

Finding a morphogen gradient using forward genetics:

Dorsal-ventral patterning by BMPs in *Drosophila* embryos

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Fruit flies...



Drosophila melanogaster

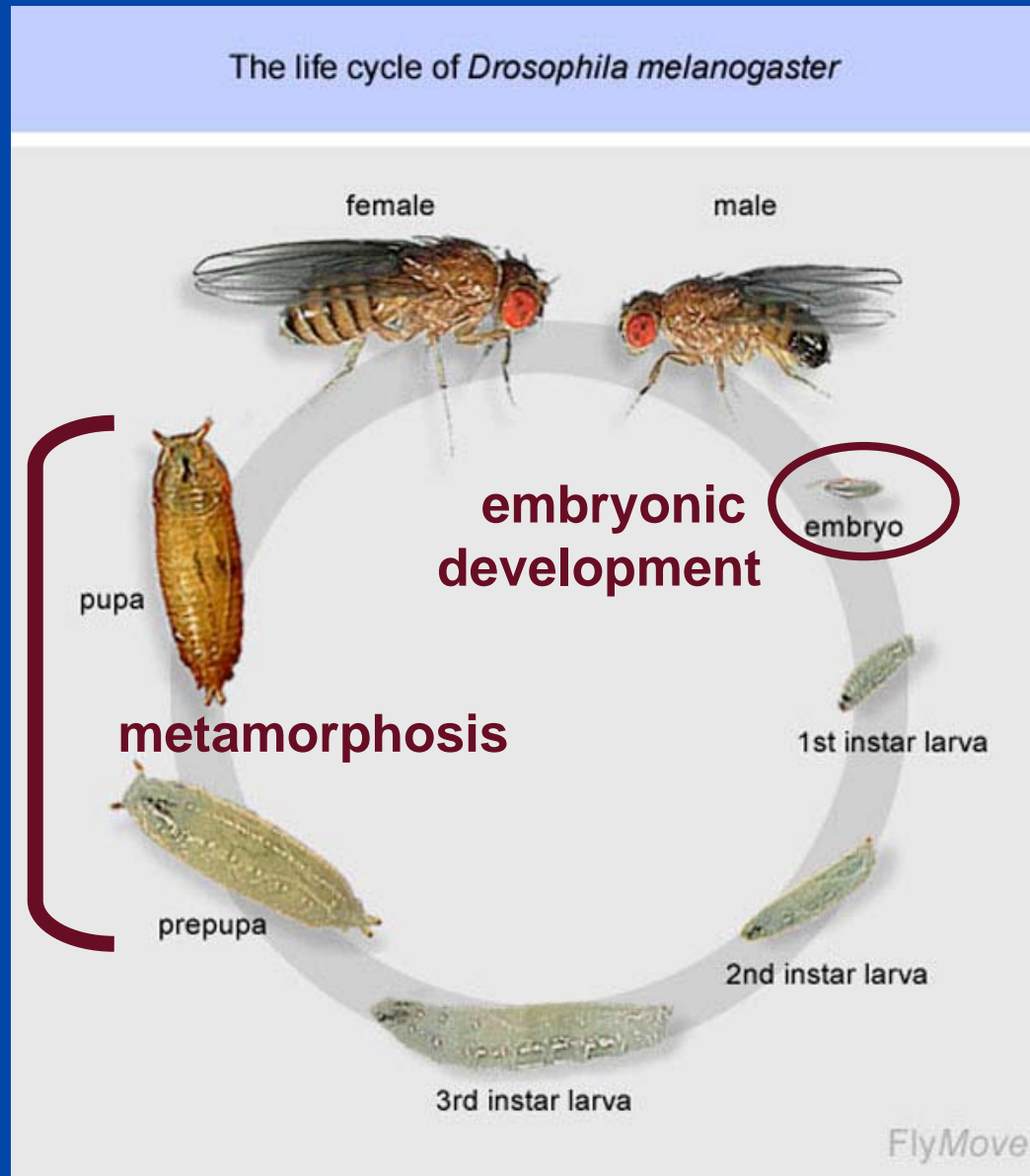
www.istoica.com/.../20060618174337_genetics.jpg



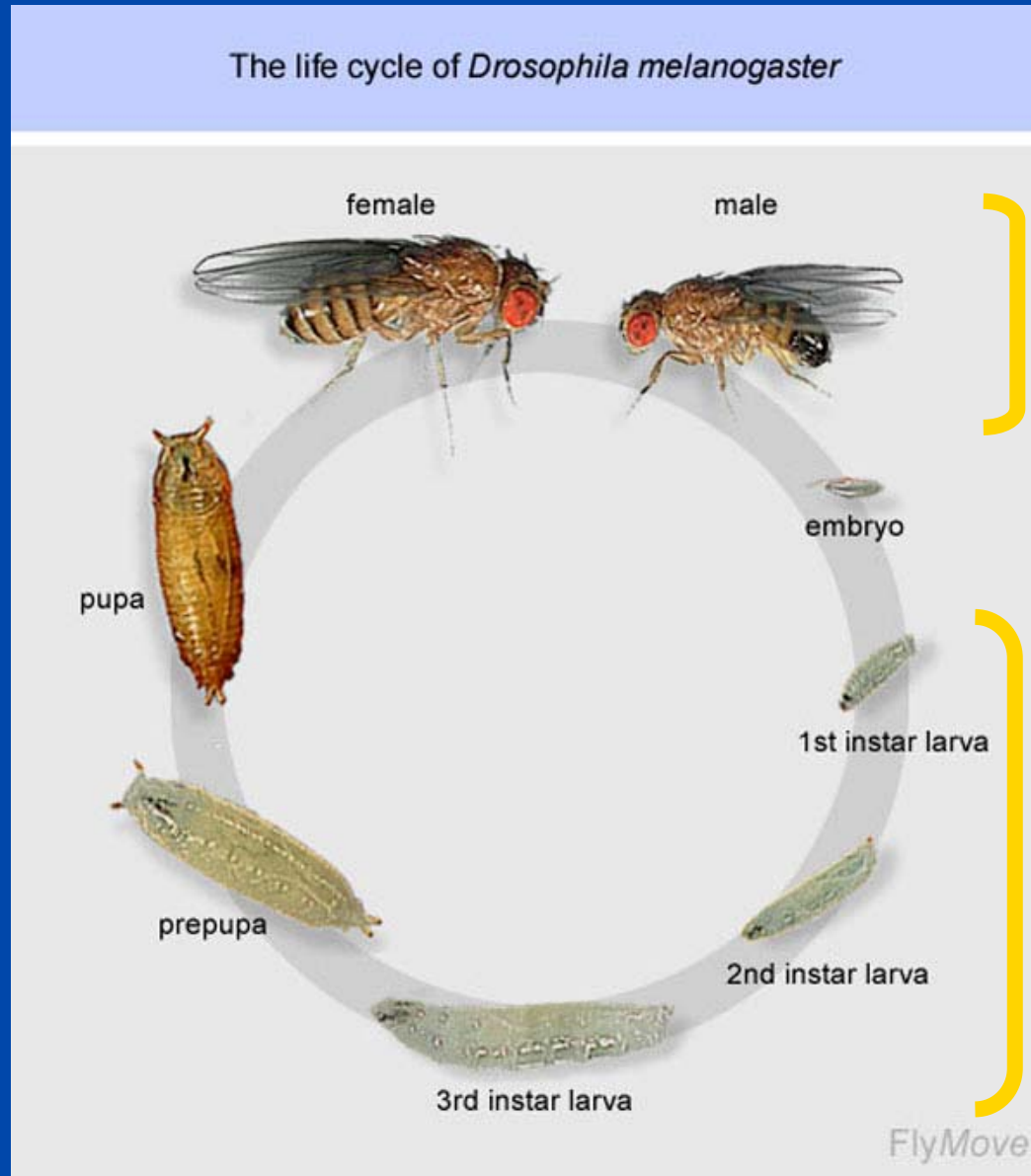
Artist's conception

www.hubbo.com/images/Fly-and-banana.jpg

Drosophila life cycle: two distinct developmental phases



Drosophila development: two different bodies for two different purposes



Dispersal and
reproduction

feeding

- **Why Drosophila genetics?**
- **What is a morphogen gradient?**
- **Inference of a morphogen gradient from genetics**
- **Identification of signaling pathway from genetics**
- **Visualization of a gradient from signal transducers**
- **Shaping the gradient**
- **Computational predictions of molecular mechanisms**

- **Why Drosophila genetics?**

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Why I chose to study *Drosophila* development:



1980

the most powerful system
to dissect the logic of
development

Nusslein-Volhard and Wieschaus

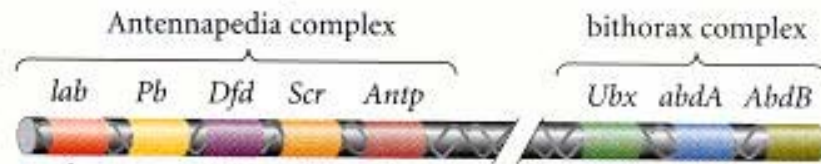
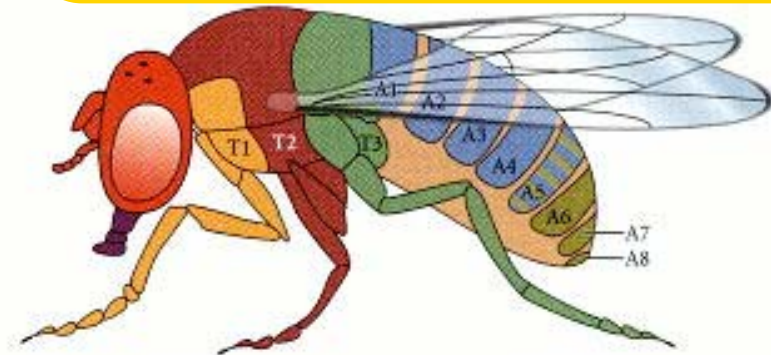


Nobel prize 1995



Ed Lewis: Homeotic mutations

"...and when that happened
a revolution occurred in the field of biology"
-Scott Gilbert



labial (lab)



Deformed (Dfd)



Sex combs reduced (Scr)



Antennapedia (Antp)



Abdominal B (AbdB)



abdominal A (abdA)

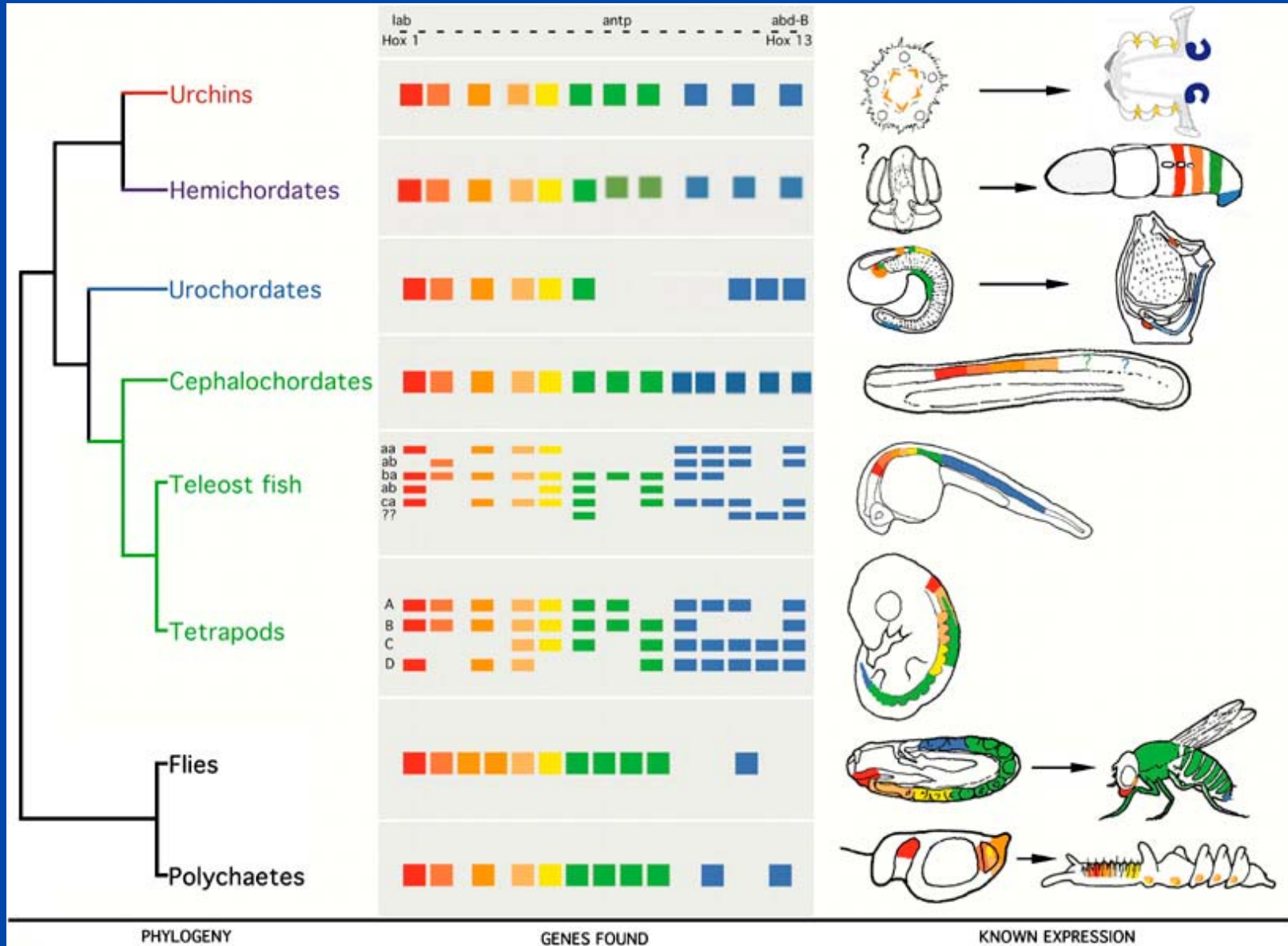


Ultrabithorax (Ubx)



From: Scott F. Gilbert, 2000. Developmental Biology 6th Edition

Homeobox genes establish body plans for all multicellular organisms



Why people choose *Drosophila* today:

- **Multicellular organisms use the same tools**
 - e.g. the gang of five for organogenesis:
 - TGF β ligands: BMPs*
 - Receptor tyrosine kinase ligands: EGFs and FGFs
 - Hedgehogs*
 - Wnts*
 - Notch ligands*
- **It is easier to understand a simple system**
 - e.g. 7 TGF β ligands in flies
 - 57 in mice
 - 34 in zebrafish

Practical reasons to study flies:

- They have a 10 day generation time
- They are inexpensive to rear
- Genetic tools are sophisticated and easy to use

- Because flies are fast and inexpensive you can
 - **Develop a developmental hypothesis**
 - **test it**
 - **Define a molecular mechanism**
 - **Test it**

all in one graduate student's thesis project

Practical reasons to study flies:

- They have a 10 day generation time
- They are inexpensive to rear
- Genetic tools are sophisticated and easy to use

- Regulators don't care about flies
- People share reagents, even before publication

- Because flies are fast and inexpensive you can
 - **Develop a developmental hypothesis**
 - **test it**
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all in one graduate student's thesis project

- Why Drosophila genetics?

- What is a morphogen gradient?

- Inference of a morphogen gradient from genetics

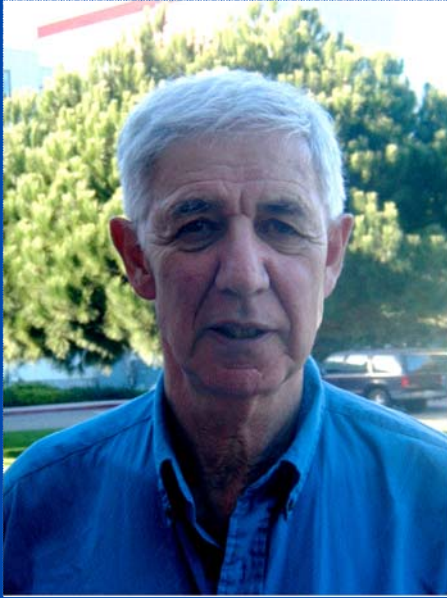
- Identification of signaling pathway from genetics

- Visualization of a gradient from signal transducers

- Shaping the gradient

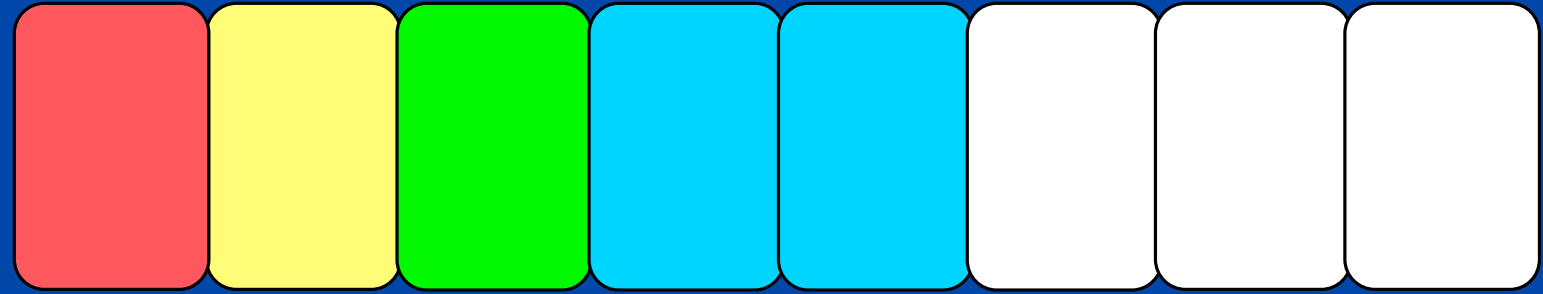
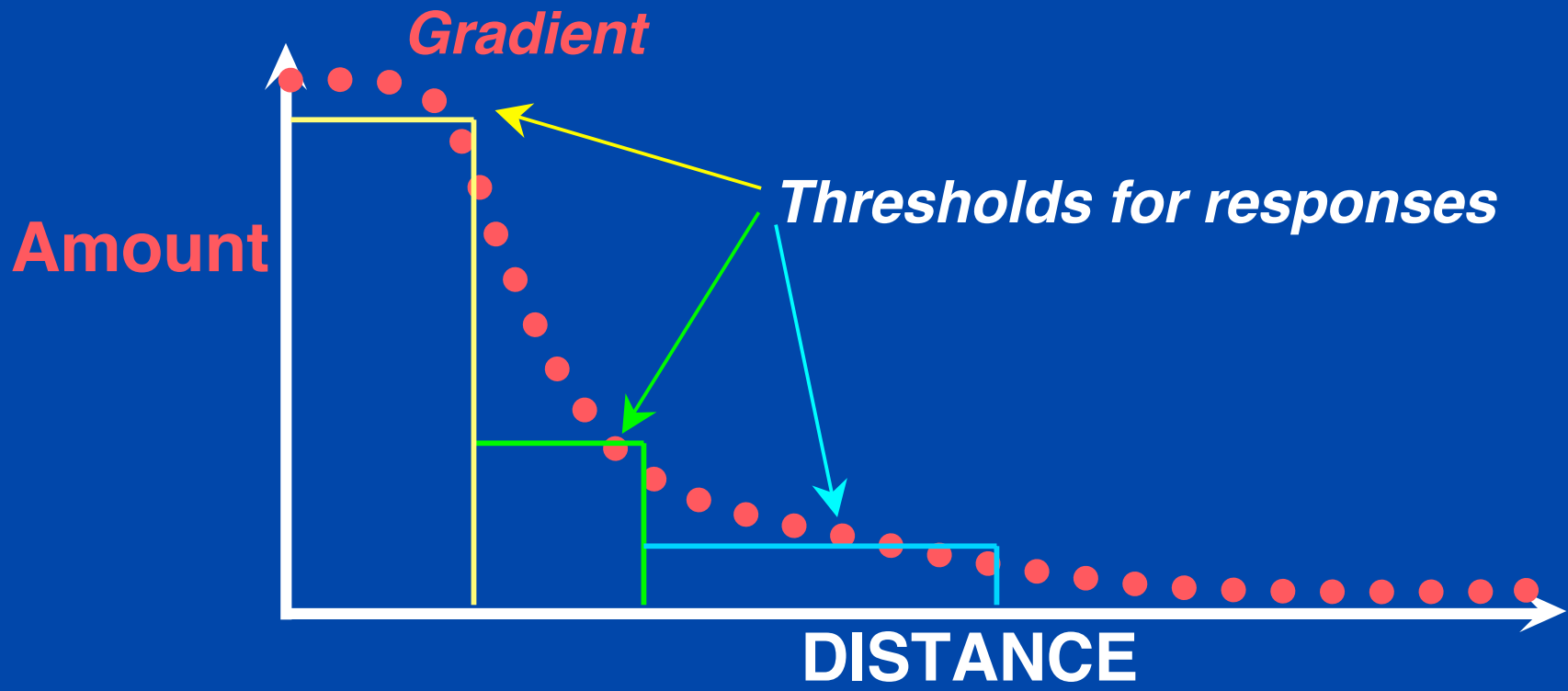
• Different cell fates can be determined by different concentrations of a **morphogen**, a diffusible molecule present at variable concentrations
(A. Turing, 1952)





2002 Curr. Biol.

A cell decides its fate based on its position within the tissue. The pattern of cell fates is established through a coordinate system of positional information.
(L. Wolpert, 1969)

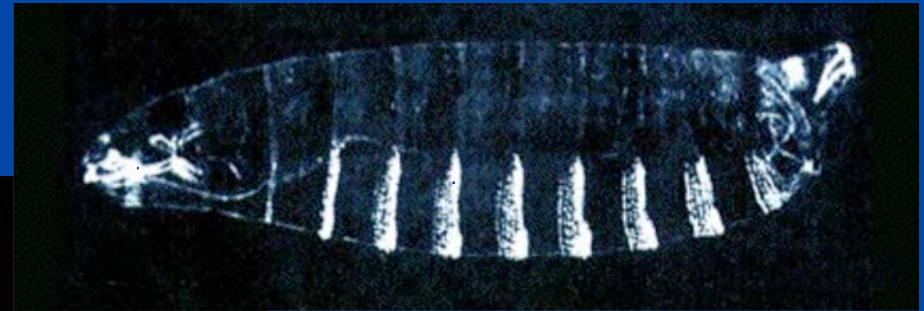


Source

Sink

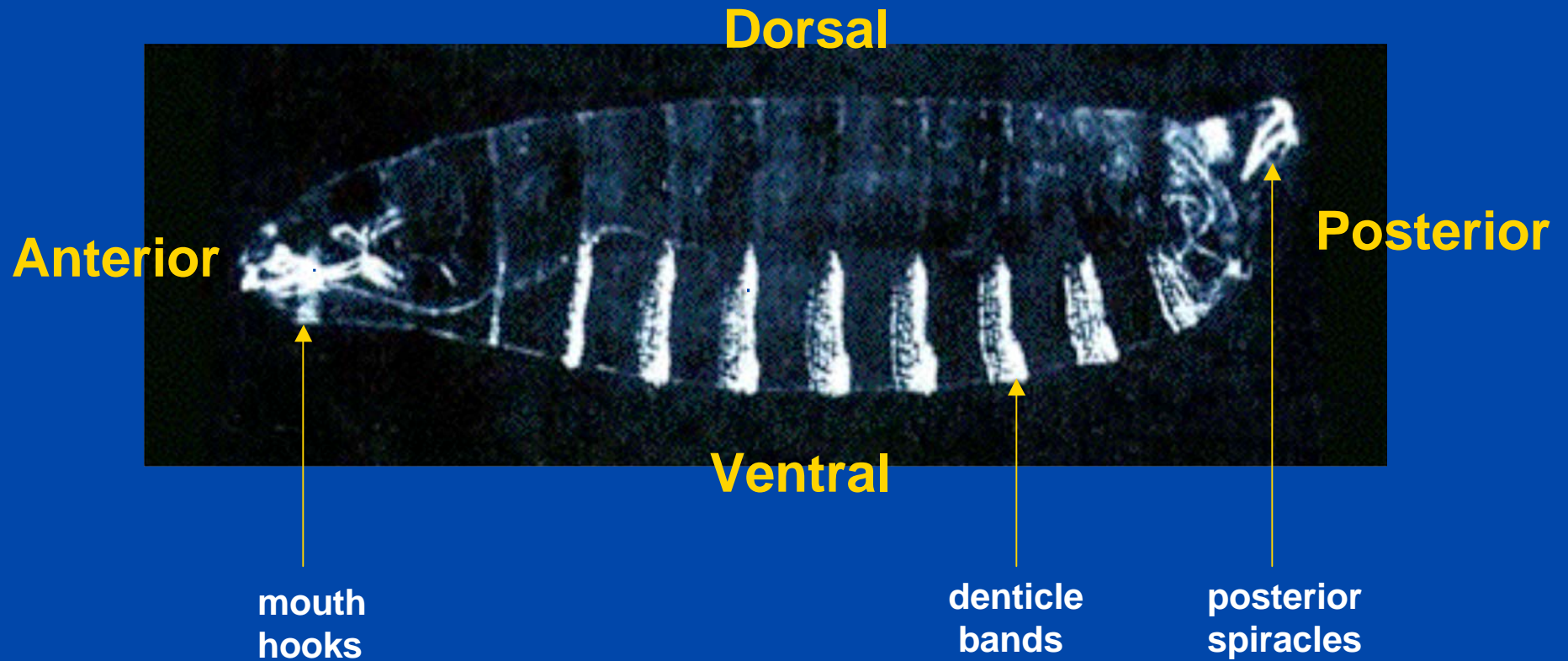
- Why *Drosophila* genetics?
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The *Drosophila* body plan is prepatterned in the egg

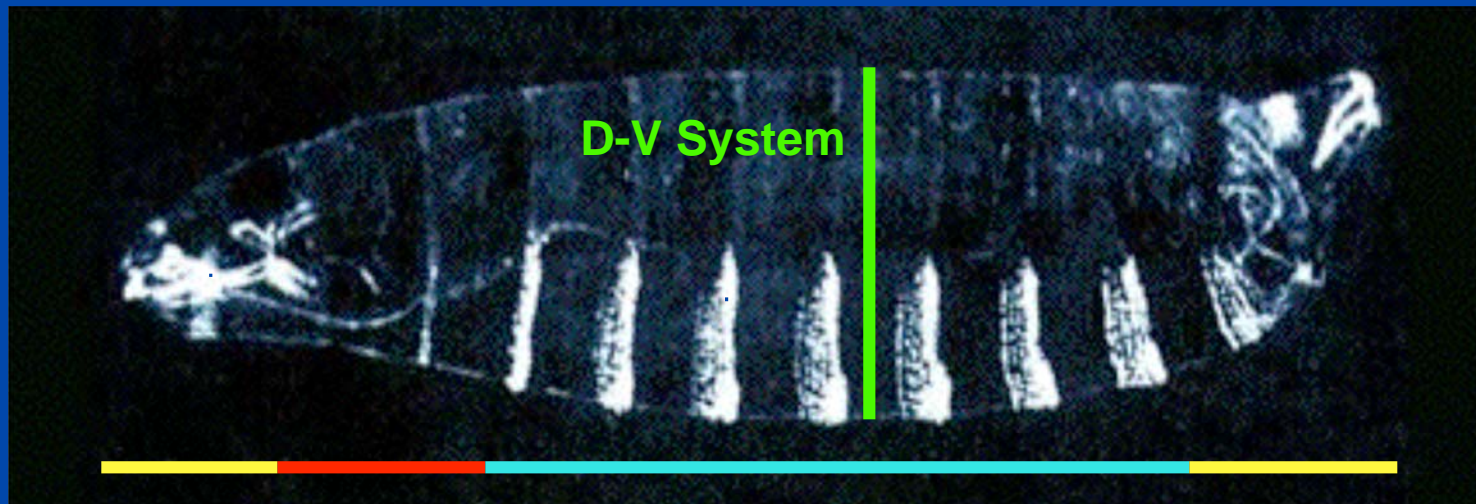


By RNA and protein from the mother

The *Drosophila* body plan is visible in the exoskeleton (cuticle)



A set of maternal genes establishes each aspect of the body plan



Anterior System

Posterior System

Terminal System

Maternal genes were identified by embryo phenotypes

Anterior
posterior
termini and
dorsal-ventral

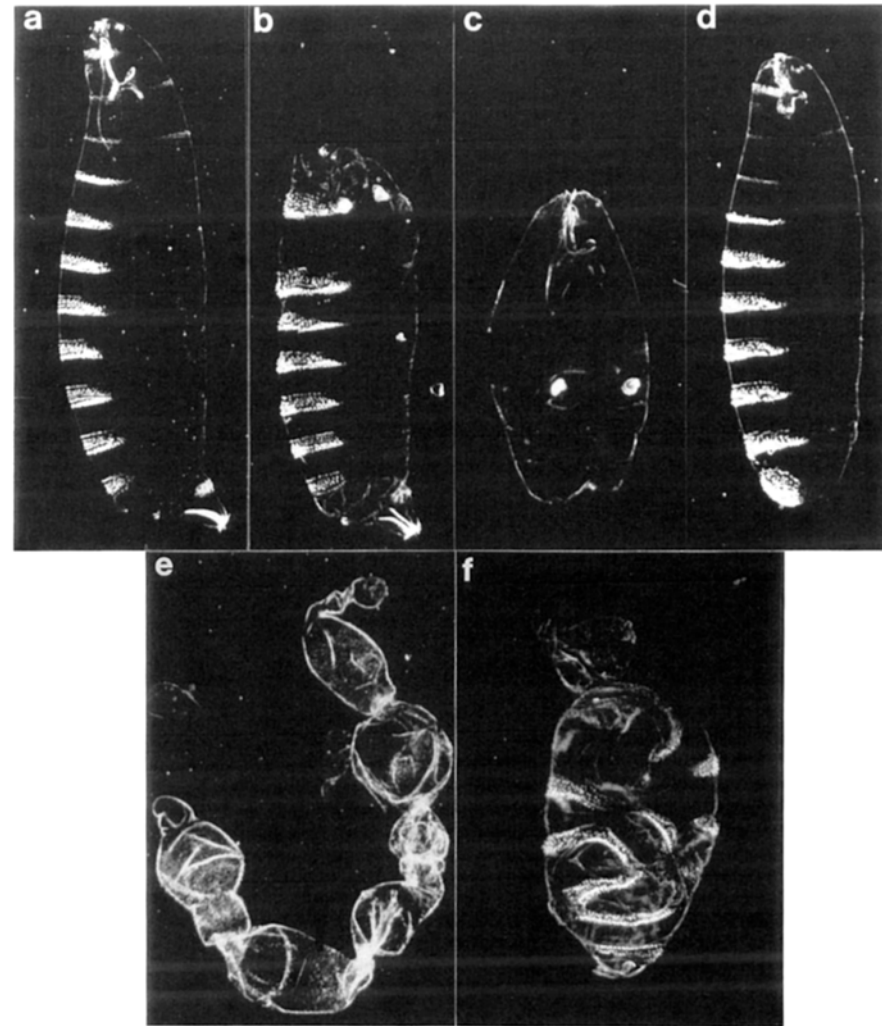


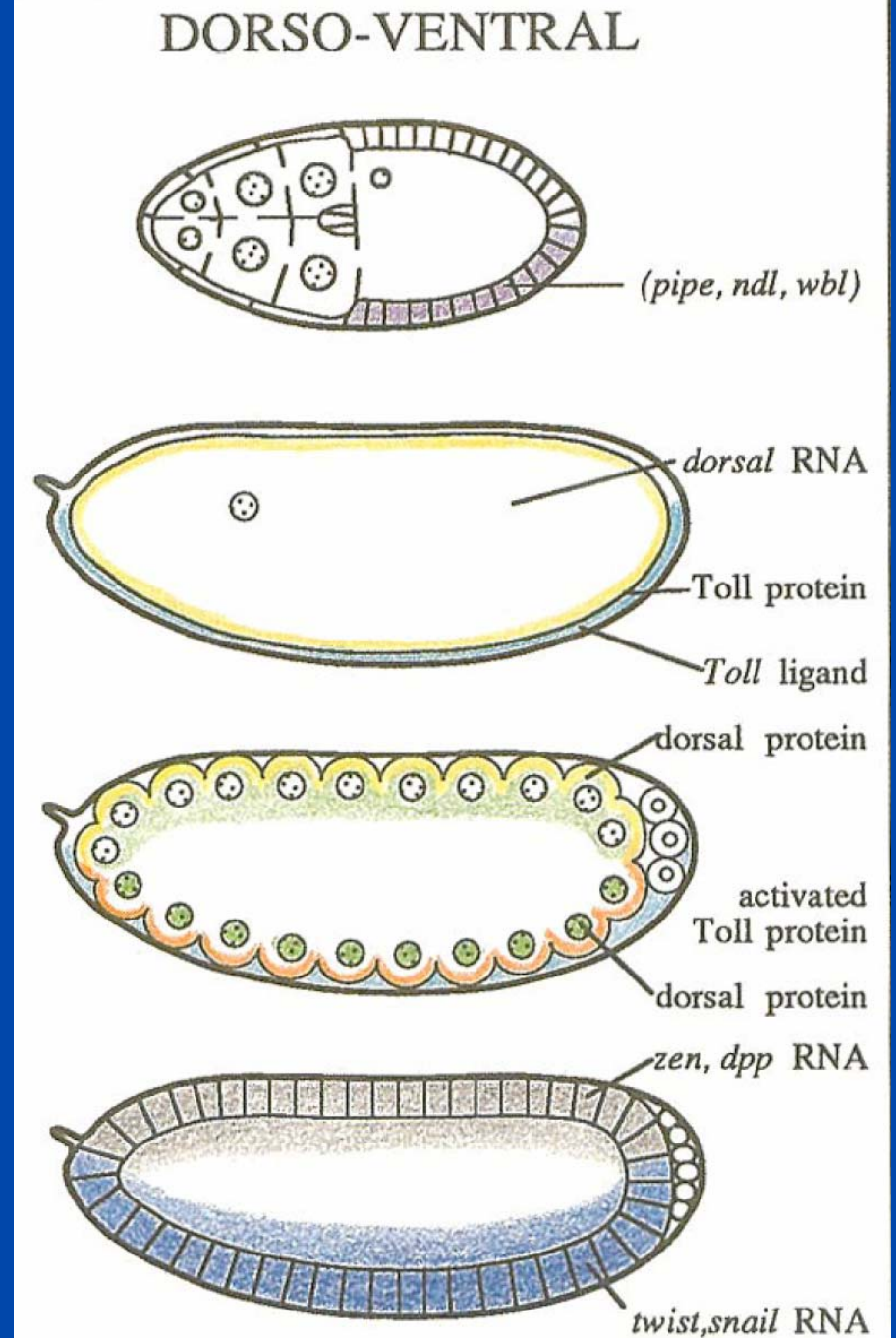
Figure 3. The Cuticular Patterns of Wild-Type and Mutant Embryos

(a) wild-type, (b) anterior (*bicoid*), (c) posterior (*oskar*), (d) terminal (*torso-like*), (e) dorsalized (*dorsal*), (f) ventralized (*cactus*).

Dorsal-ventral axis is patterned by sequential morphogen gradients

**Maternal genes:
NF- κ B gradient subdivides mesoderm/ectoderm**

**Zygotic genes:
BMP gradient subdivides ectodermal territories**



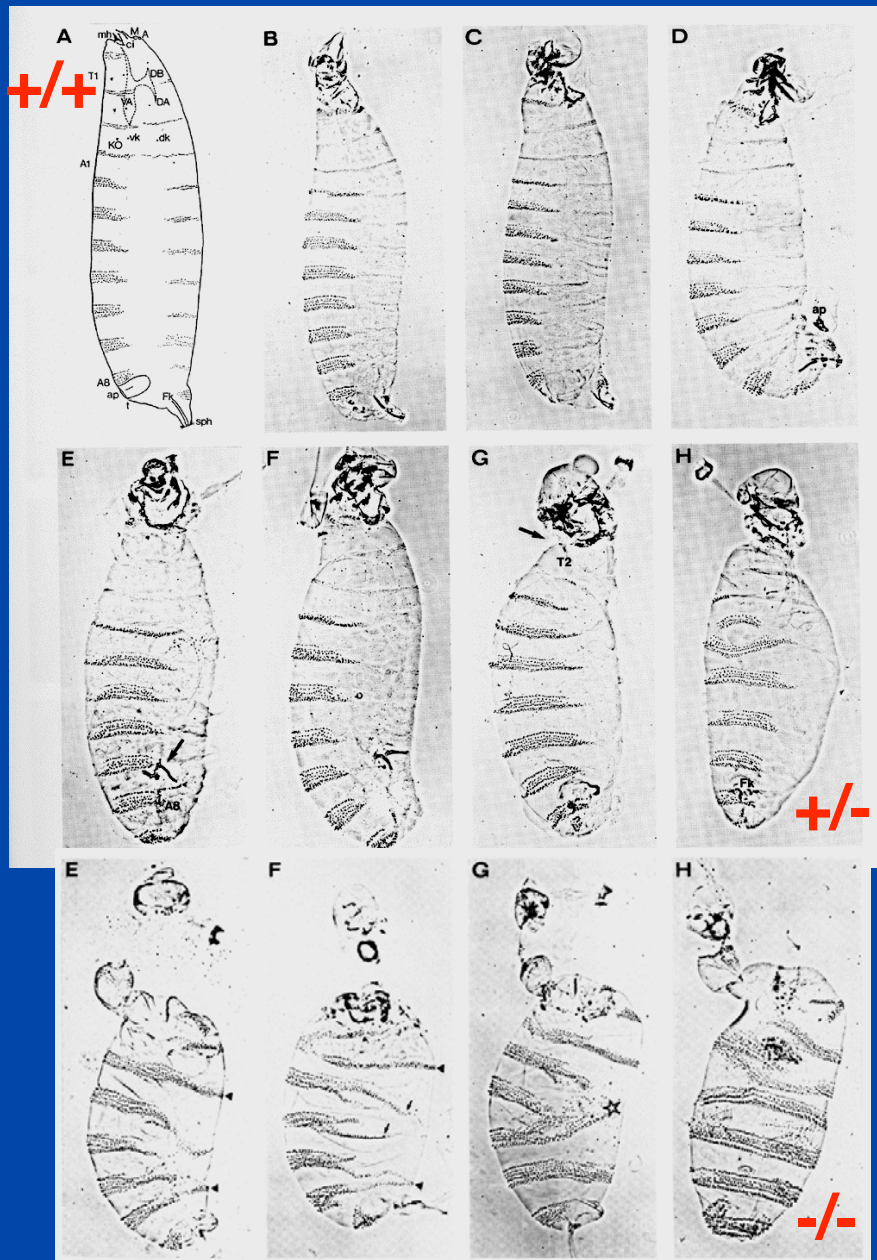
Why a *gradient* of Dpp in embryos?

- Uniform RNA levels, protein undetectable
- ⇒ **An inference from the genetic analysis of ectoderm patterning**

• Dpp was known to be a BMP, and hypothesized as a morphogen for appendage development.

Why a *gradient* of Dpp in embryos?

- Uniform RNA levels, protein undetectable
 - ➡ An inference from the genetic analysis of ectoderm patterning
- Different Dpp mutations cause loss of more or fewer dorsal ectoderm structures
 - Wharton et al, 1993, Development
- Dpp was known to be a BMP, and hypothesized as a morphogen for appendage development.



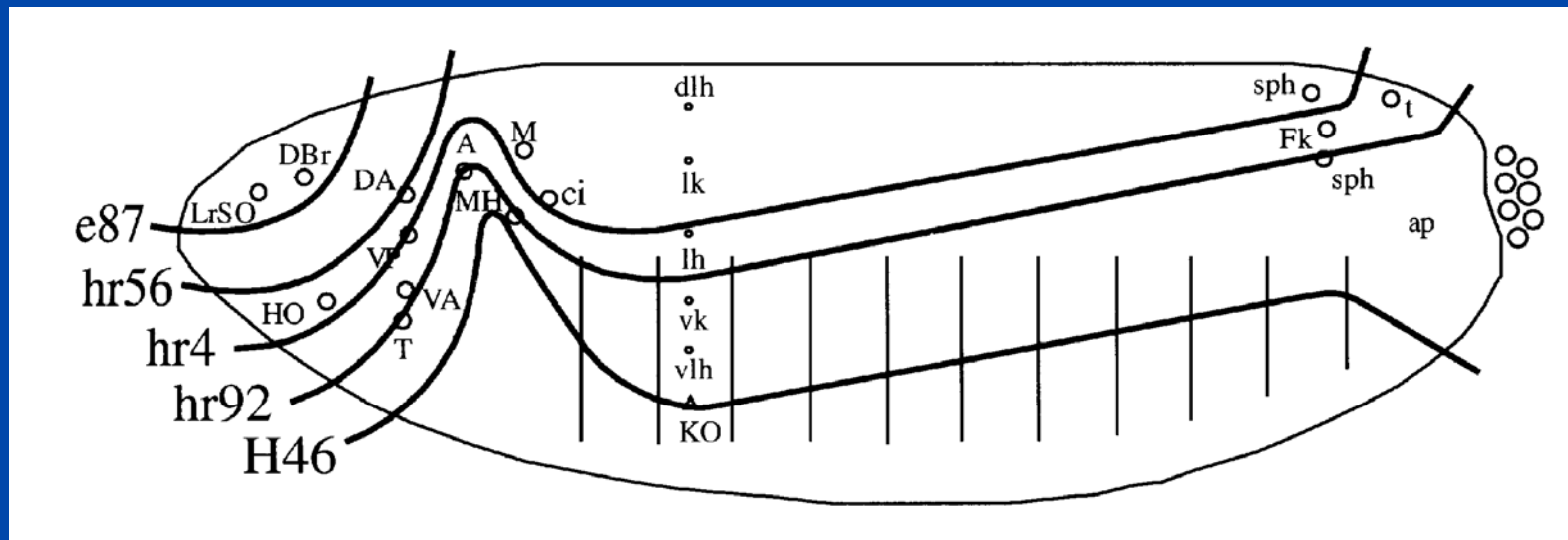
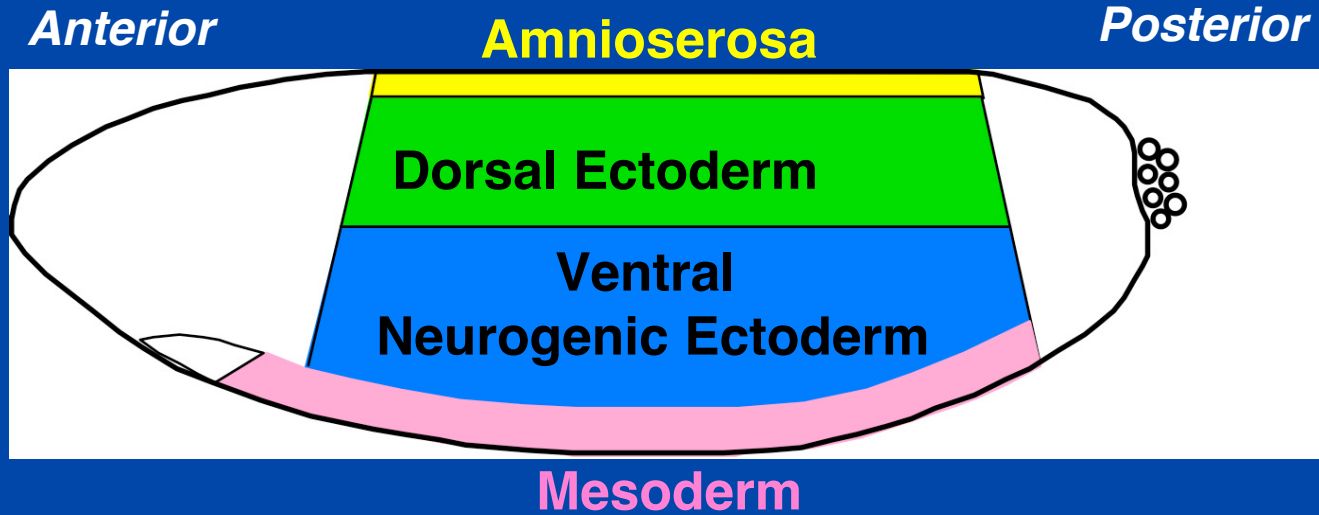
Different mutations in *dpp* eliminate different amounts of dorsal fates

Dpp is *necessary* to make dorsal ectoderm and amnioserosa

Irish & Gelbart, 1987, *Genes Dev.* 1, 868
 Wharton et al. 1993, *Development* 117, 807

The less DPP, the fewer dorsal structures

Lateral View

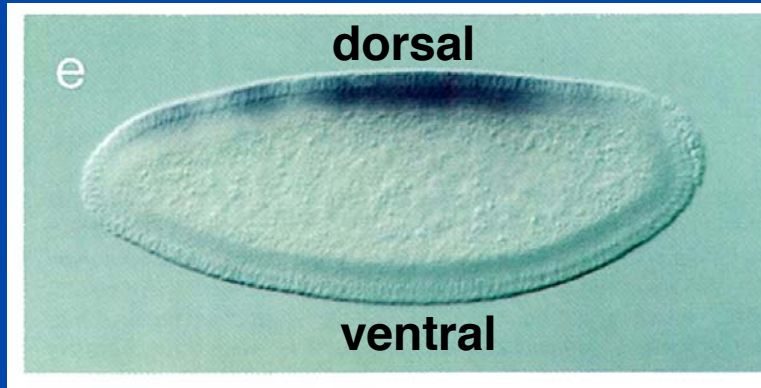


Why a *gradient* of Dpp in embryos?

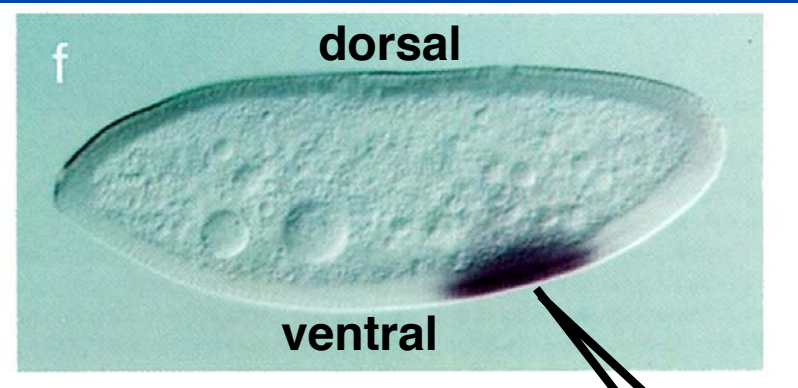
- **Uniform RNA levels, protein undetectable**
- **Different Dpp mutations cause loss of more or fewer dorsal ectoderm structures**
 - Wharton et al, 1993, Development
- **Injections of Dpp RNA induced dorsal fates**
 - Ferguson and Anderson, 1992, Cell

Dpp RNA is *sufficient* to induce dorsal fates

zen is expressed in
amnioserosa primordium



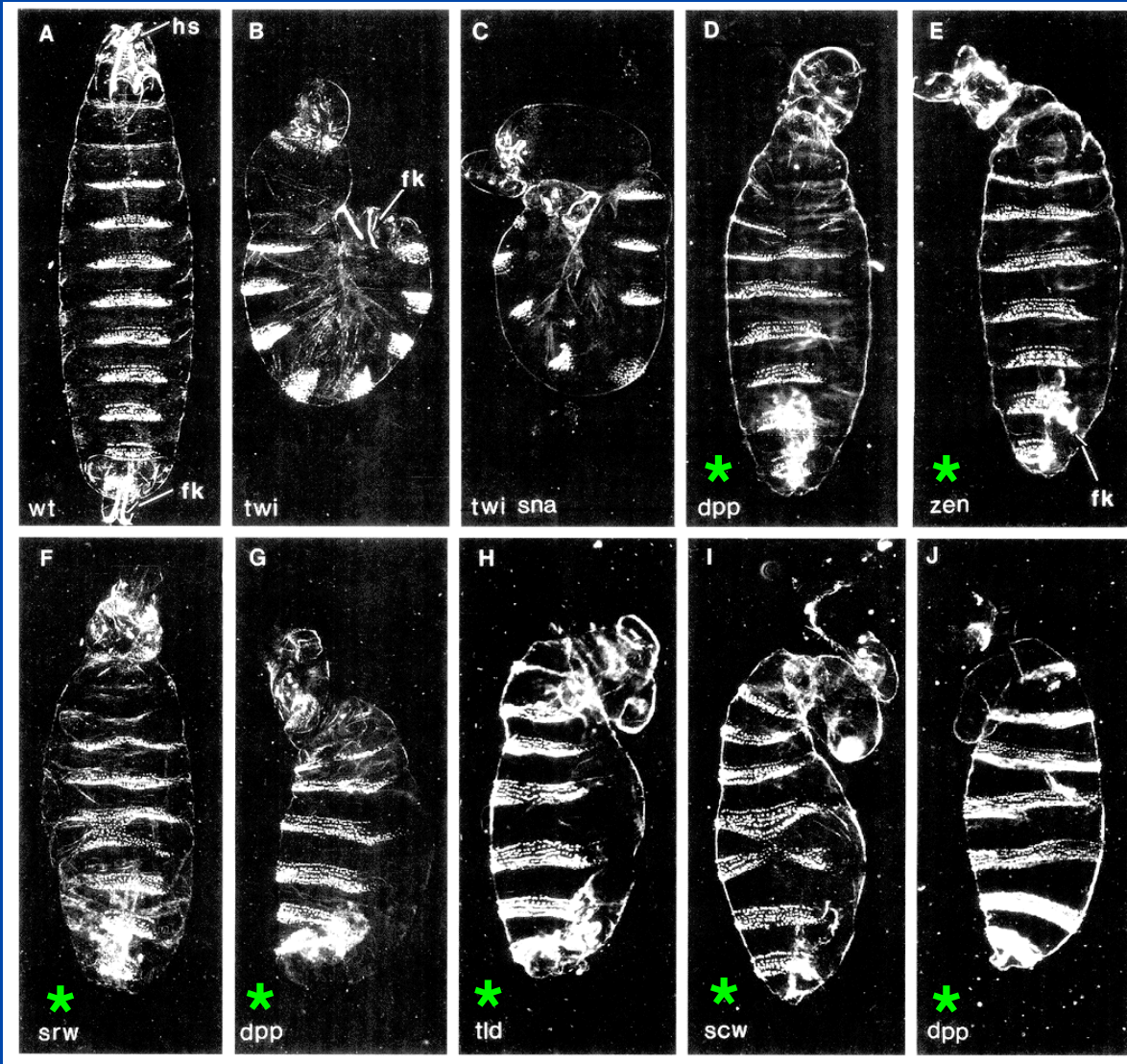
zen expression is induced in
ventral cells by Dpp RNA



Dpp
RNA

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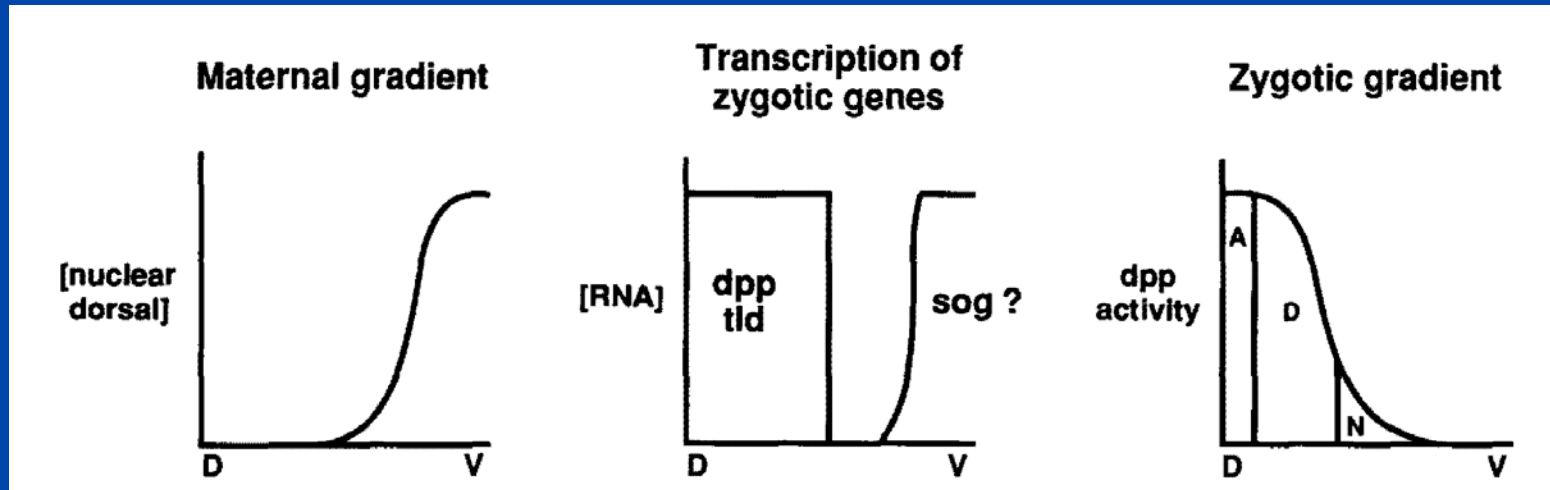
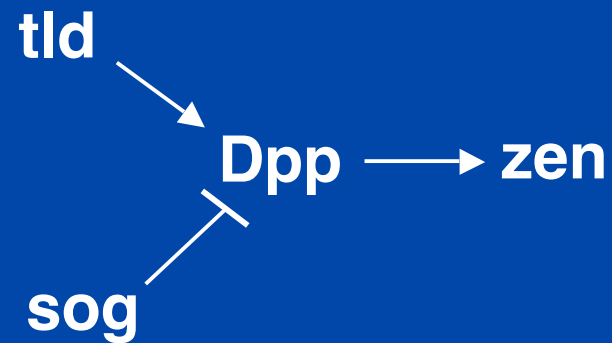
Mutational screens identified classes of genes with shared D-V patterning defects

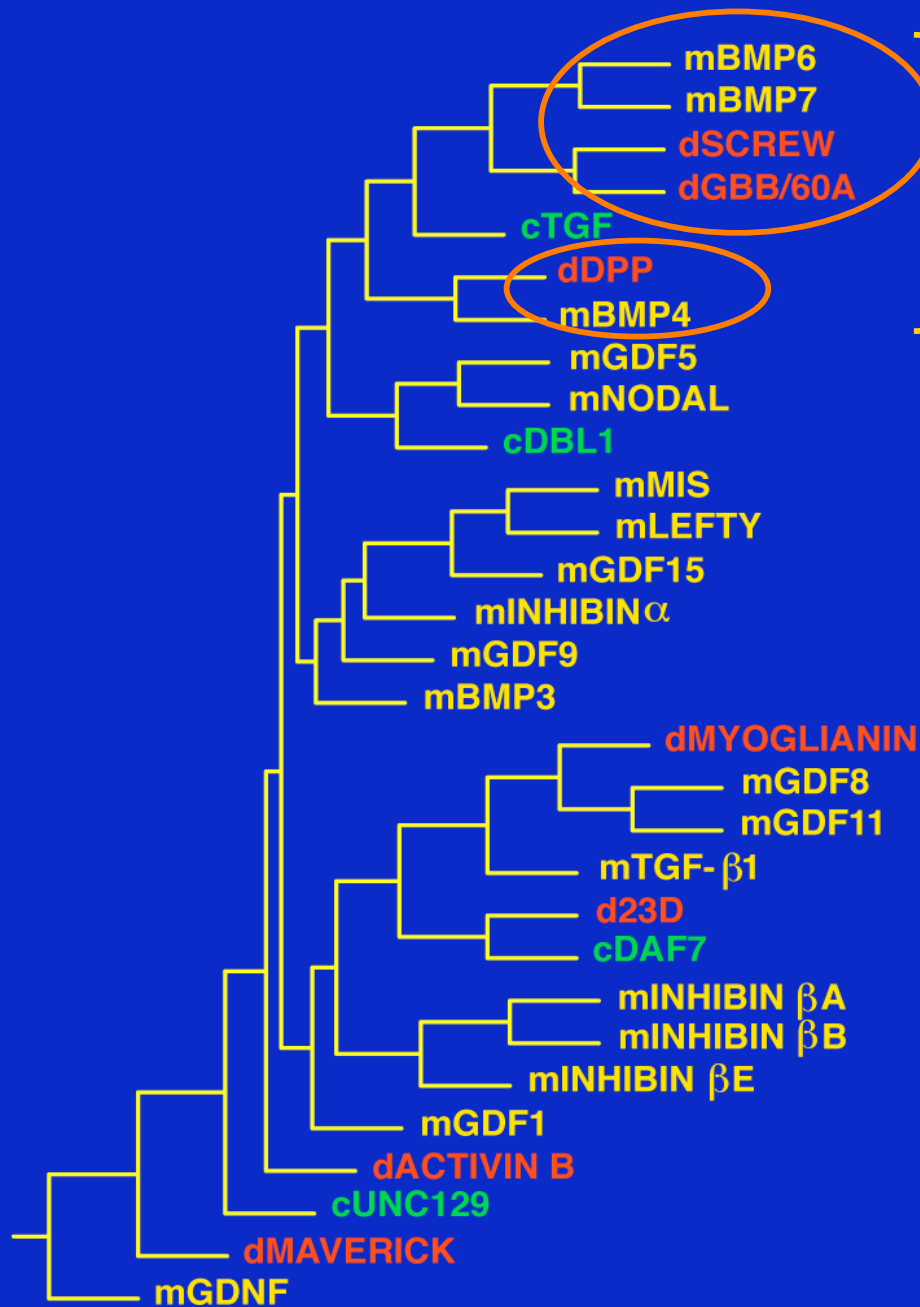


tld and *scw*
are almost as
severe as *dpp*

Arora et al, 1992,
Development 114, 1003

Dpp activity is modulated by *tld* and *sog*





Dpp and Screw are BMPs

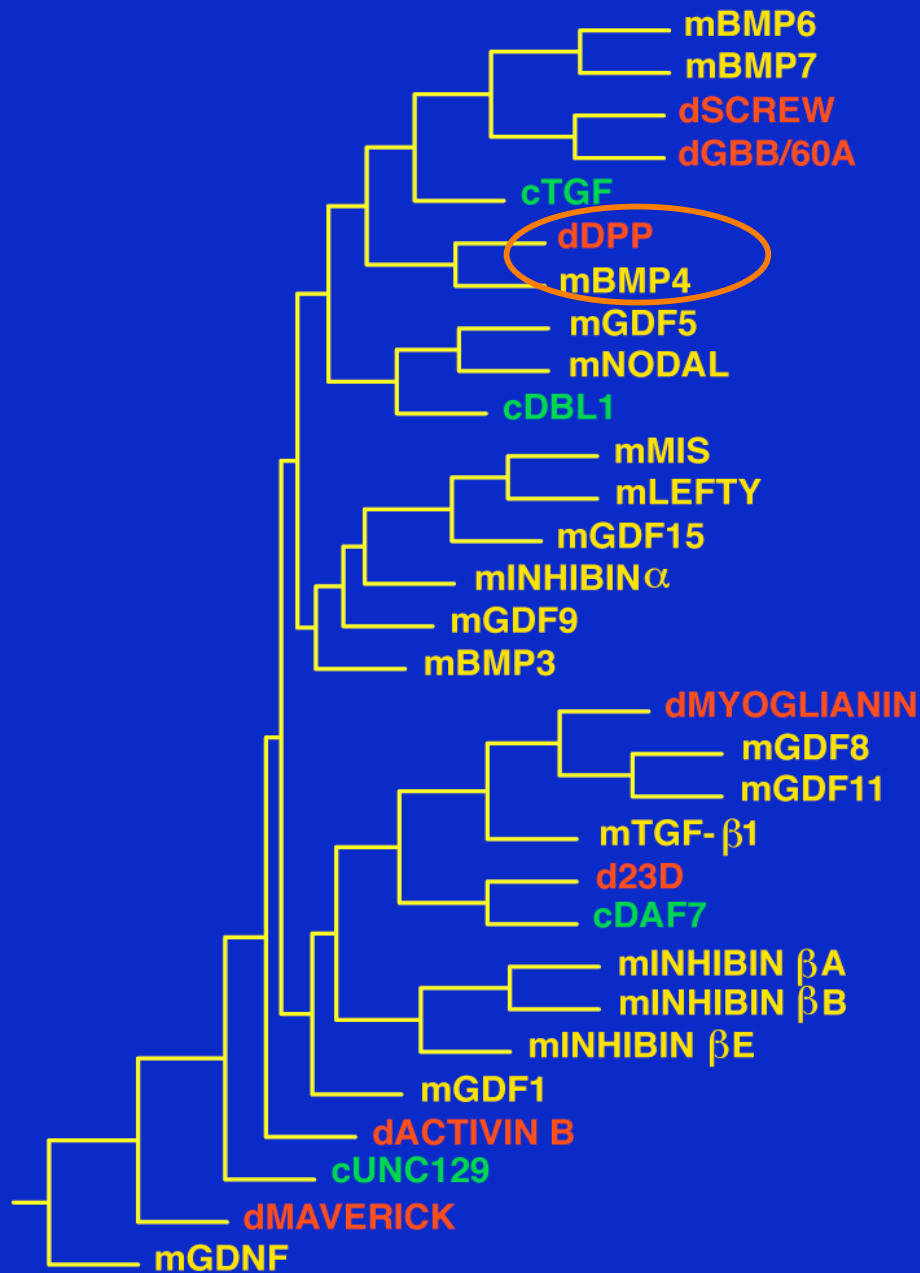
(bone morphogenetic proteins)

TGF β proteins organized according to sequence similarity: A phylogenetic tree

Mouse 57

Fly 7

Worm 4



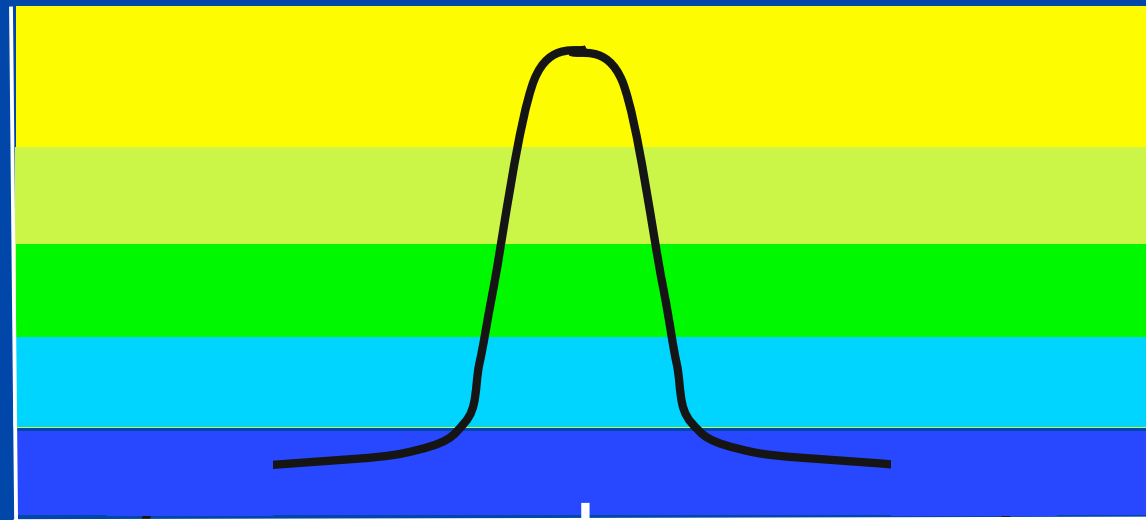
Experimental tests say that the most similar BMPs are from different species:

Fly DPP can induce bone formation in rats.

Mouse BMP2 can replace fly DPP in dorsal-ventral patterning.

DPP, BMP2, BMP4 are orthologs

BMPs regulate nested patterns of gene expression



Race

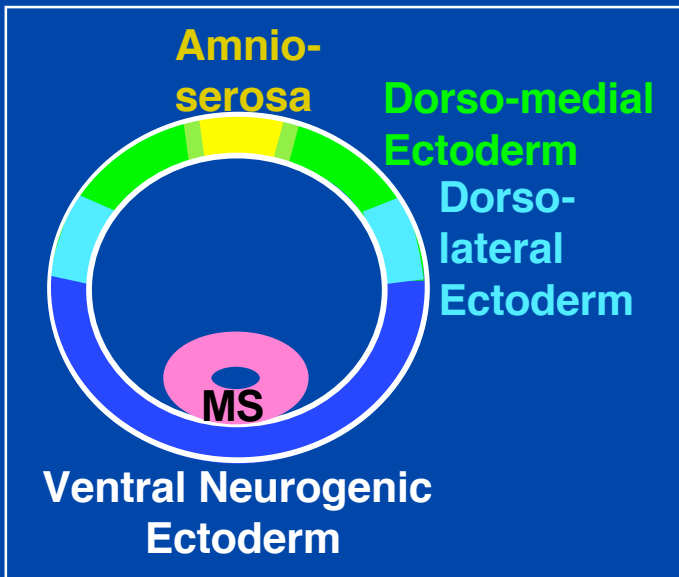
hindsight

ush

tup

pannier

dpp



reviewed by
Raftery & Sutherland, 2003 Trends Genet. 19, 701

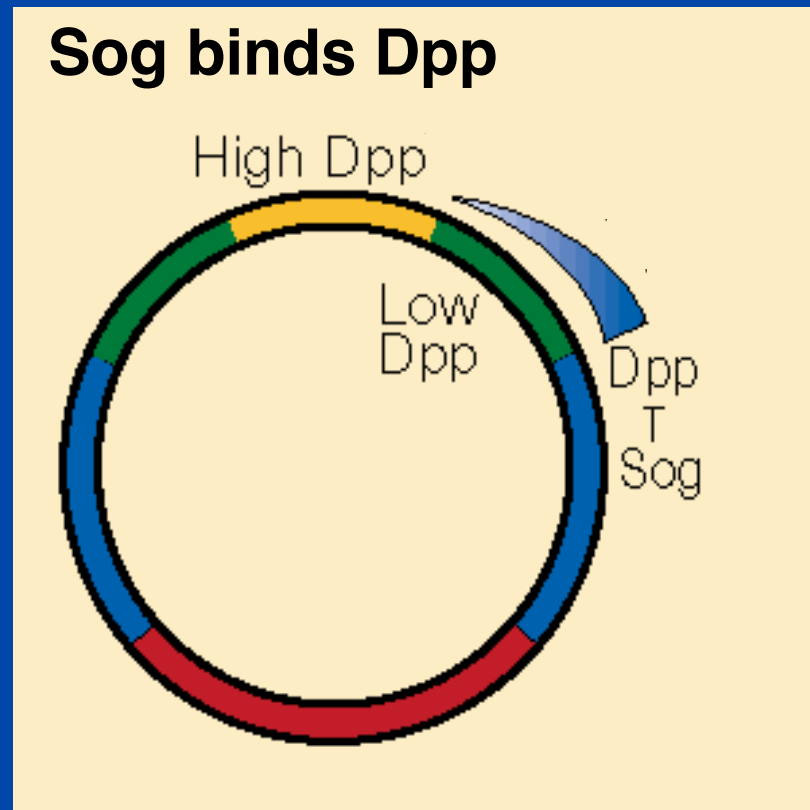
Model: Sog removes BMPs at low end of gradient

Screw is everywhere

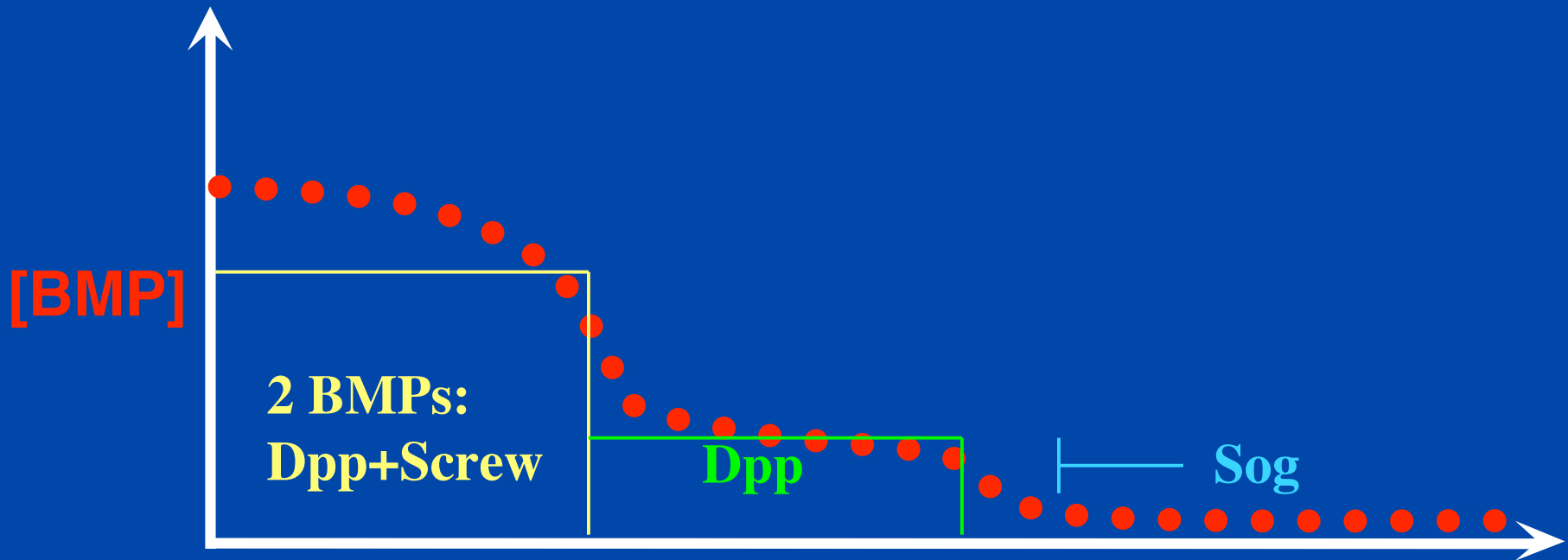


Jazwinska et al, 1999,
Development 126, 3323

Sog binds Dpp



Model: a BMP activity gradient at blastoderm induces dorsal ectoderm fates



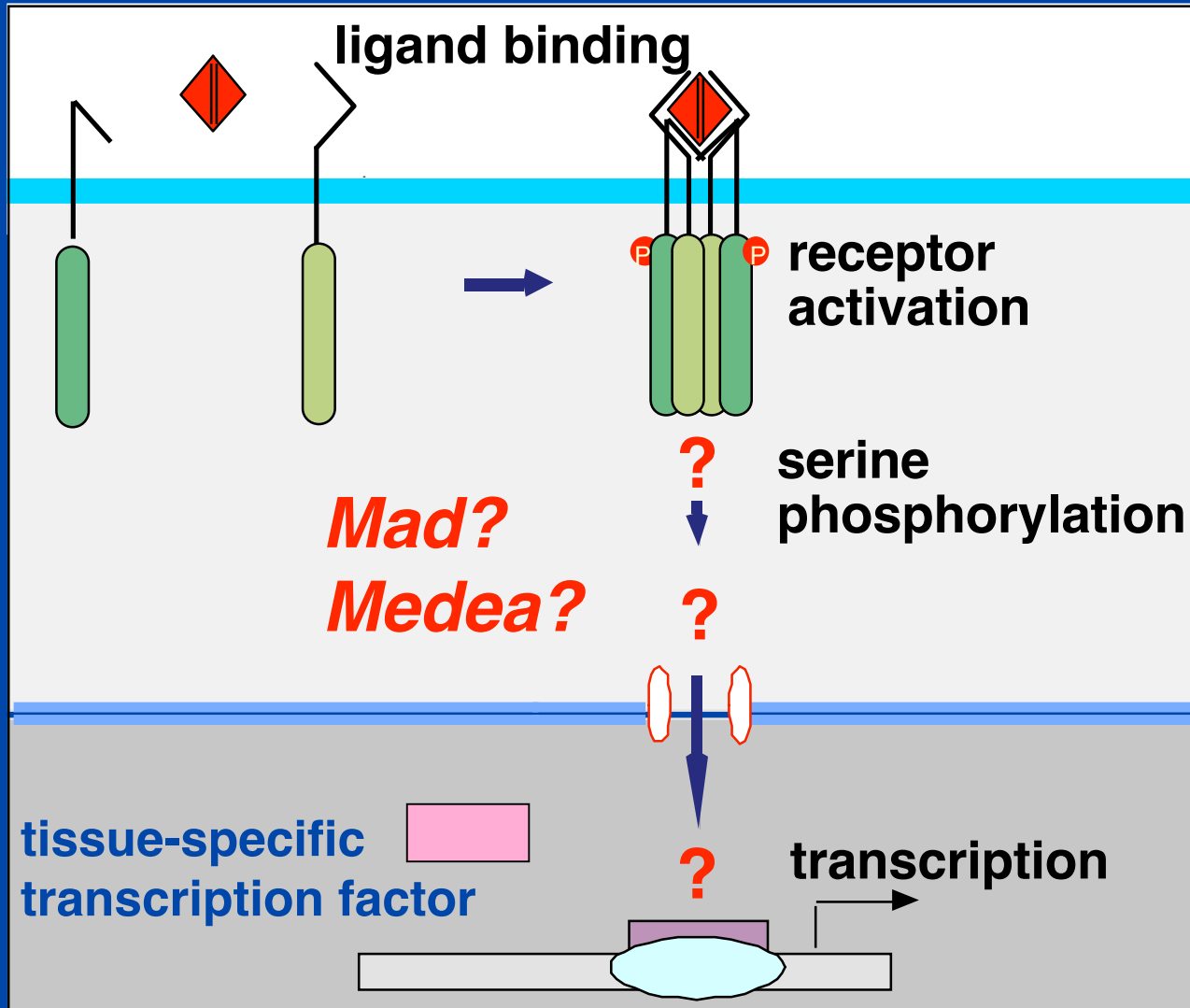
Amnioserosa

**Dorsal
Ectoderm**

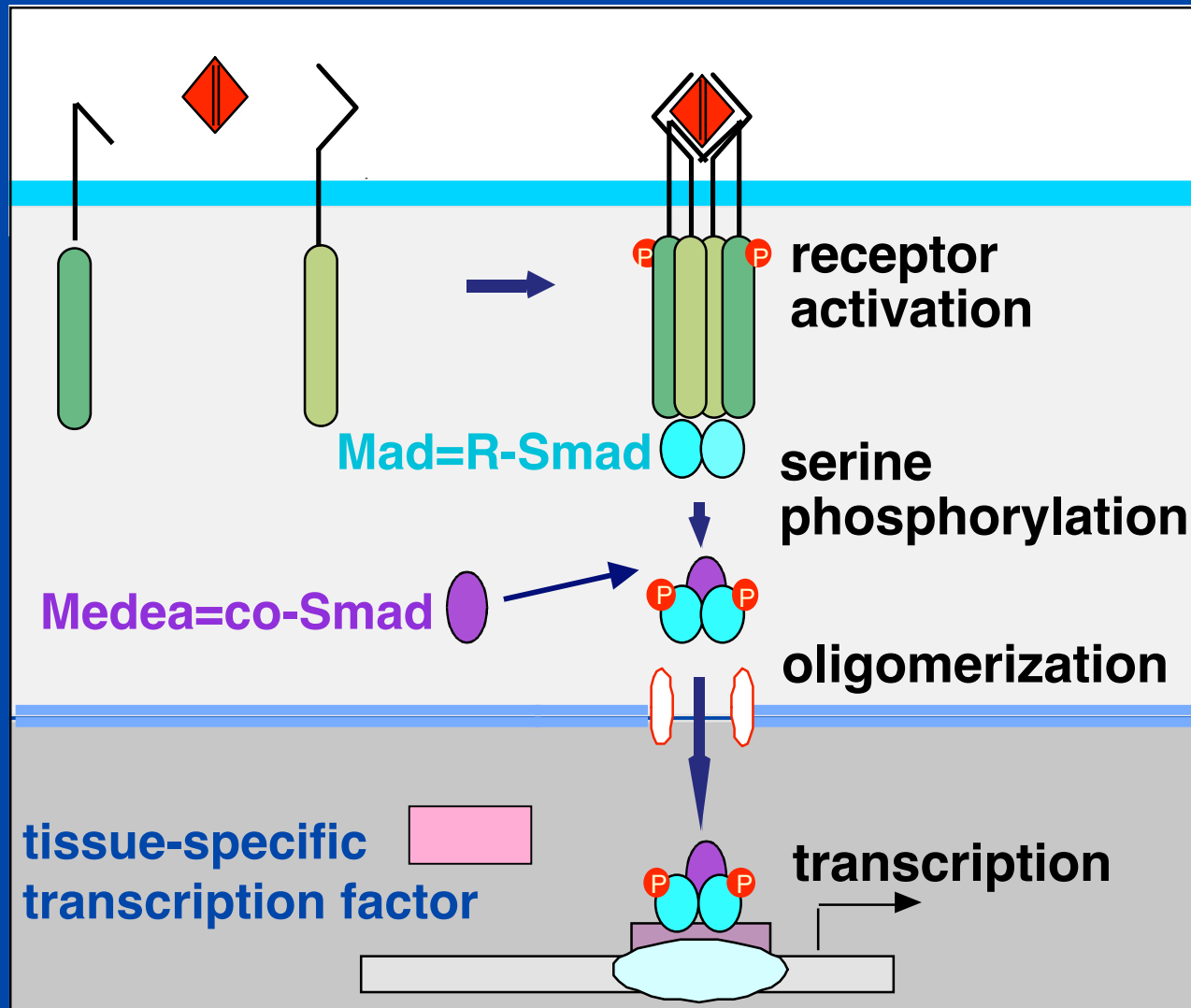
**Ventral
Neurectoderm**

How do cells respond to the BMP gradient?

Mad and Medea identified in screens for genes that interact with Dpp



Smad proteins transduce TGF β family signals to the nucleus



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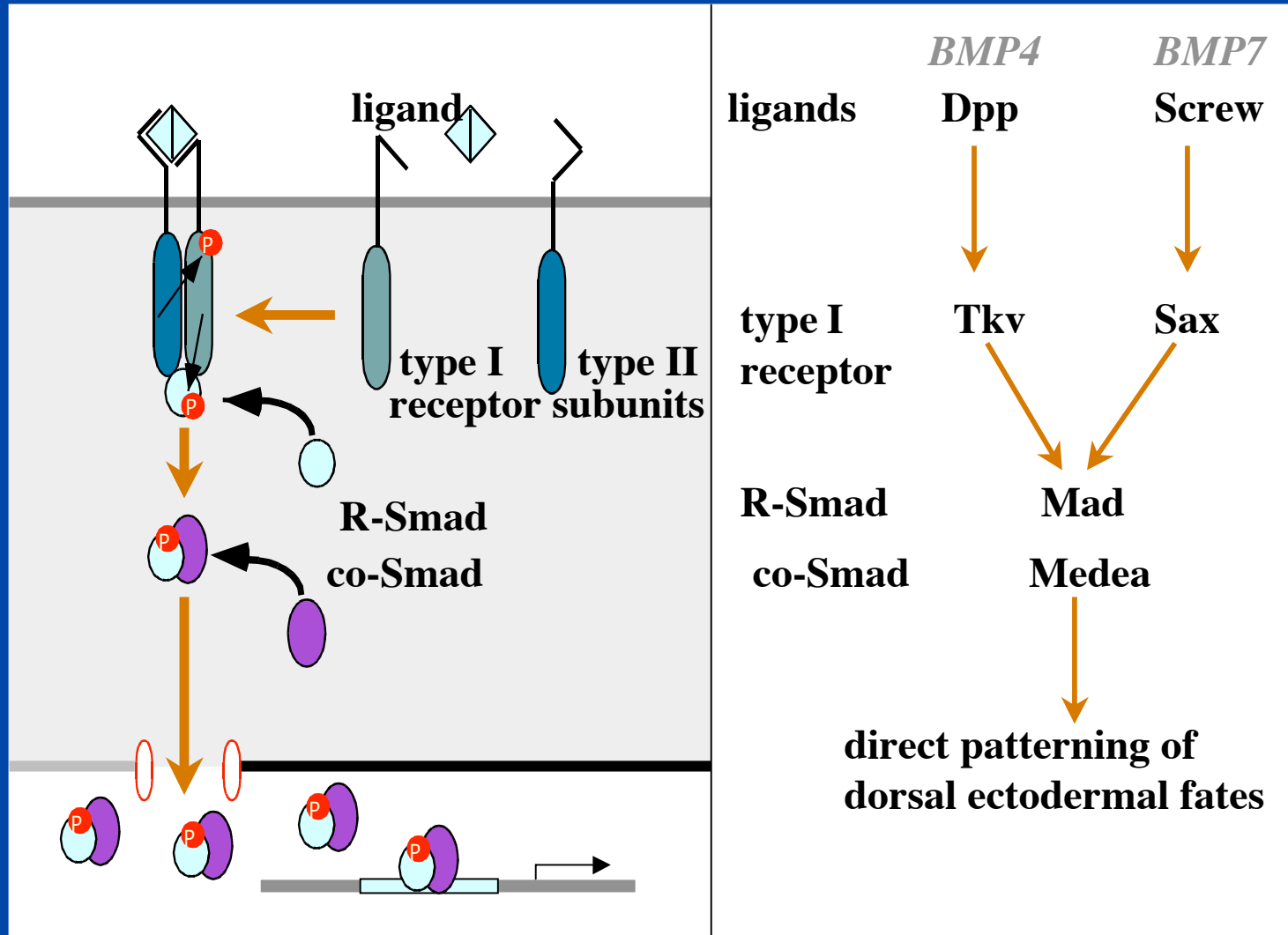
How can we visualize a BMP activity gradient?

- Dpp RNA expression is low
- Dpp protein hasn't been visualized
- Graded distribution of diffusible antagonist, Sog

Srinivasan et al, 2002, Dev. Cell 2, 91

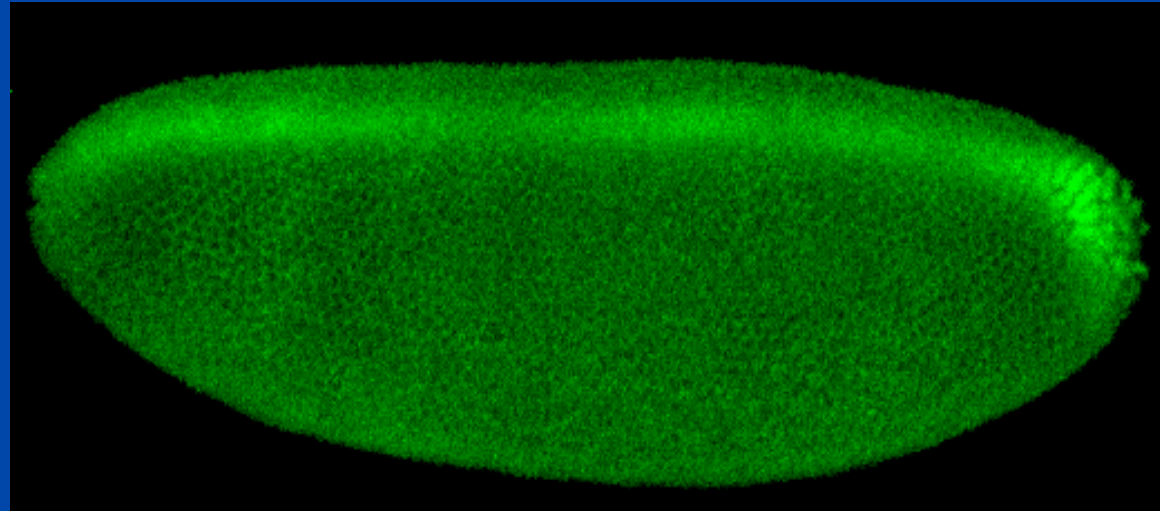
 Look at distribution of activated signal transducers

Medea nuclear localization and phospho-Mad predicted to indicate level of BMP activity *in vivo*

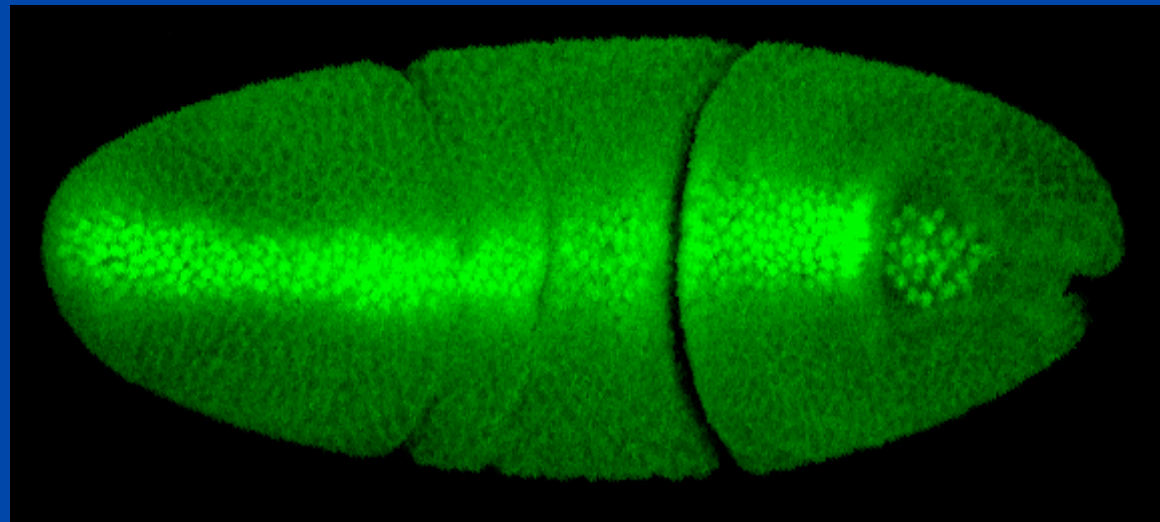


A dorsal midline stripe of strong nuclear Medea

End blastoderm
Side view
2.75hr



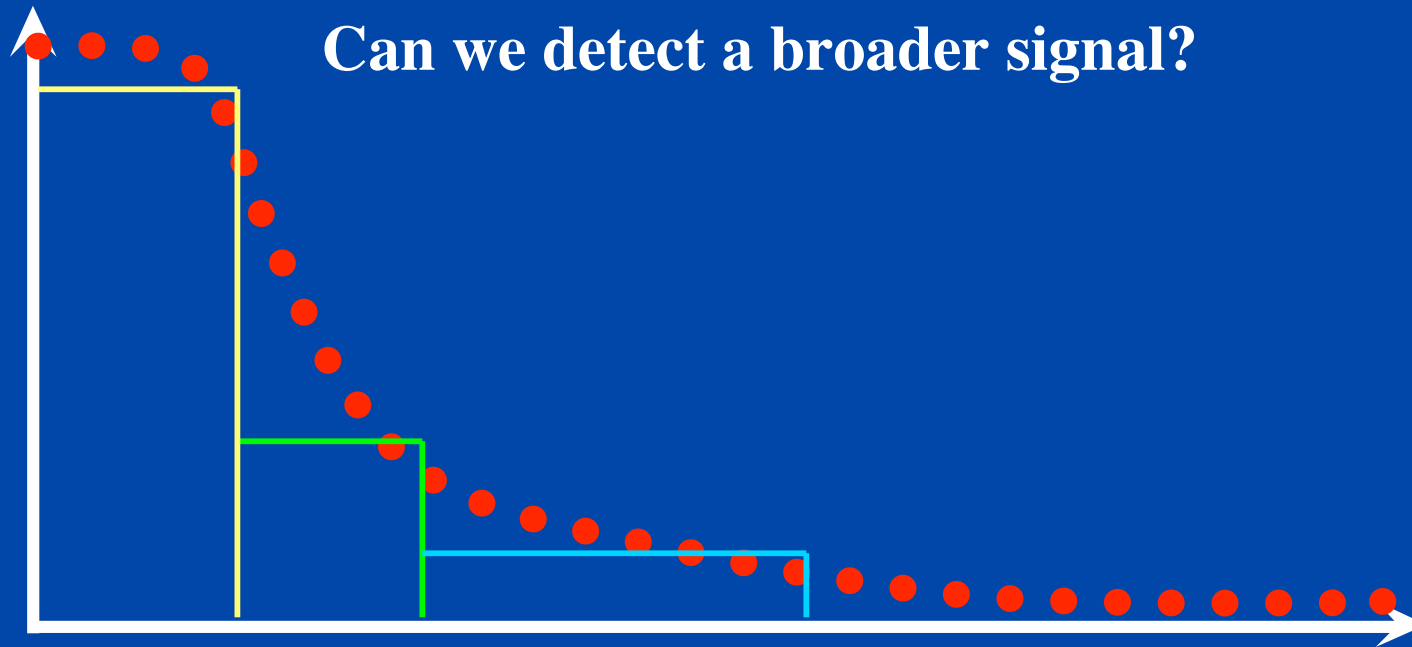
Gastrulation
Dorsal view
3.25hr



BUT...?

Stripe of intense response is only wide enough to account for amnioserosa induction...

Can we detect a broader signal?

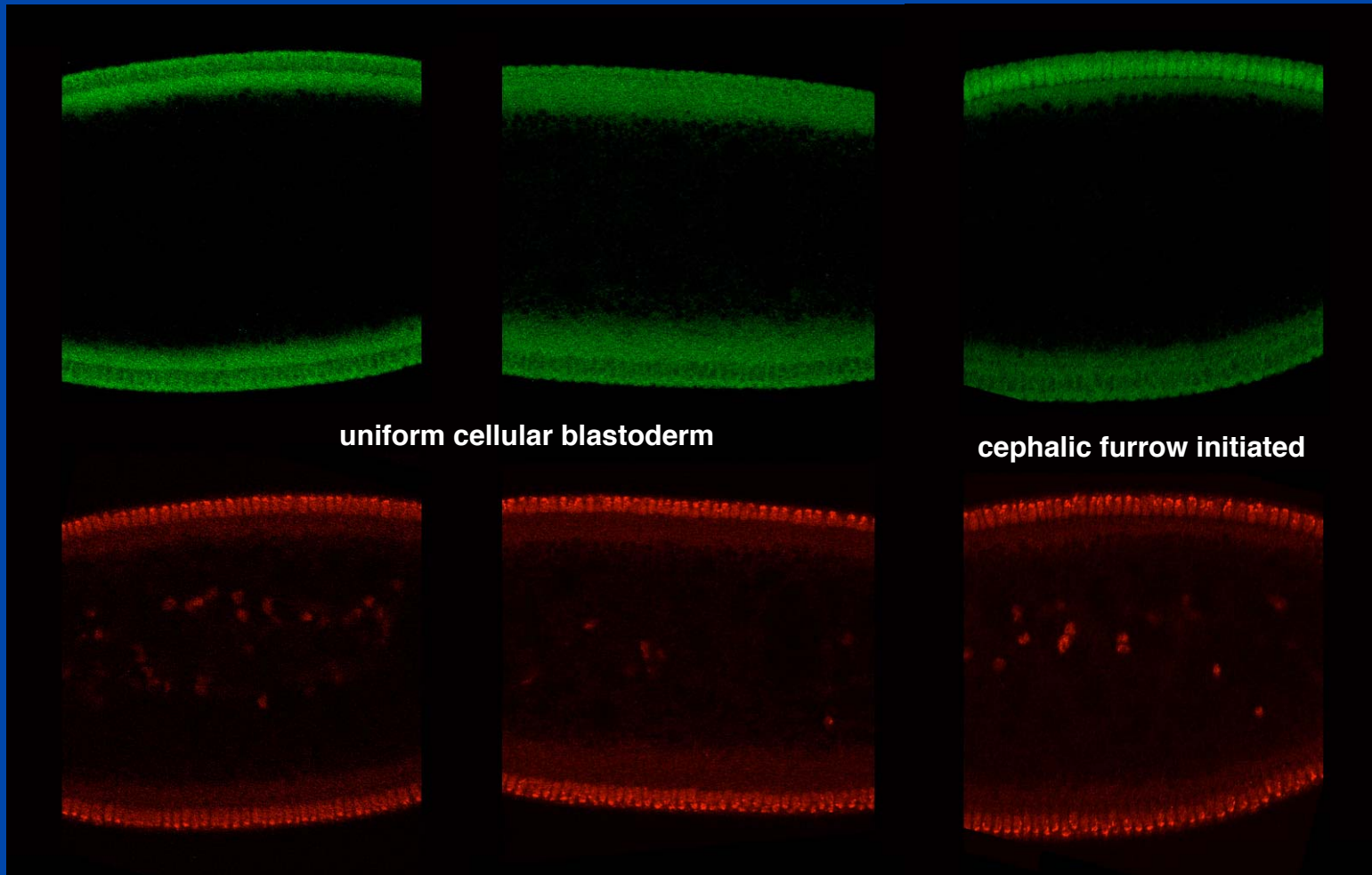


A low level signal in some blastoderm embryos

32 cells wide

3-9 cells wide

Medea

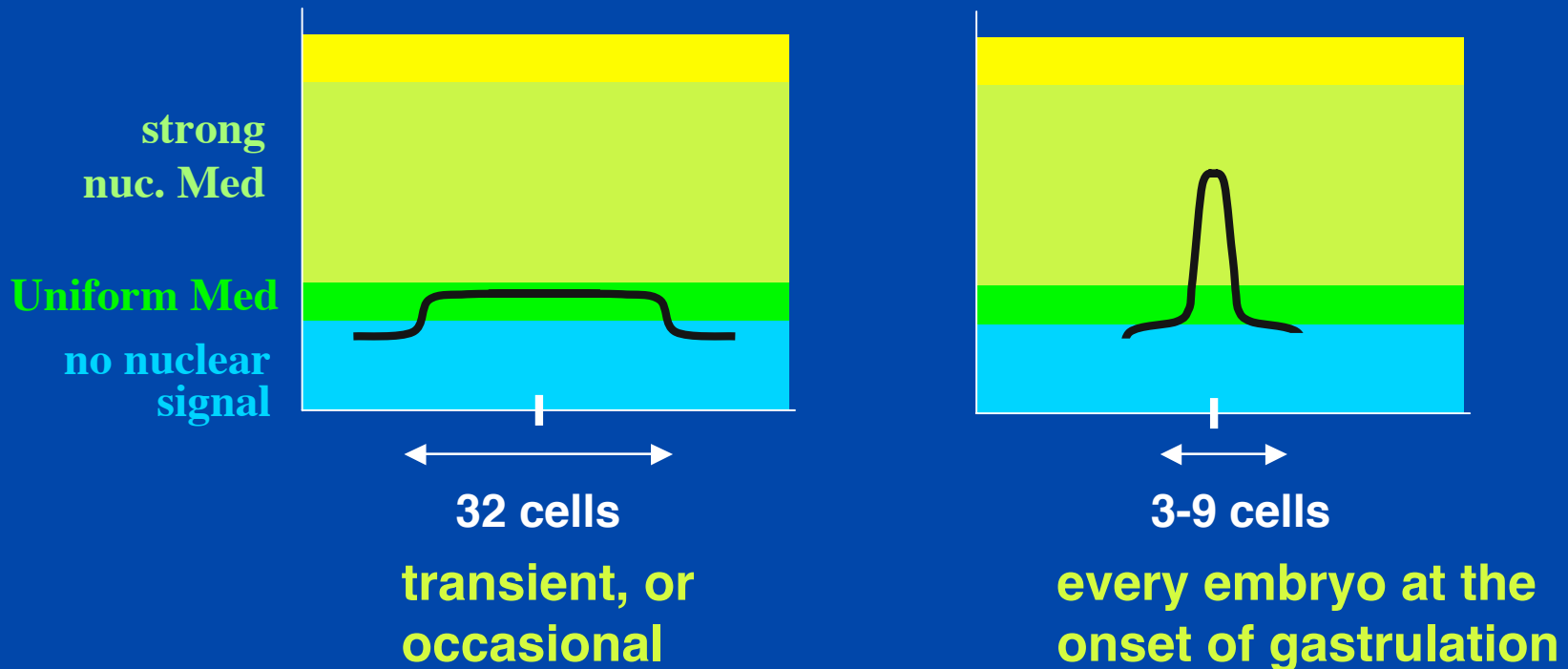


uniform cellular blastoderm

cephalic furrow initiated

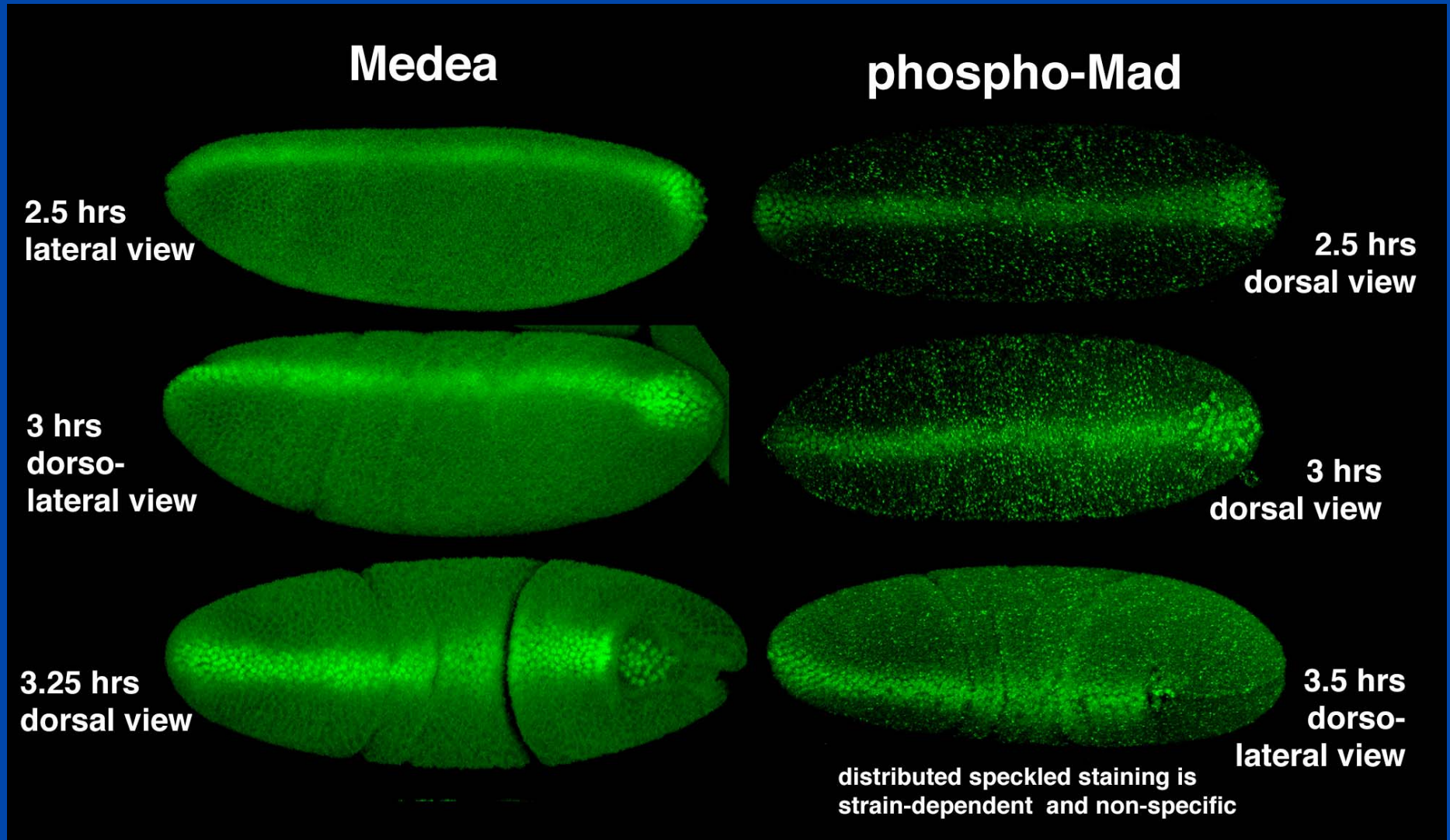
DNA

co-Smad response domain narrows as peak forms at the dorsal midline

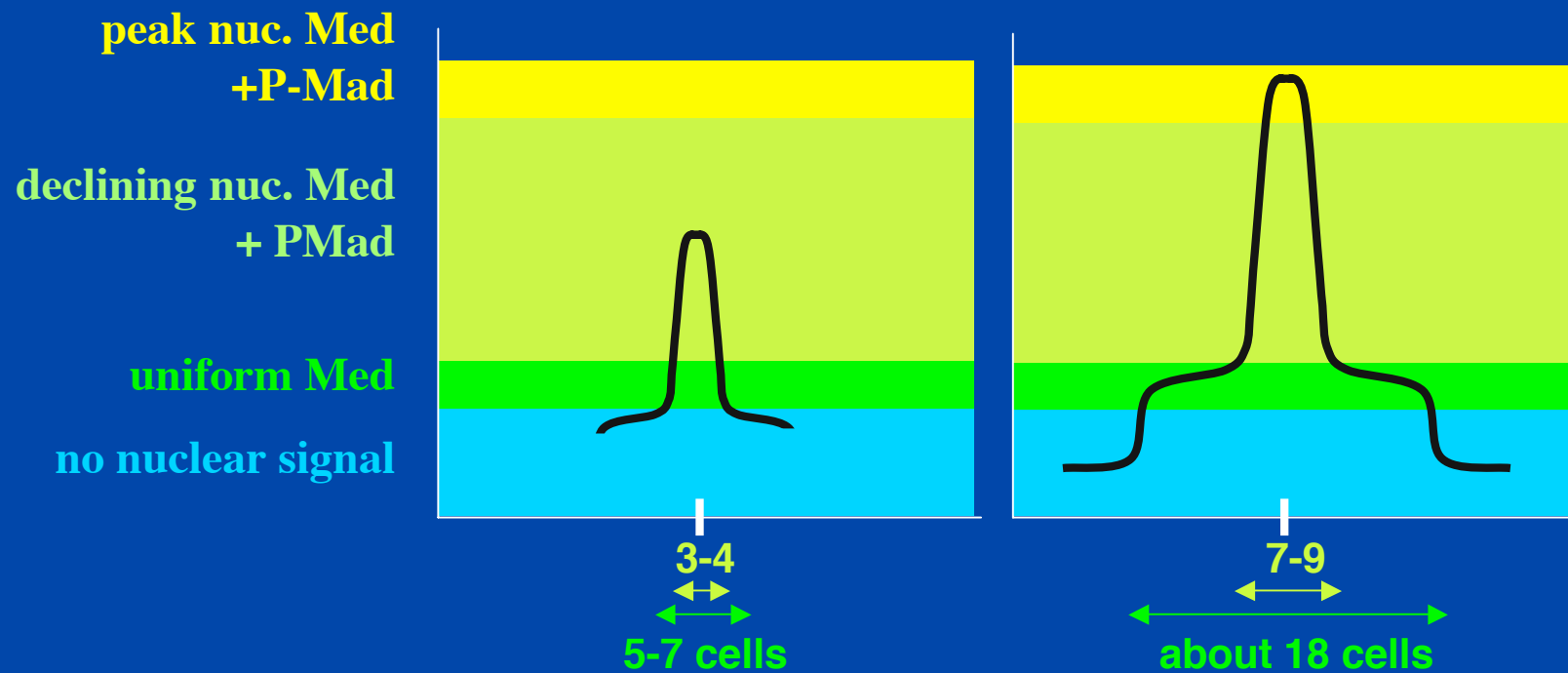


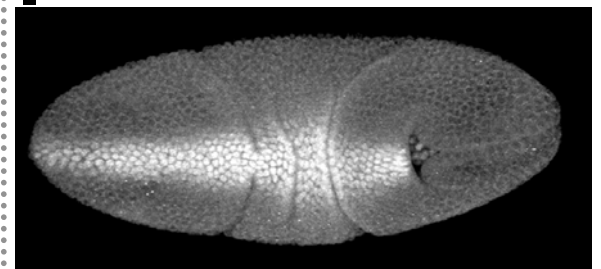
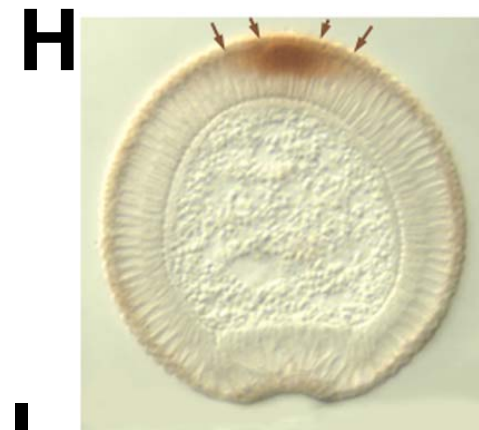
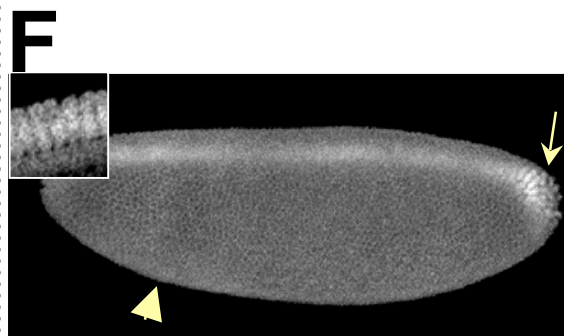
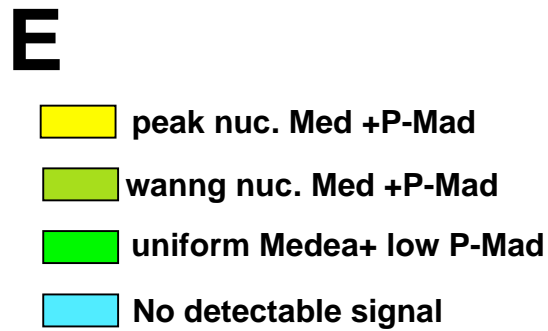
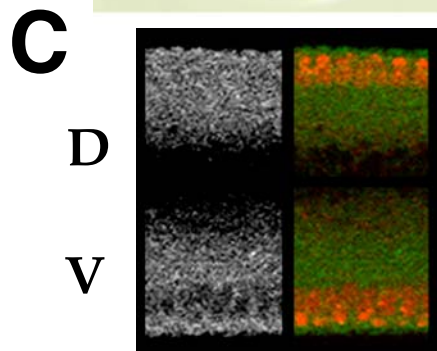
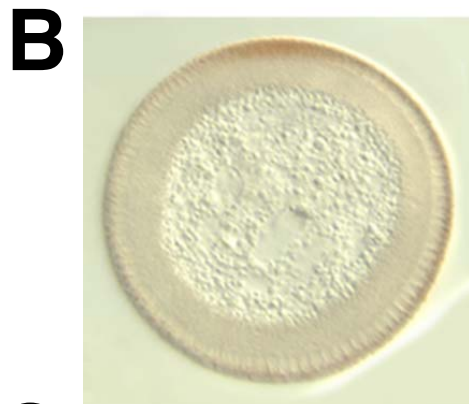
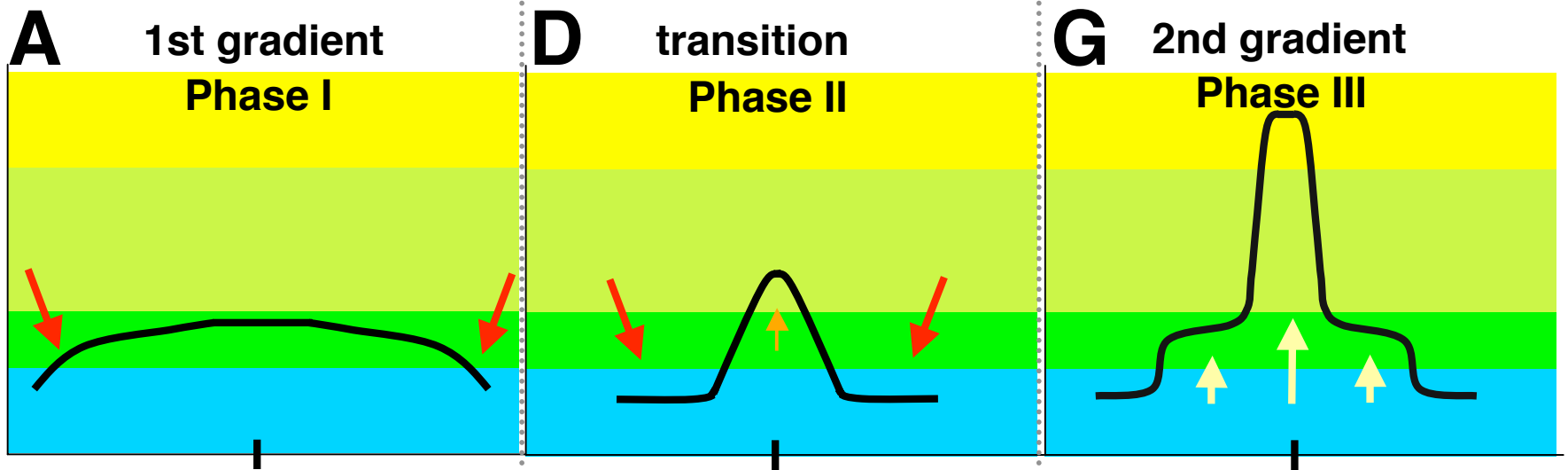
Ross et al, 2001 and Rushlow et al, 2001 have similar observations for PMad

Both co-Smad and phospho-Mad responses elevate during gastrulation



During gastrulation, peak co-Smad response is further elevated and more cells are involved





- Why *Drosophila* genetics?
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How is the BMP gradient shaped?

Mid-blastoderm

Wide band

weak signal

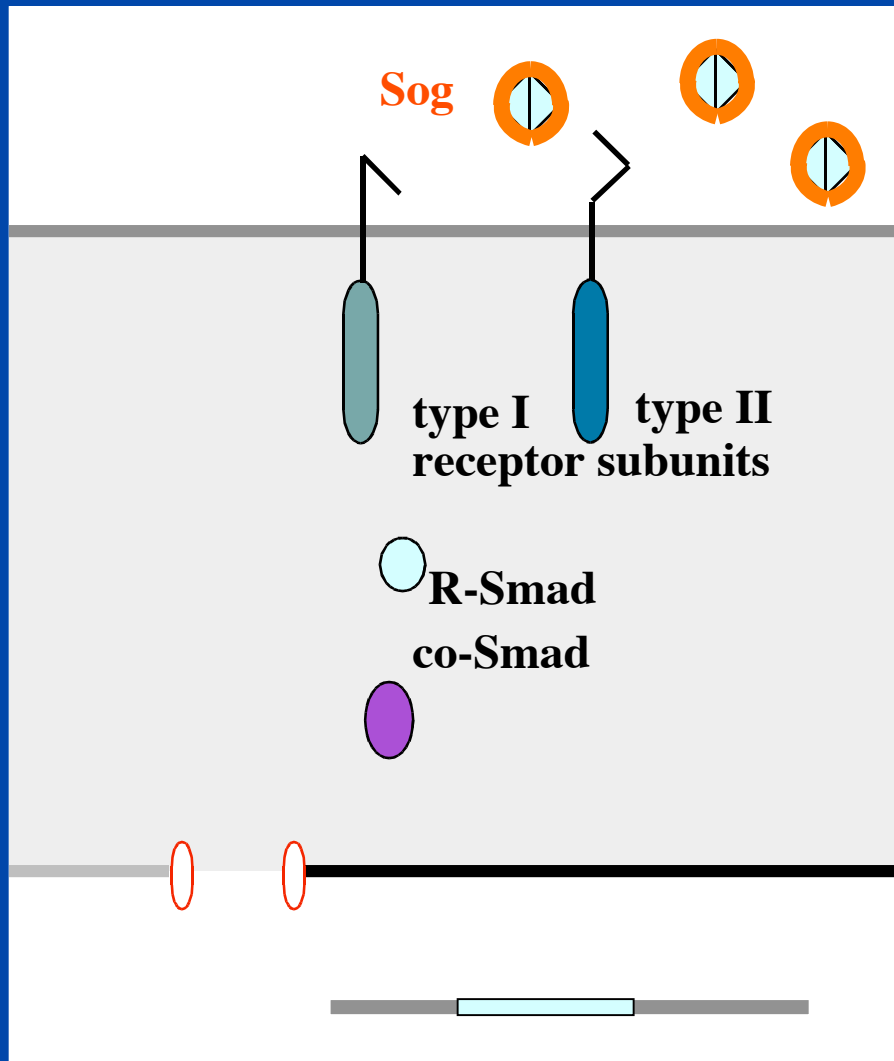


Gastrulation

Narrow stripe

strong signal

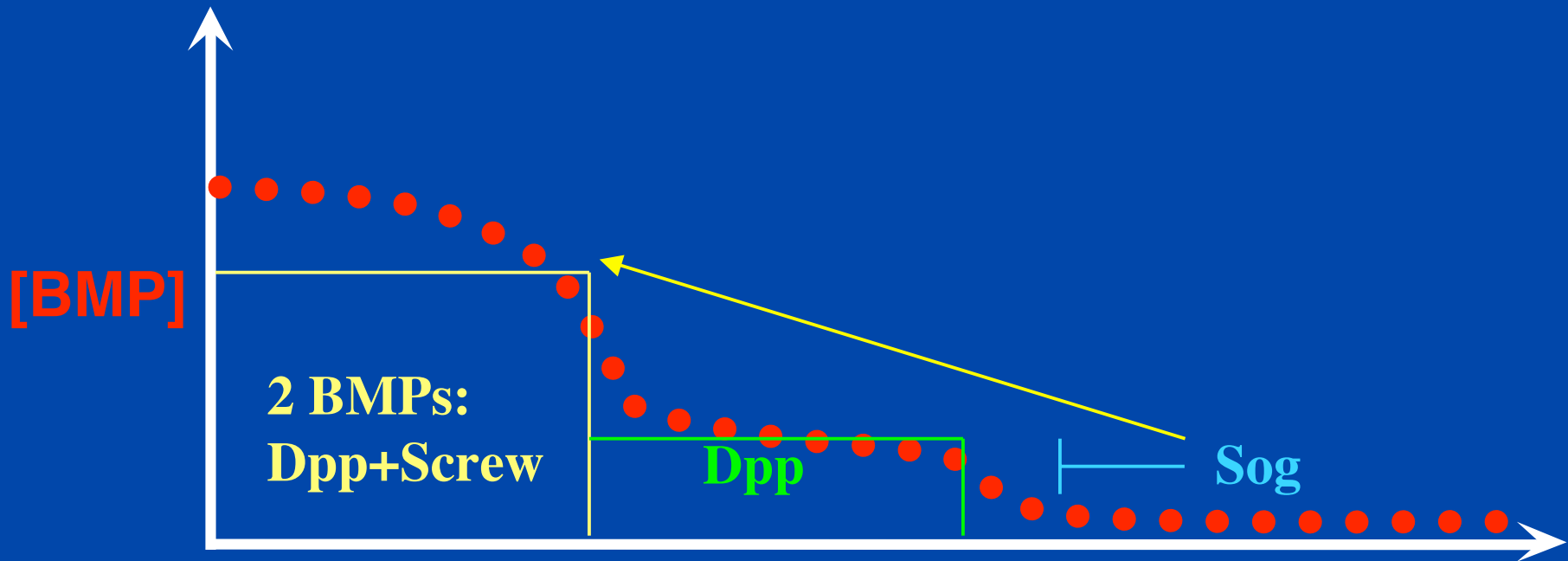
Sog is an extracellular BMP binding protein that can block signaling



Jazwinska et al, 1999,
Development 126, 3323

but, Sog is also required
to get a normal number of
amnioserosa cells

How does Sog affect the nuclear co-Smad signal?

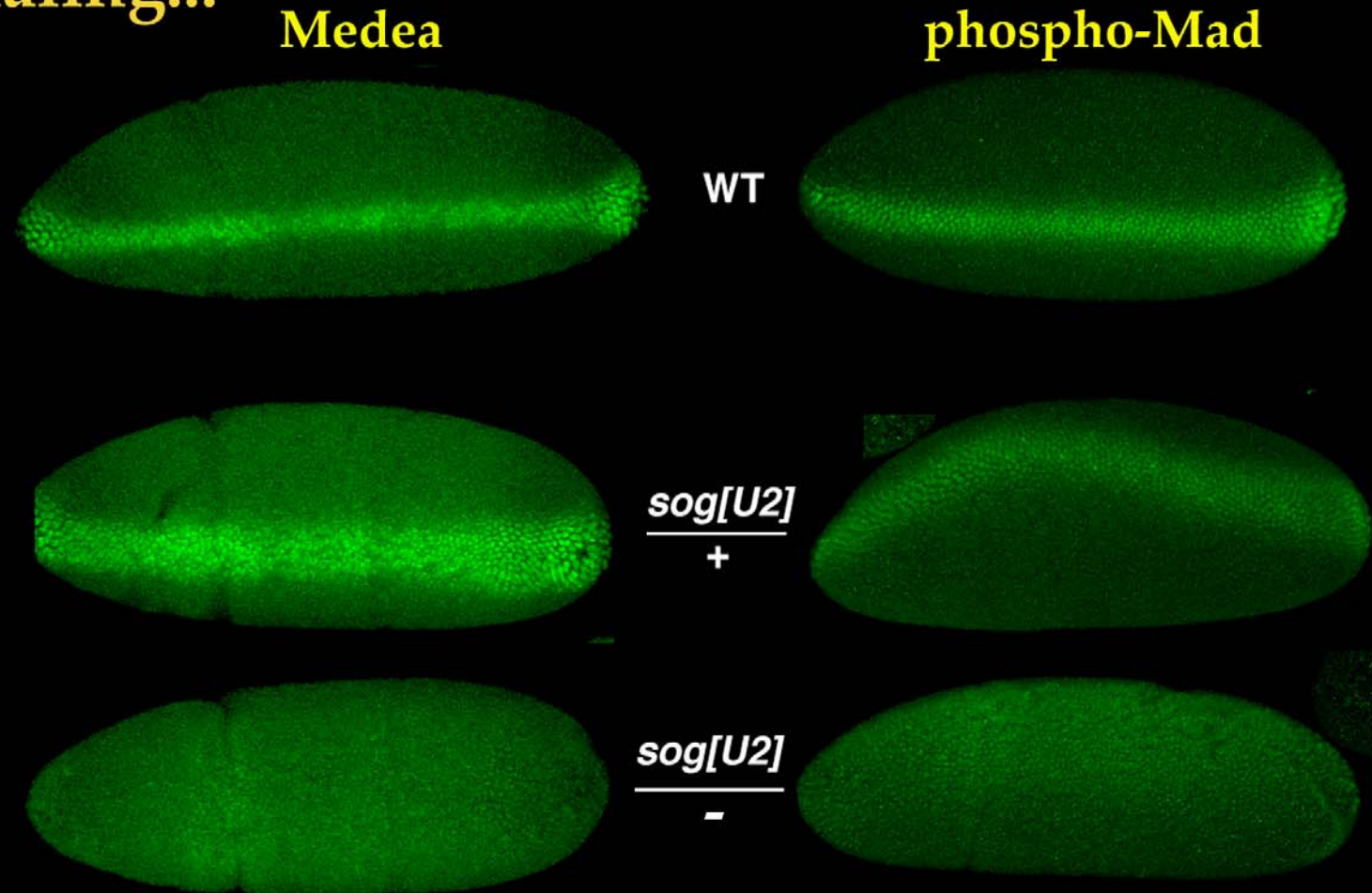


Amnioserosa

**Dorsal
Ectoderm**

**Ventral
Neurectoderm**

The BMP antagonist Sog limits the domain of signaling...



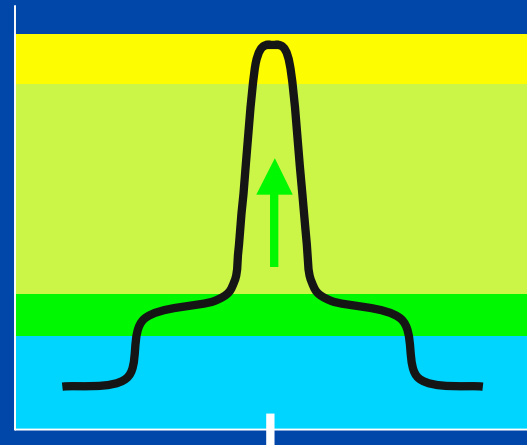
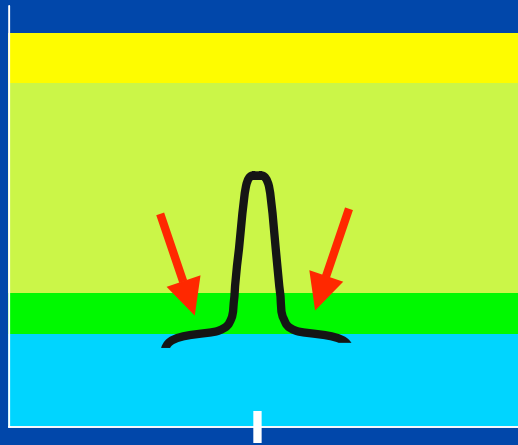
but Sog is also necessary for the stripe of peak signal.

Sog shapes the BMP activity gradient over time

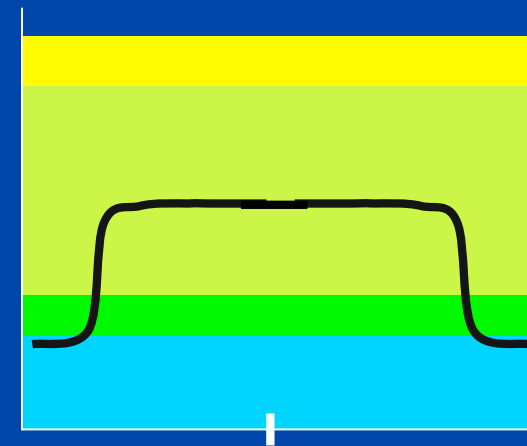
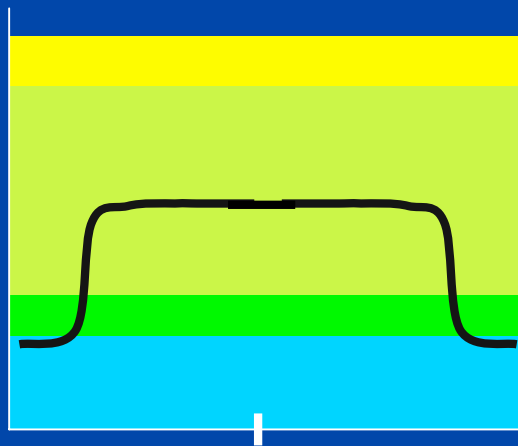
quenches

promotes

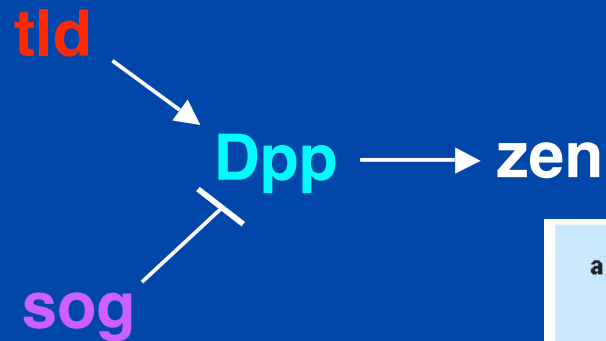
WT



sog⁻



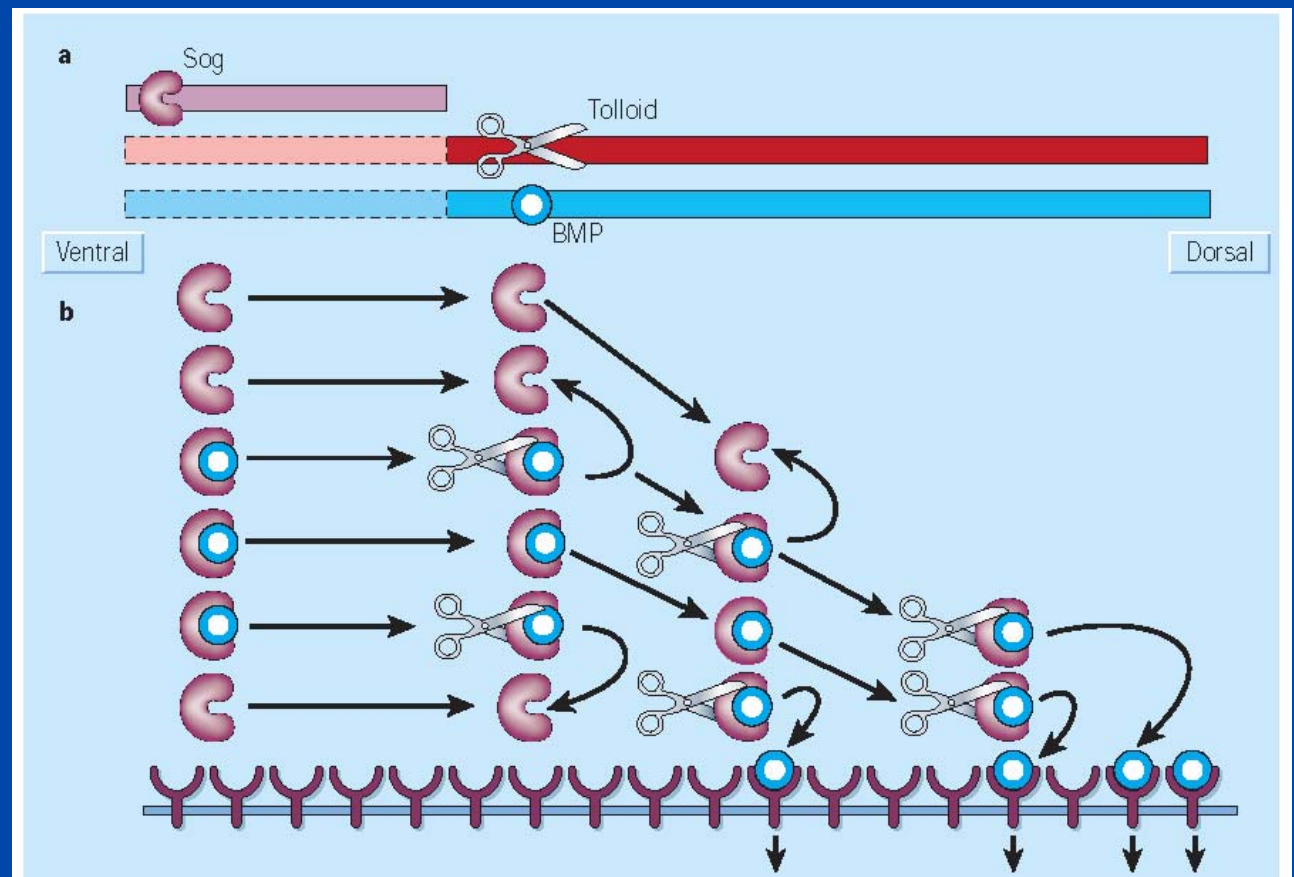
Modulation of BMP levels: 1992 to 2002



Tld=
Metalloprotease

Sog=
BMP binding protein

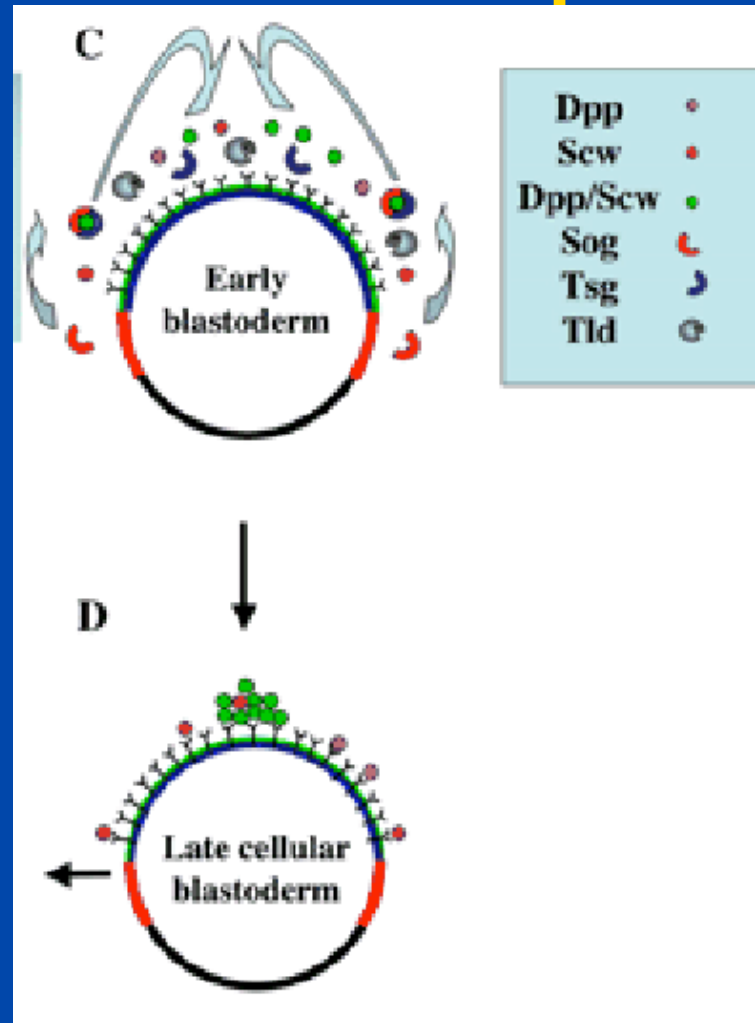
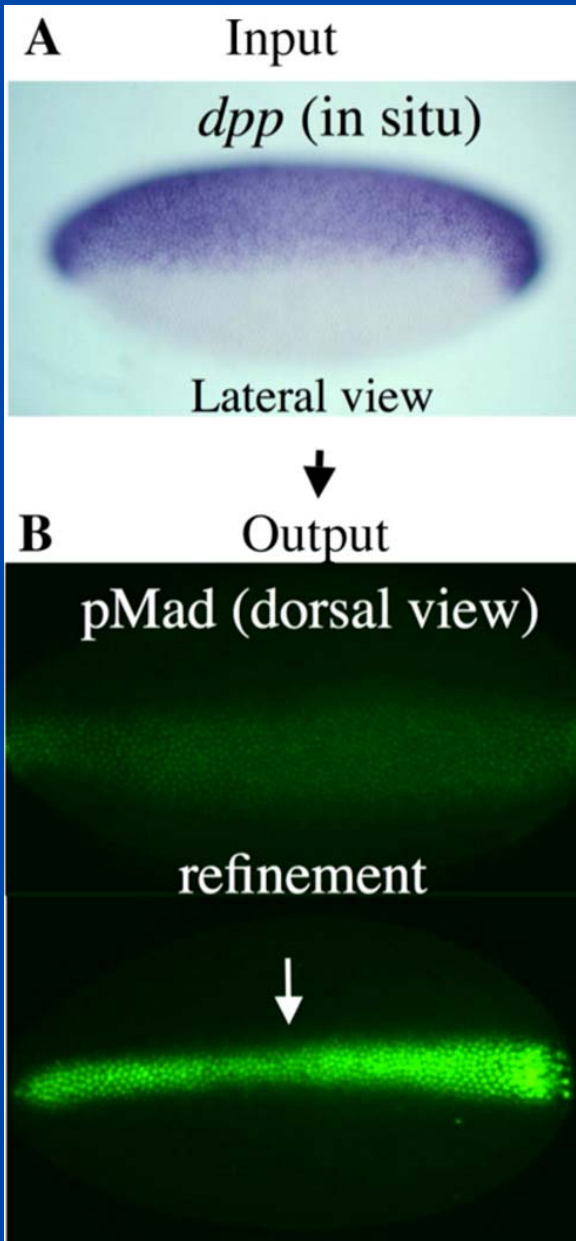
Zen=
transcription factor,
Partner to Smads



Reviewed by Meinhardt & Roth, 2002, 1999, Dev. Biol. 210, 251

But how does this work?

A BMP transport model:



O'Connor, M. B. et al. Development 2006;133:183-193



Development

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Changing BMP gradient can be described mathematically

Free ligand concentration :

$$\frac{\partial[L]}{\partial t} = D_L \frac{\partial^2[L]}{\partial x^2} - k_{on}[L](R_0 - [LR]) + k_{off}[LR] - j_{on}[L][ST] + (j_{off} + \tau)[LST] + V_L(x)$$

Ligand bound to receptor:

$$\frac{\partial[LR]}{\partial t} = k_{on}[L](R_0 - [LR]) - (k_{off} + k_{deg})[LR]$$

Sog concentration over time:

$$\frac{\partial[S]}{\partial t} = D_S \frac{\partial^2[S]}{\partial x^2} - n_{on}[S][T] + n_{off}[ST] + V_S(x)$$

Sog bound to Tld:

$$\frac{\partial[ST]}{\partial t} = D_{ST} \frac{\partial^2[ST]}{\partial x^2} + n_{on}[S][T] - n_{off}[ST] - j_{on}[L][ST] + j_{off}[LST]$$

Tld concentration over time:

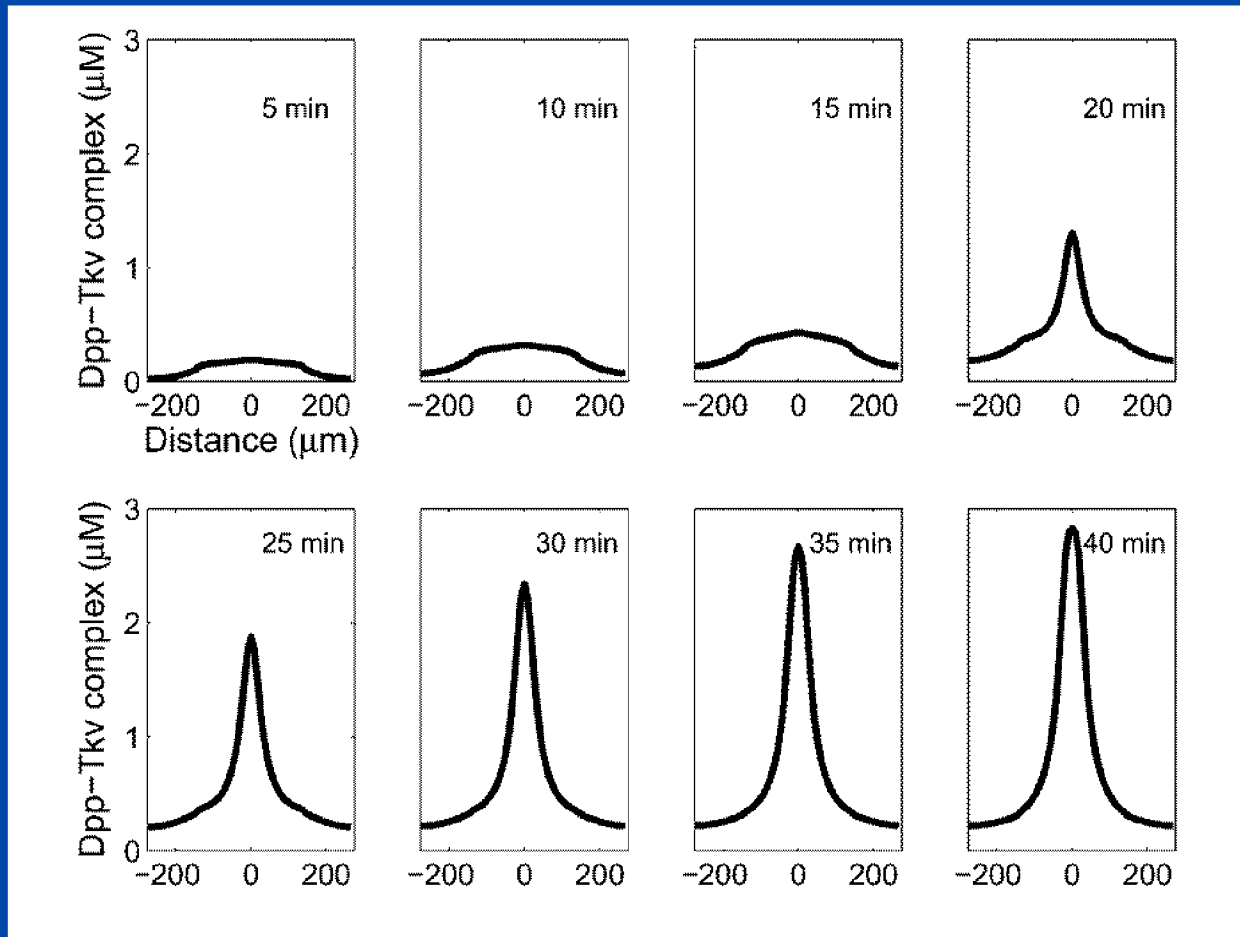
$$\frac{\partial[T]}{\partial t} = D_T \frac{\partial^2[T]}{\partial x^2} - n_{on}[S][T] + n_{off}[ST] + \tau[LST] + V_T(x)$$

Ligand bound to Sog and Tld:

$$\frac{\partial[LST]}{\partial t} = D_{LST} \frac{\partial^2[LST]}{\partial x^2} + j_{on}[L][ST] - (j_{off} + \tau)[LST]$$

Mizutani et al, 2005, Dev Cell 8:915-924

Computational modeling of the changing pattern of responses...



Can be used to predict important parameters

Computational modeling to test the importance of molecular mechanisms...

- **Degradation of ligand-receptor complexes**

Mizutani et al, 2005, Dev Cell 8:915-924

- **Positive feedback to stabilize narrow, high level gradient**

O'Connor et al., 2006, Development 133:183-193

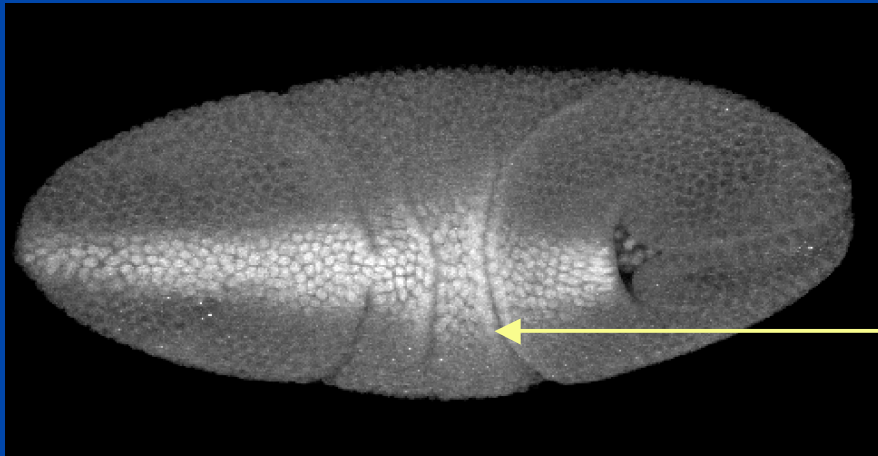
- **Contribution of Screw ligand: heterodimers versus two homodimers**

Shimmi et al. 2005, Cell 120: 873-886

- **Pathway genetics**
- **Prediction of a morphogen gradient**
- **Identification of signaling pathway from genetics**
- **Visualization of a gradient from signal transducers**
- **Computational predictions of molecular mechanisms**

We still have questions...

How is a threshold response organized to make a tissue boundary?



amnioserosa/
dorsal ectoderm boundary

