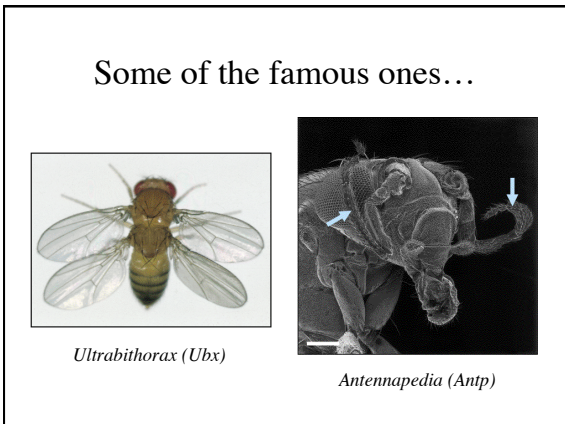
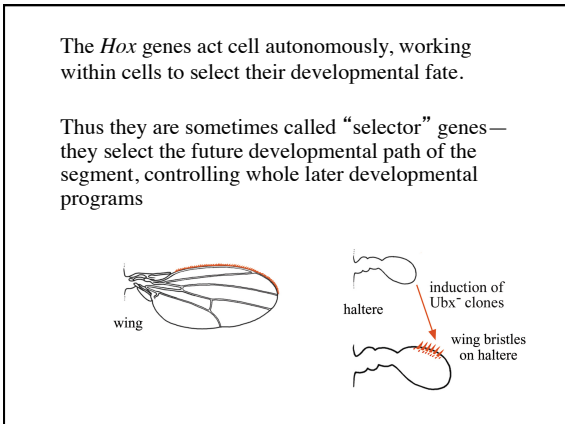


Homeotic mutation (Bateson, 1894):
A mutation in which one normal body part is replaced with another



The *Hox* genes specify the identity of a field (segment), not the formation of the field.

Contrast this to the segmentation genes, which establish the segments themselves.



The *Hox* genes are all *transcription factors* and are related in their DNA binding domains, the *homeobox*

```

lab      NNSGRITNFNKQLTELEKEPHFNRYLTRARRIEIANTLQINETOVKIWFQNRMRKQKKRV
pb       FRRIRTAITNTQLLEKEEHPFNKYLGRPRRIEIASLDLTERQWKIWFQNRMRKHKBQT
Dfd      FRRQRTATRHQLLEKEEHPFNRYLTRRRRIEIAHLVLSEKQIKIWFQNRMRKWKBN
Scr      TRQRTYTRYQTLLEKEEHPFNRYLTRRRRIEIAHALCLTERQIKIWFQNRMRKWKKH
Antp     RRRGRQTYTRYQTLLEKEEHPFNRYLTRRRRIEIAHALCLTERQIKIWFQNRMRKWKKN
Ubx      RRRGRQTYTRYQTLLEKEEHPFNRYLTRRRRIEIAHALCLTERQIKIWFQNRMRKWKKEI
abd-A    RRRGRQTYTRYQTLLEKEEHPFNRYLTRRRRIEIAHALCLTERQIKIWFQNRMRKWKKEI
abd-B    VKKKRKPYSKQTLLEKEEFLPNAVYSKOKRWELARNLQITERQWKIWFQNRMRKWKKNS
    
```

In the fly, there are two main clusters of *Hox* genes: the Antennapedia Complex (ANT-C) and the Bithorax Complex (BX-C). Strikingly, their order on the chromosome is identical to their order of expression in the A/P axis.

In a *Ubx* mutant, *Antp* expands into T3, causing a homeotic mutation of T3 into T2 (which has wings). More posterior segments are unaffected due to repression of *Antp* by *Abd-A*.

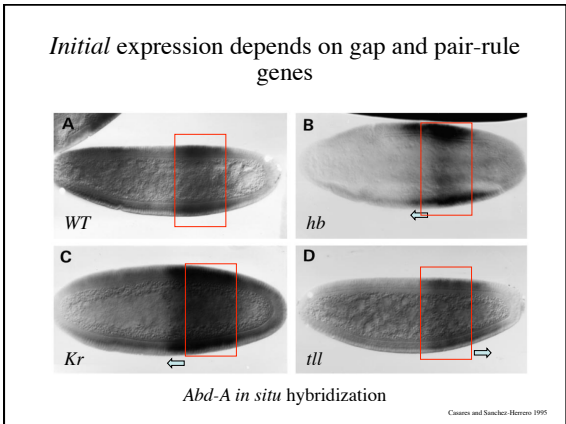
Although there are only 3 genes in the BX-C, there is a complex regulatory structure that causes segment-specific phenotypes in various mutants.

For example, the *bxd/pbx* mutation causes a transformation of A1-> T3 and a partial transformation of more posterior segments.

The *Hox* genes control the expression of important segment-specific and organ-specific genes, usually other transcription factors, which in turn control whole developmental programs.

For example, *Scr* **activates** the *forkhead* tx factor, which controls development of the salivary gland.

Ubx and *Abd-A* **repress** *distal-less* in abdominal segments, preventing the development of appendages. This also requires segmentation gene input.



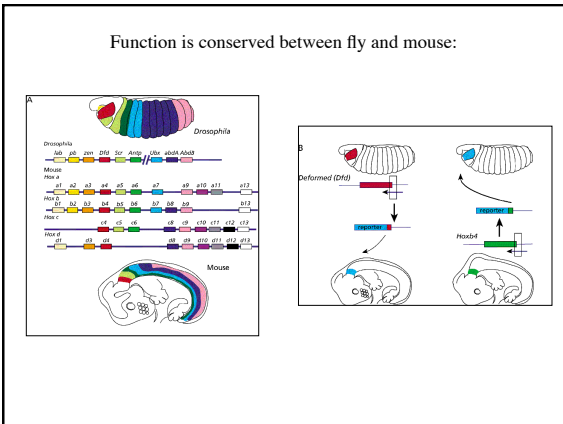
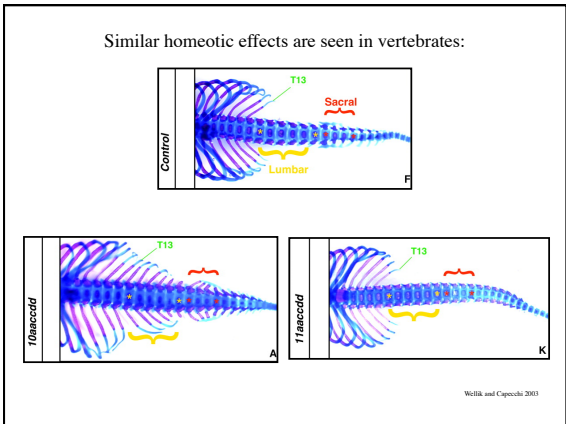
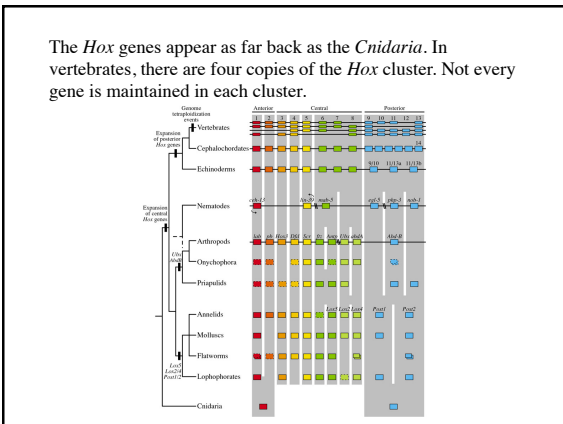
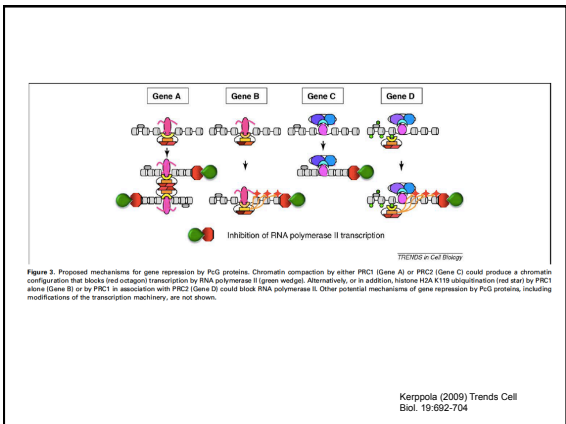
Maintenance of expression is via epigenetic mechanisms.

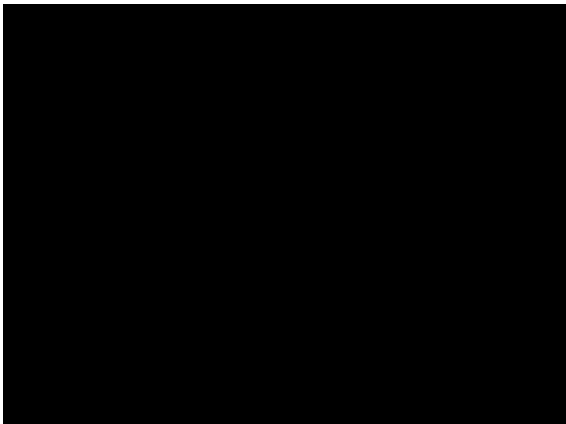
Polycomb group (Pc-G) genes keep homeotic genes repressed in segments where they were not initially activated.

Trithorax group (trx-G) genes maintain the expression of the homeotic genes in their appropriate segments.

Both groups act by modifying chromatin conformation. PcG complexes promote H3K27 tri-methylation. TrxG complexes can promote H3K4 methylation. Many details remain to be established.

In this way, patterning information conveyed by the early activity of the segmentation genes is preserved throughout the life cycle.





Strategies for Patterning

- ubiquitous receptor, localized activation of ligand (dorsal/ventral system, terminal system)
- localized RNA (anterior/posterior)
- translational inhibition (posterior)
- morphogen gradients (d/v, a/p)
- mutual repression (segmentation)
- cell-cell signaling (segment polarity)
- chromatin remodeling (Hox genes)

Maternal localization of *sqf* RNA in zebrafish presages the dorsal axis

Gore et al. 2005

The developing vertebrate limb

- Interactions between Wg (Wnt) and HH (Shh), like we saw in the fly embryo

Some major players in human disease

- Wg: Wnt/ β -catenin/APC (oncogenes)
- Dpp: TGF- β (oncogene)
- Dorsal: $\text{rel/NF}\kappa\text{B/TLR/IL-1}$ (immune response)
- Ras (oncogene)
- HH/*ptc/smo/gli* (cancer, birth defects)
- Runt (hematopoiesis, leukemias)
- Hox cluster (various birth defects)

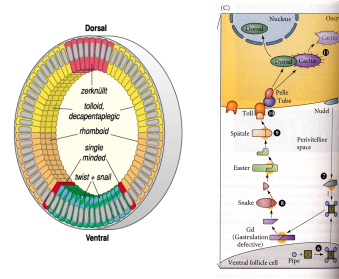
Locus	Map position*	No. of alleles†
<i>cubitus interruptus</i> ² (<i>ci</i> ²)	4-0	(2)
<i>wingless</i> (<i>wg</i>)	2-30	6
<i>sonoshears</i> (<i>shh</i>)	2-104	1
<i>hedgehog</i> (<i>hh</i>)	3-90	2
<i>fused</i> (<i>fu</i>) ²	1-59.5	(9)
<i>patch</i> (<i>pat</i>)	2-55	8

Sonic Hedgehog

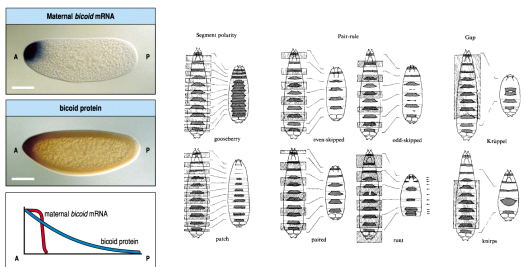
- Holoprosencephaly
- Smith-Lemli-Opitz
- Gorlin Syndrome
- Basal Cell Carcinoma
- Medulloblastoma
- Trichoeptiloma
- Esophageal SCC
- Bladder Transitional CCA
- Basal Cell Carcinoma
- Medulloblastoma
- Basal Cell Carcinoma
- Medulloblastoma
- Basal Cell Carcinoma
- Glioblastoma
- Rhabdomyosarcoma
- Osteosarcoma
- Predicts Sarcoma Grade
- Greig Syndrome
- Pallister-Hall Syndrome
- Postaxial Polydactyly A
- Preaxial Polydactyly IV
- Postaxial Polydactyly A/B
- Rubinstein-Taybi
- Saethre-Chotzen

We've now seen:

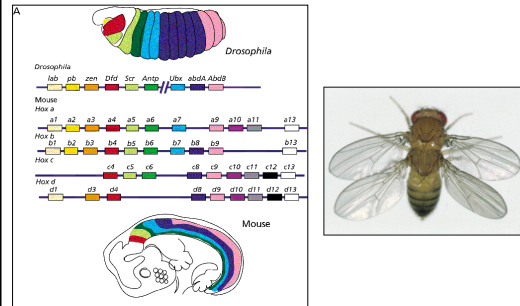
Dorsal-Ventral Patterning:



Anterior-Posterior patterning and Segmentation:

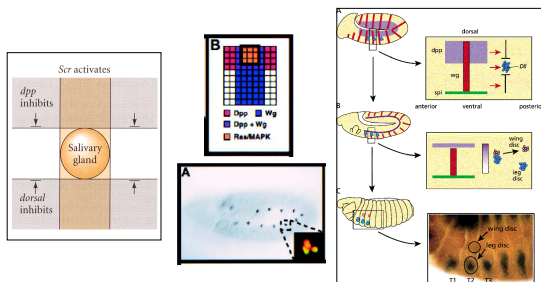


Segment Identity specification:

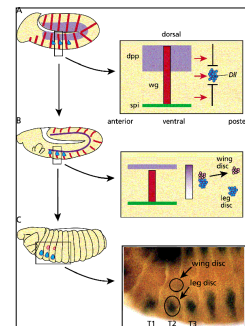


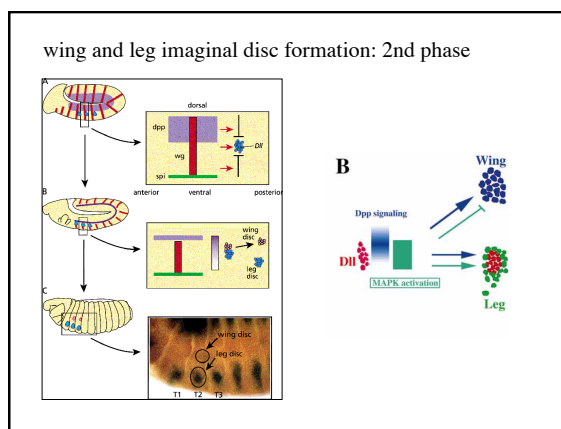
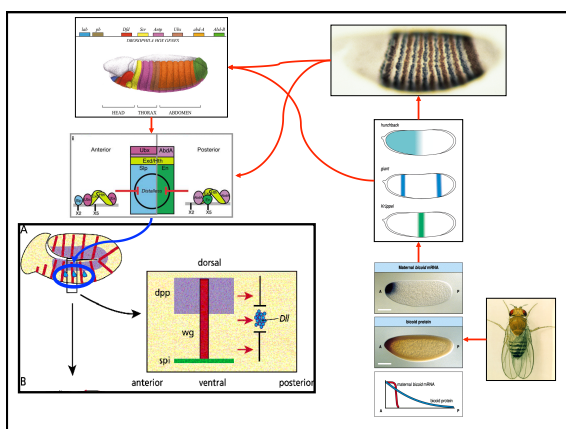
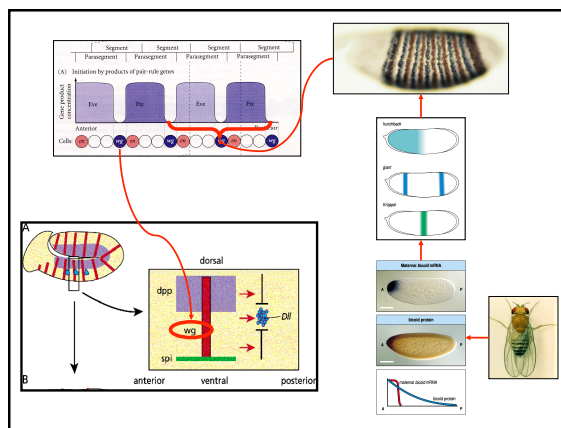
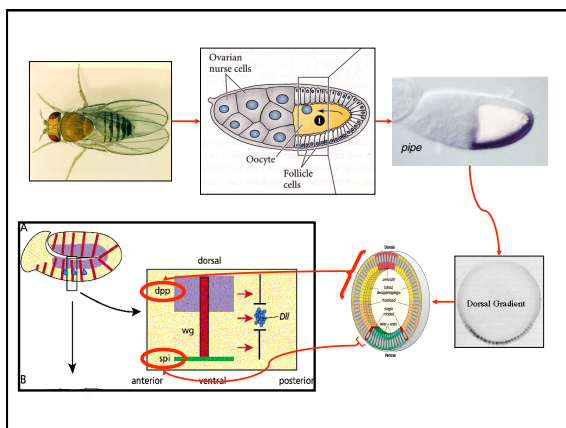
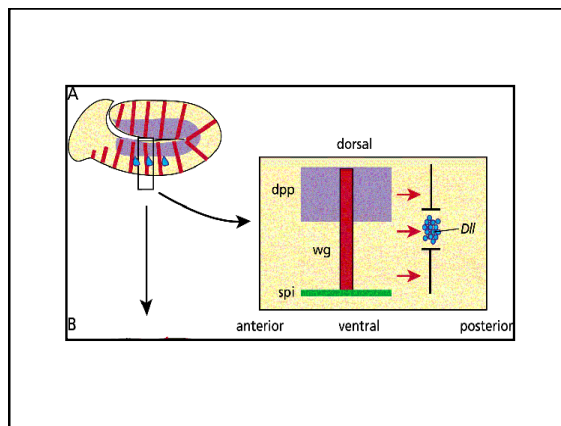
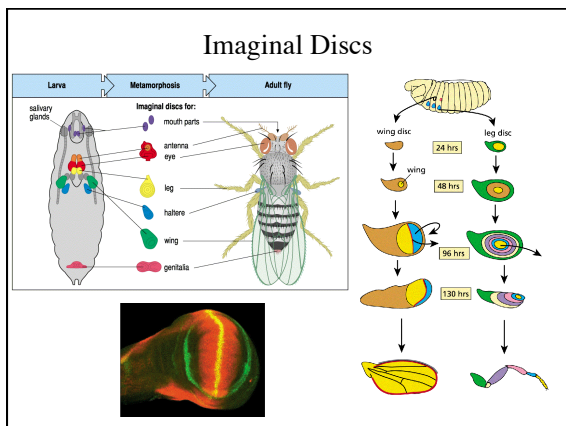
What does this mean for organogenesis?

- These early axial patterning and segmentation events set up a coordinate grid on which organs are built

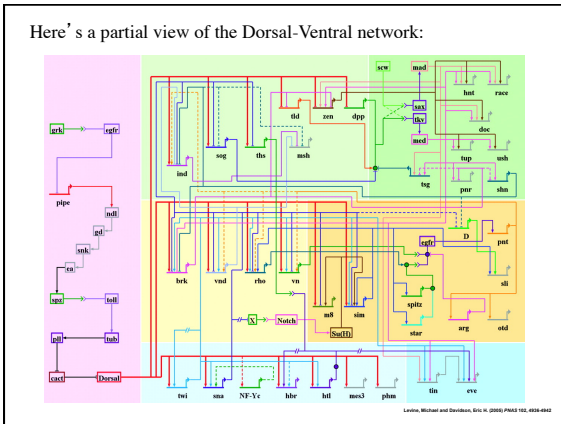


Let's look at wing and leg imaginal disc formation:





Here's a partial view of the Dorsal-Ventral network:



Musical Interlude

Regulatin' Genes

(http://www.youtube.com/watch?v=9k_oKK4Teco&feature=youtu.be)