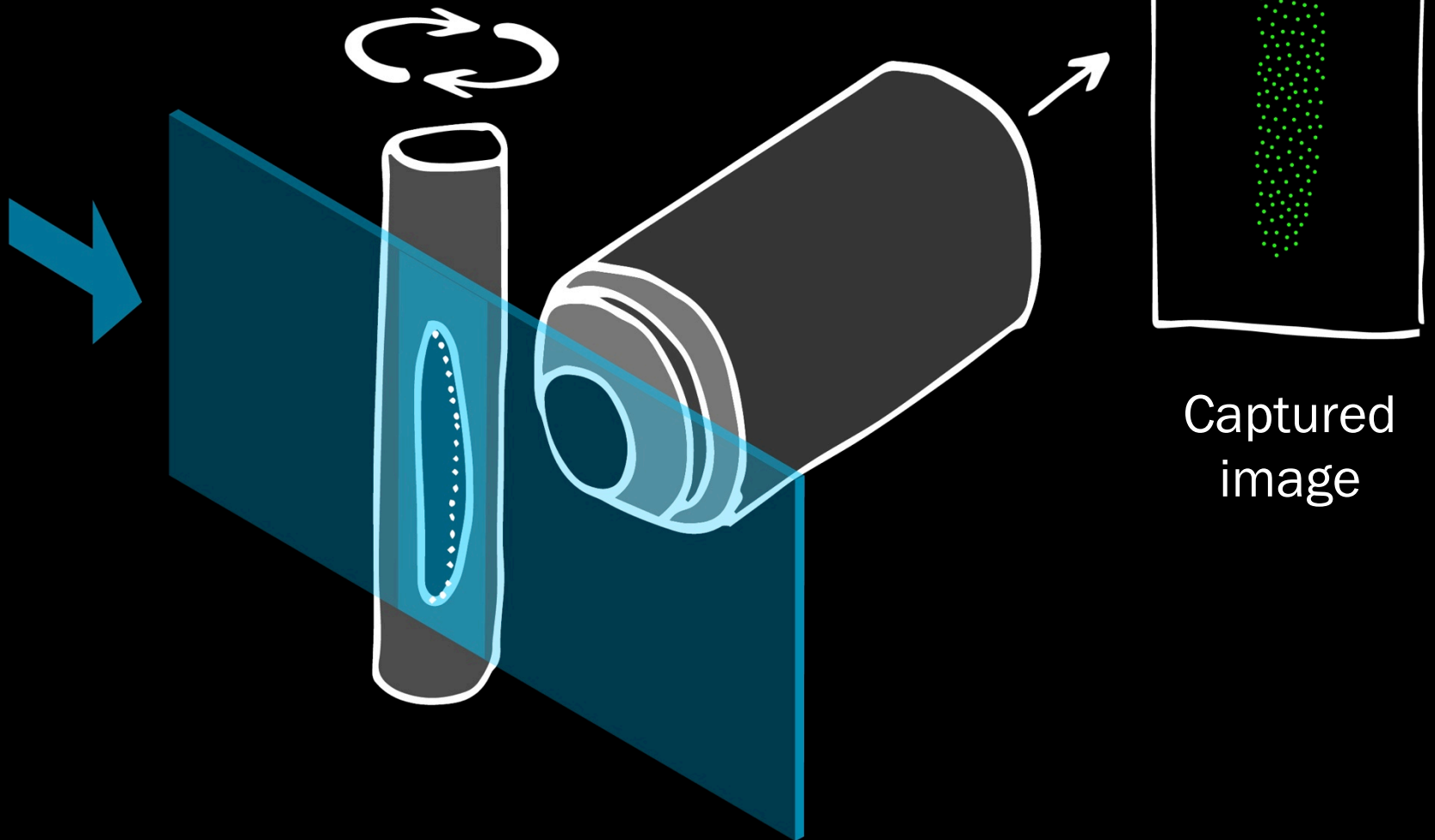
**axial expansion**



**embryonic coalescence**

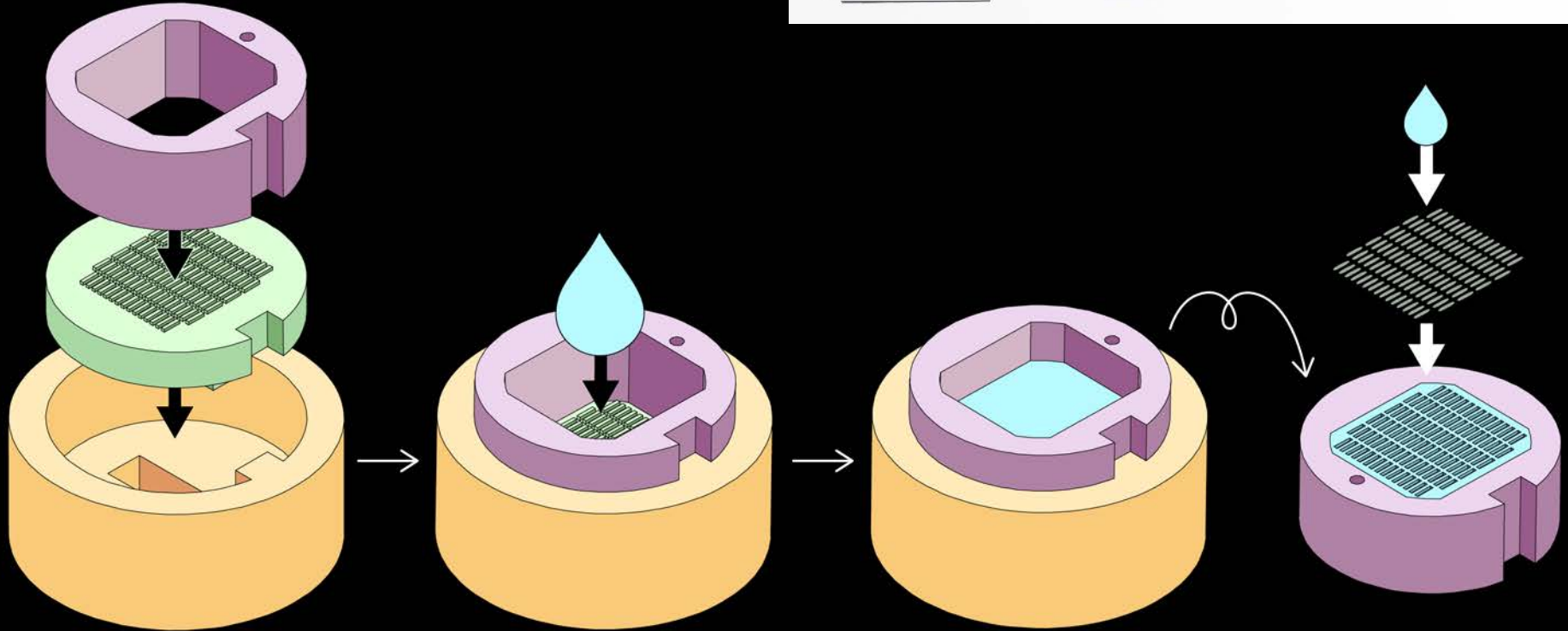


# Method 1: Lightsheet Z.1

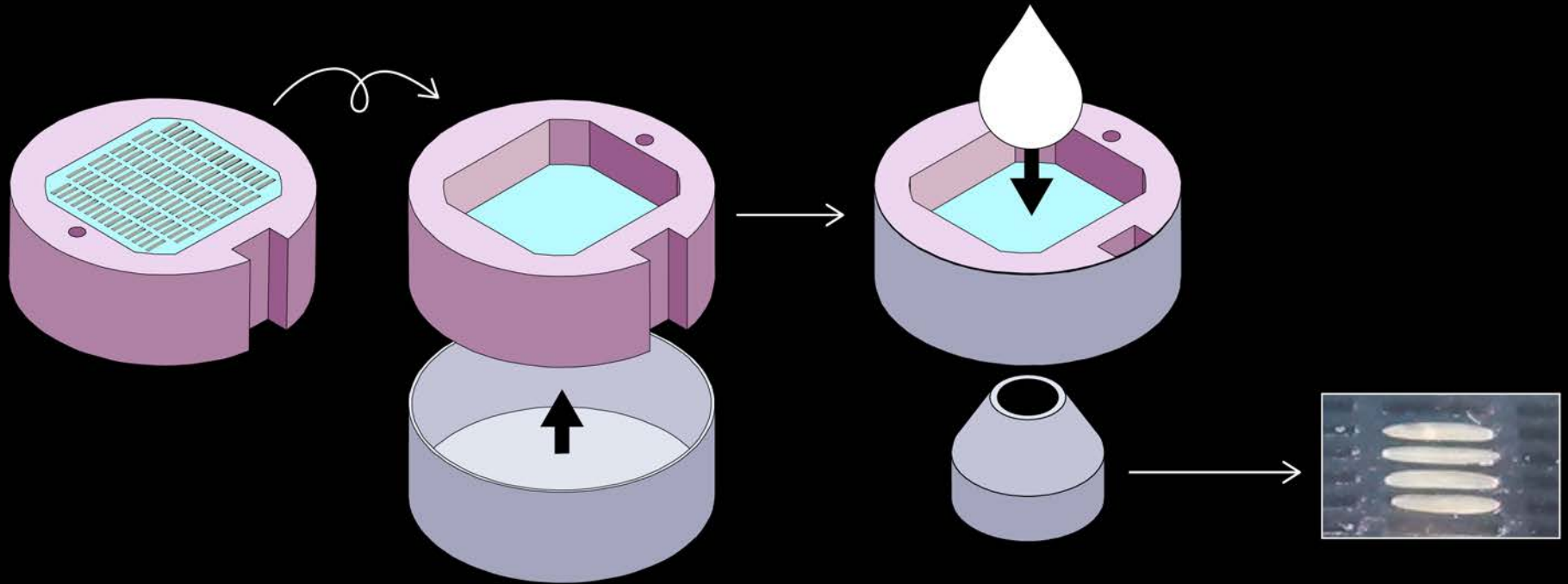


Captured  
image

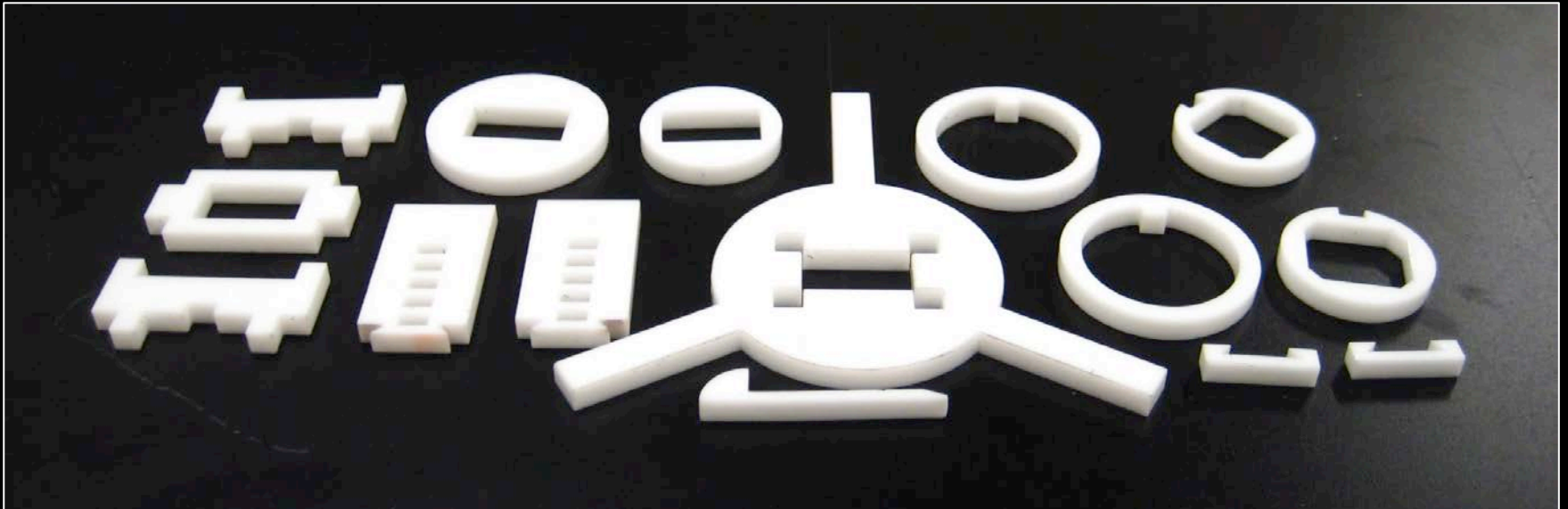
## Method 2: Cell Observer



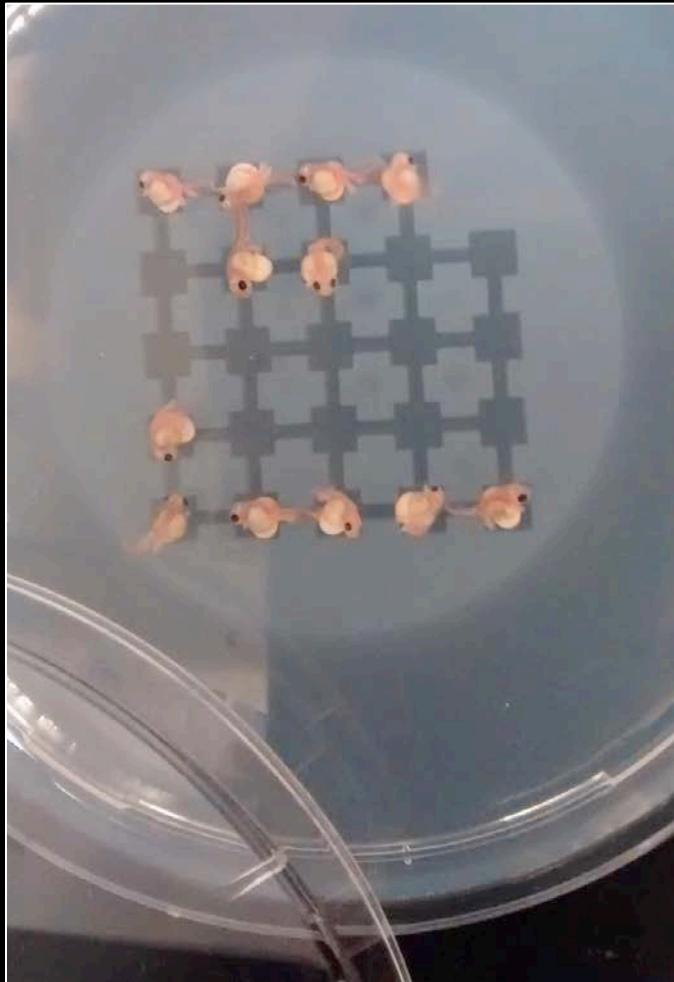
## Method 2: Cell Observer



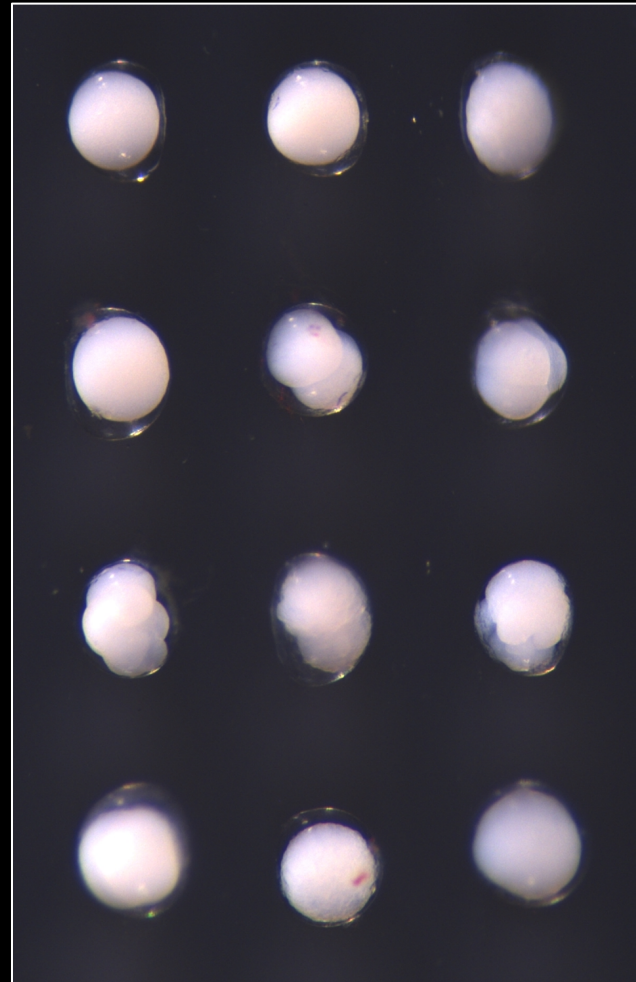
Modular device, easy to make with a laser cutter



## Easy to design inserts for different species



*Eleutherodactylus coqui* (frog)



*Hofstenia miamia* (acoel)

# Lightsheet Z.1



- 3 dimensional nucleus position information
- 1 Embryo at a time
- 4 angles, ~500 time points

A

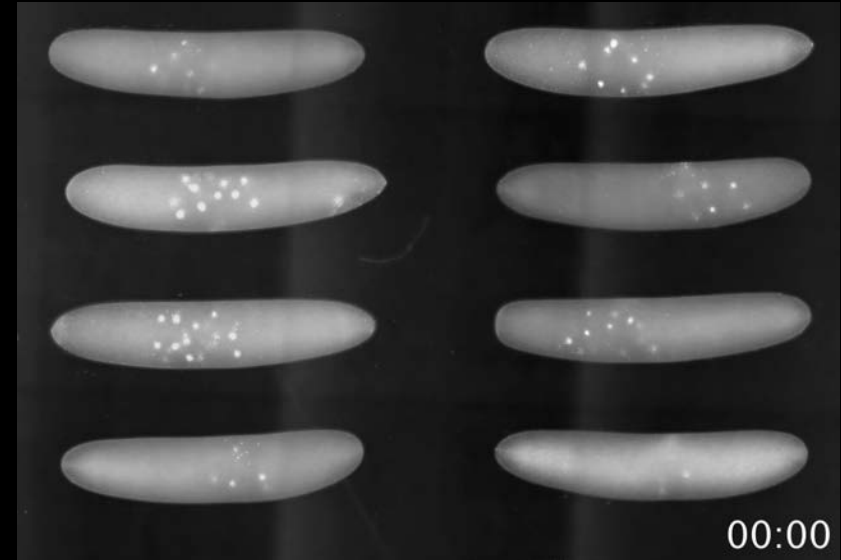


P

# Cell Observer



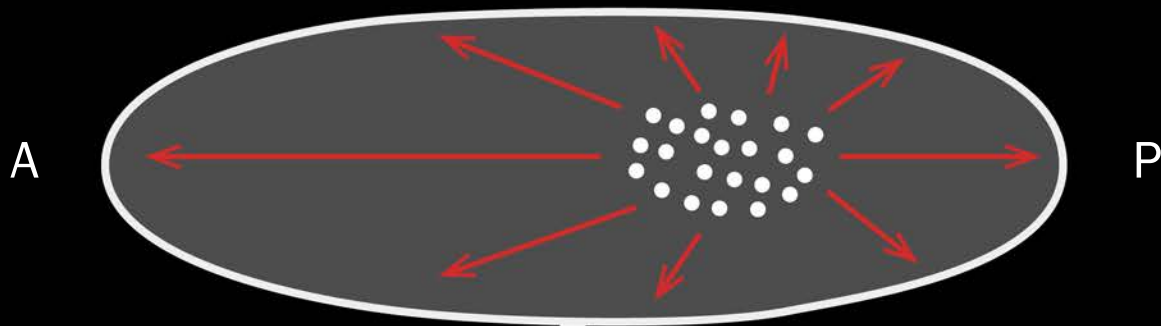
- 2 dimensional nucleus position information
- ~20 embryos at a time
- 1 angle, ~500 time points



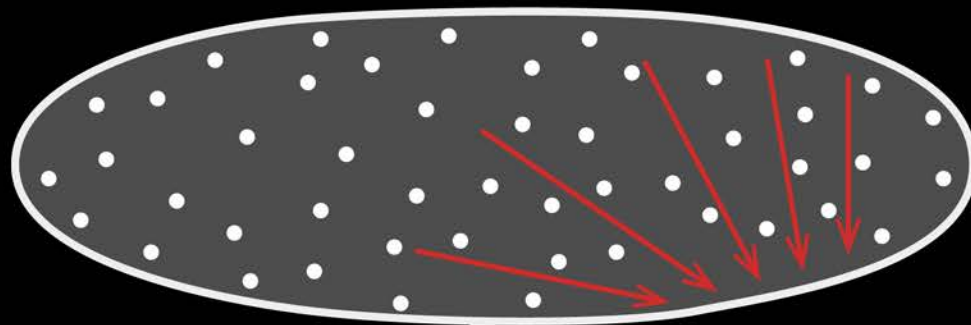
A

P

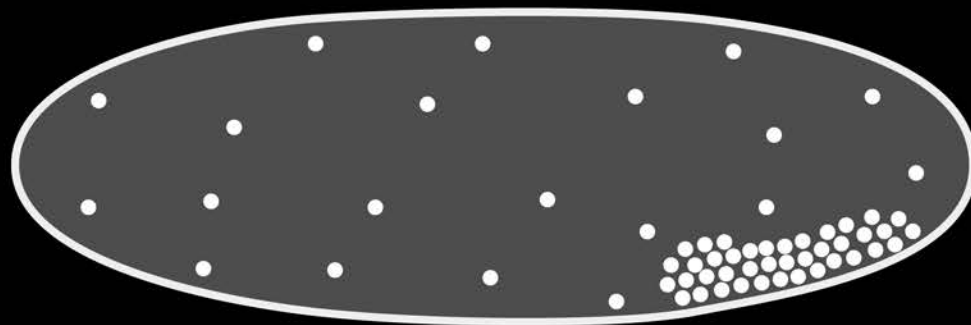


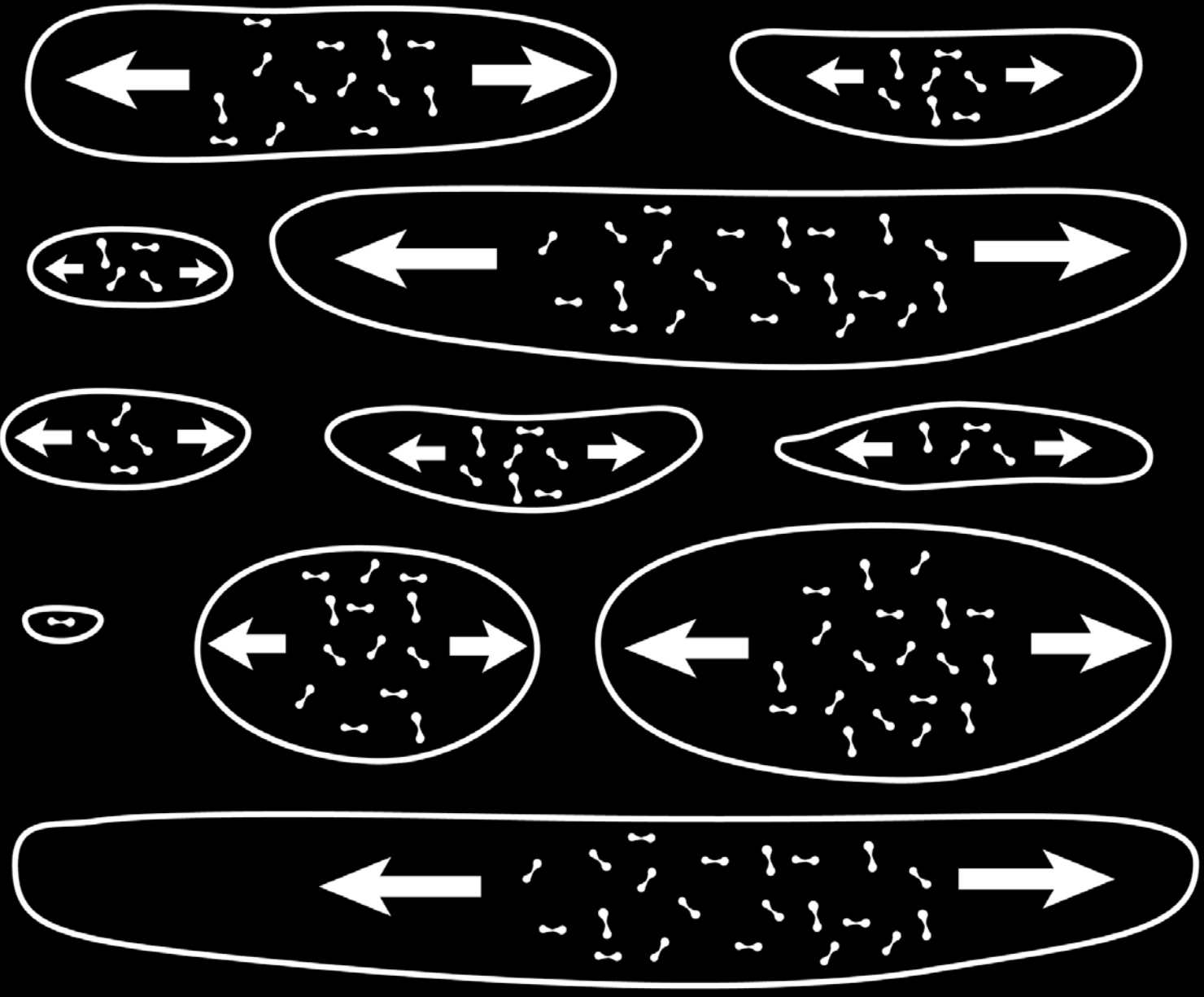


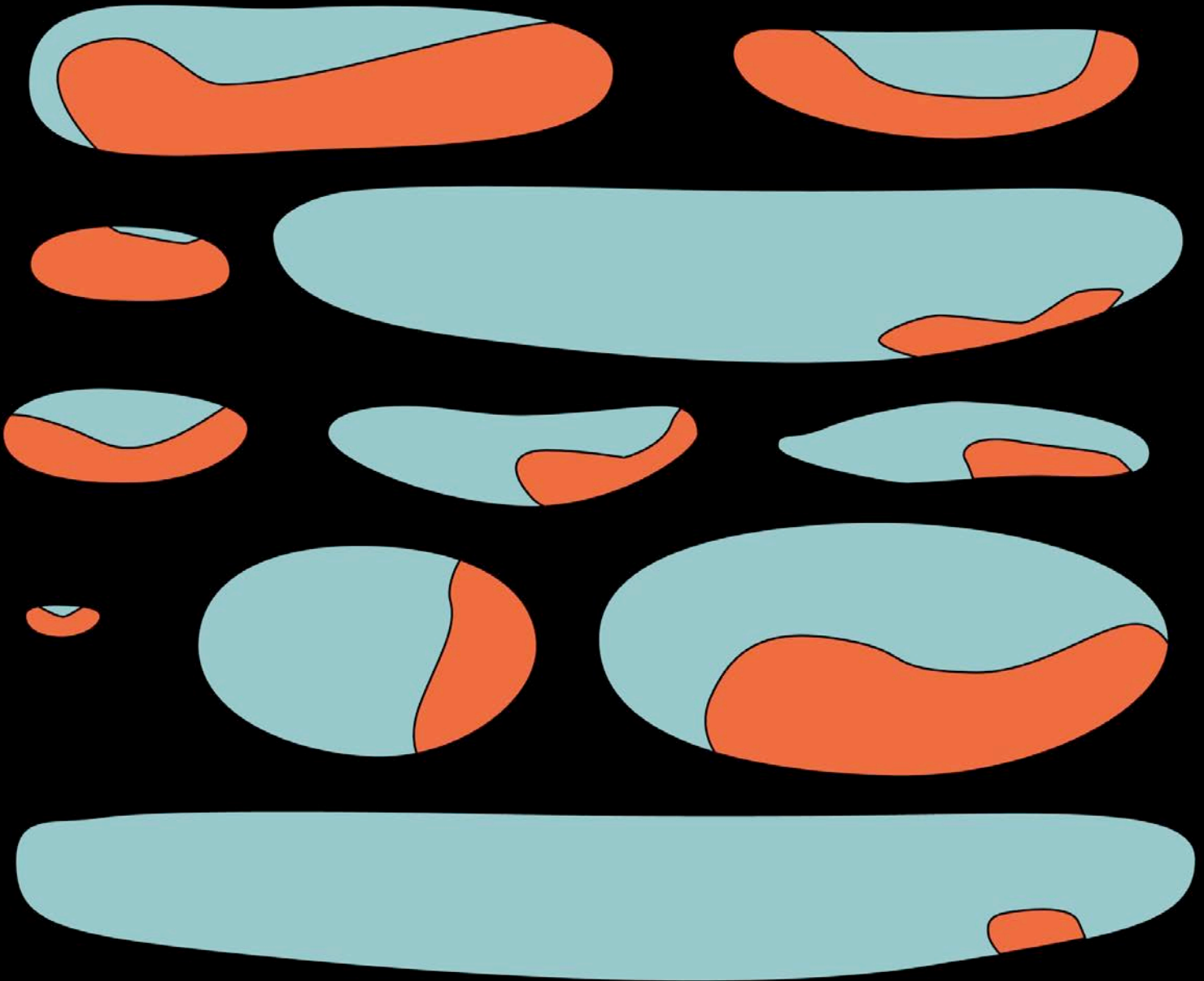
axial expansion



embryonic coalescence







extra-embryonic tissue

embryonic rudiment