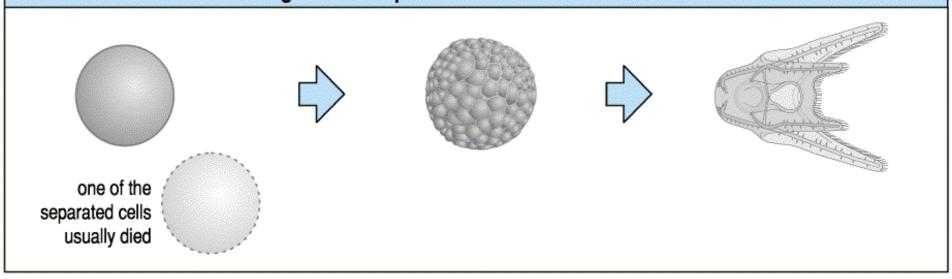
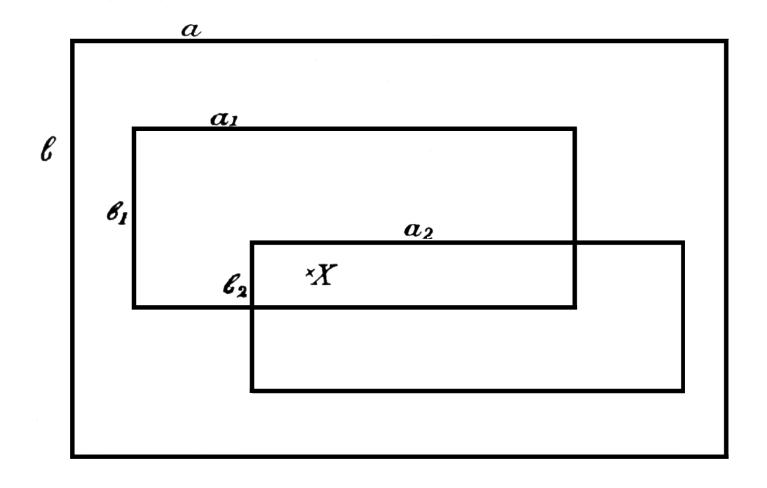
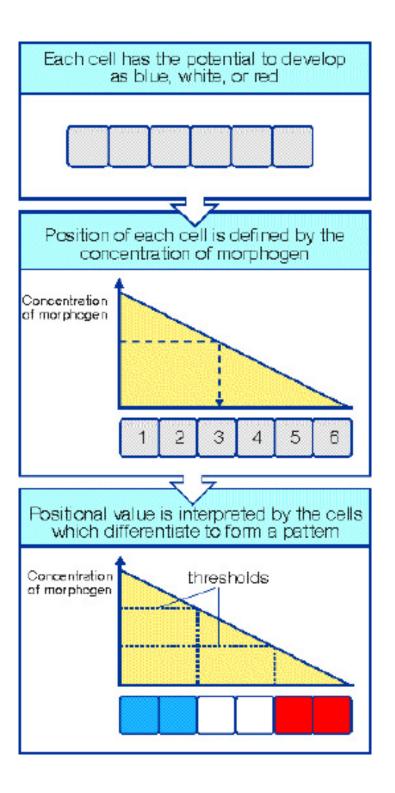
Normal development of sea urchin larva from two-cell stage

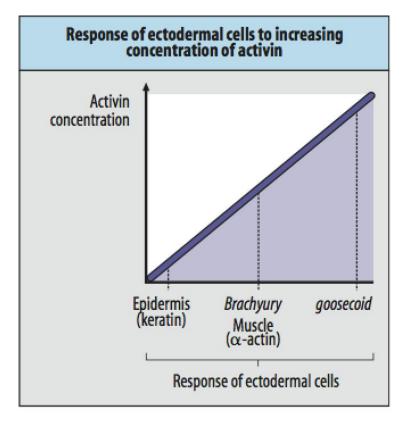
Driesch's separation of cells at two-cell stage resulted in the death of one cell. The surviving cell developed into a small but otherwise normal larva

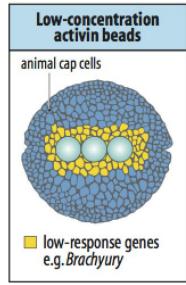


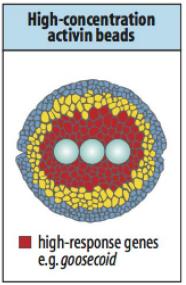
Driesch's entelechy

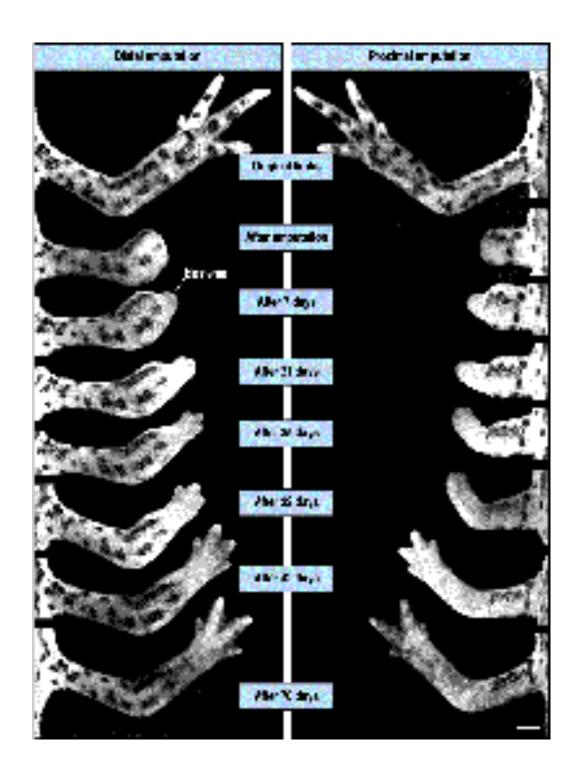




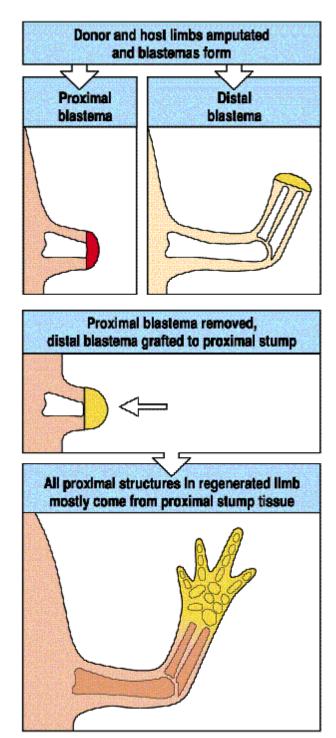








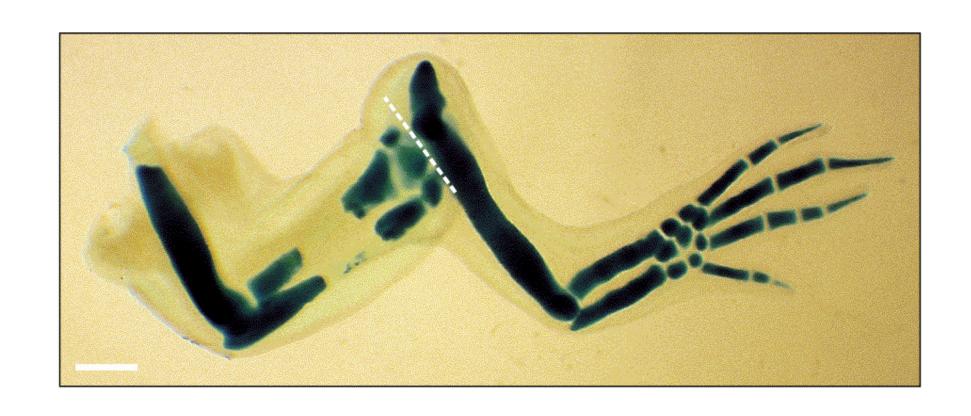
What is the molecular basis of positional values?



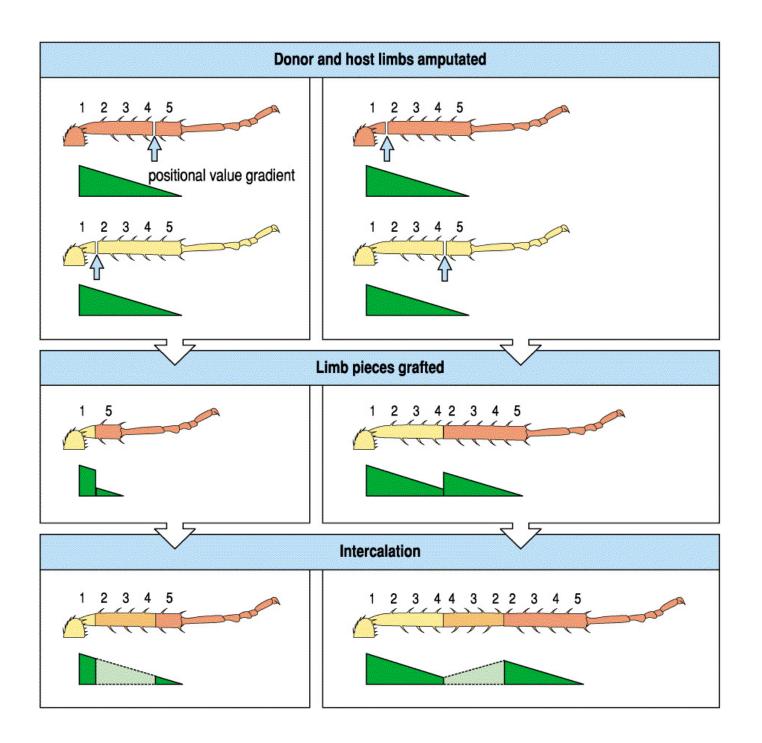
Prod 1 has a 5-fold difference along the

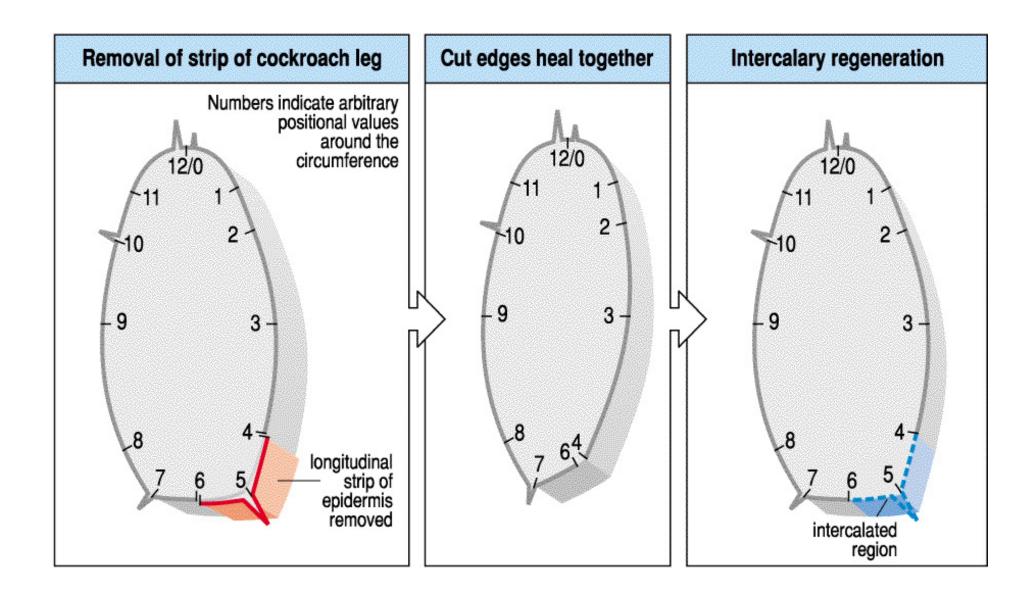
Newt limb - this is equivalent to about

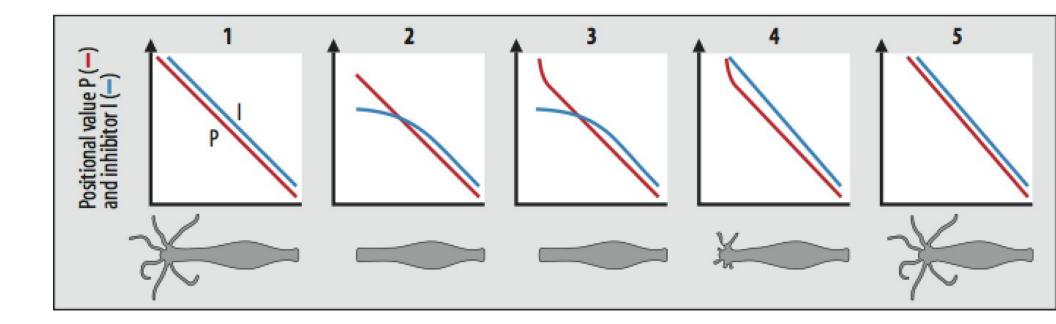
a 2% difference over each cell

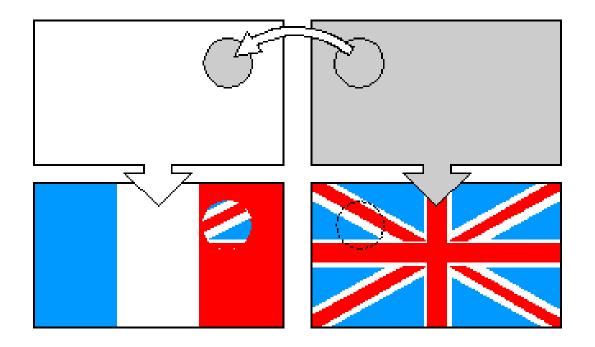


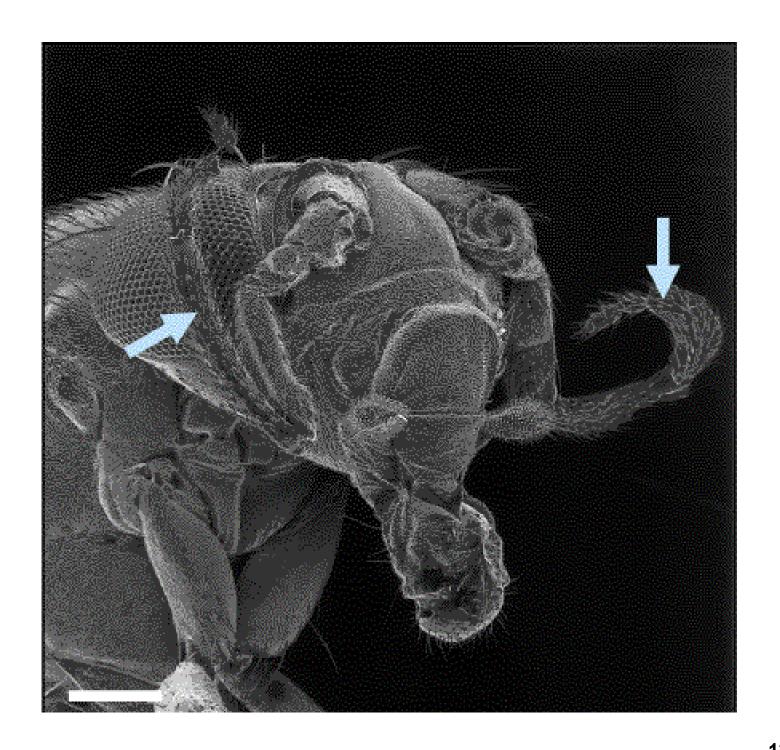
Retinoic acid alters positional values continuously





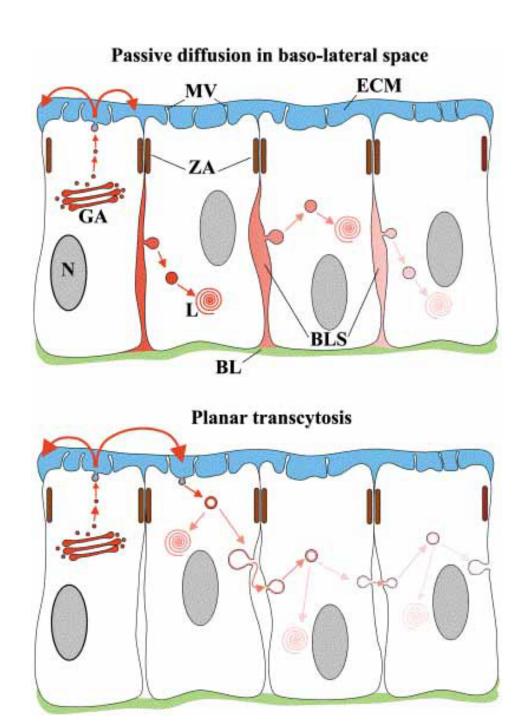


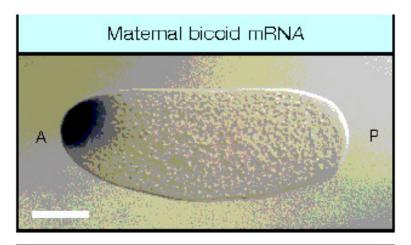


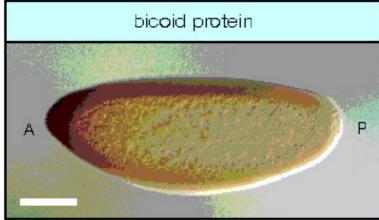


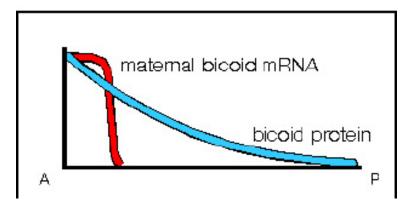
How can positional information be specified? Most models are based on diffusion of a morphogen which seems unreliable. Some models for the fly involve endocytosis and others for the limb, time.

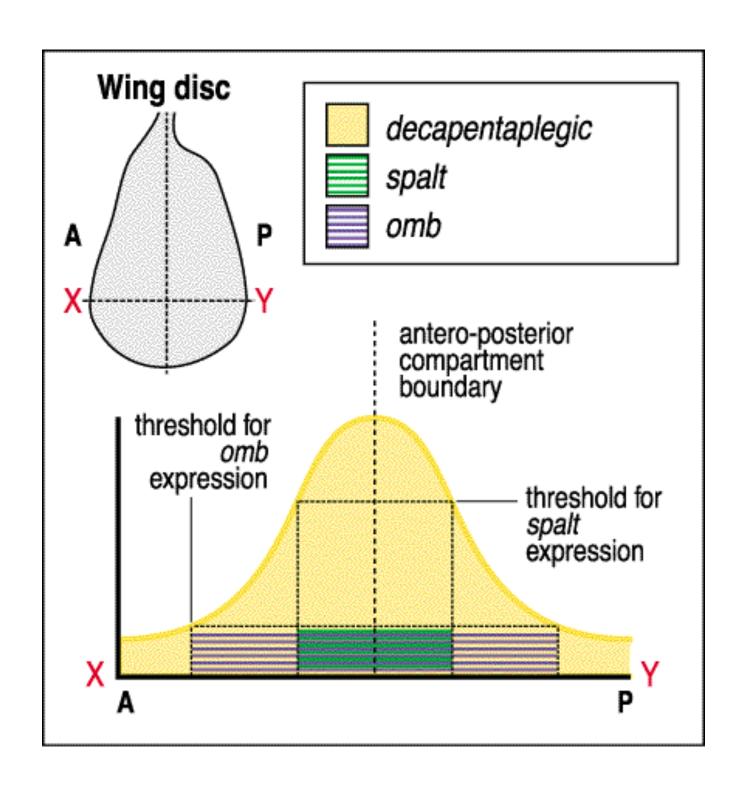
There is no good quantitative evidence for any of the many gradients that have been described.eg.Dpp,Chordin, BMP, Hedgehog







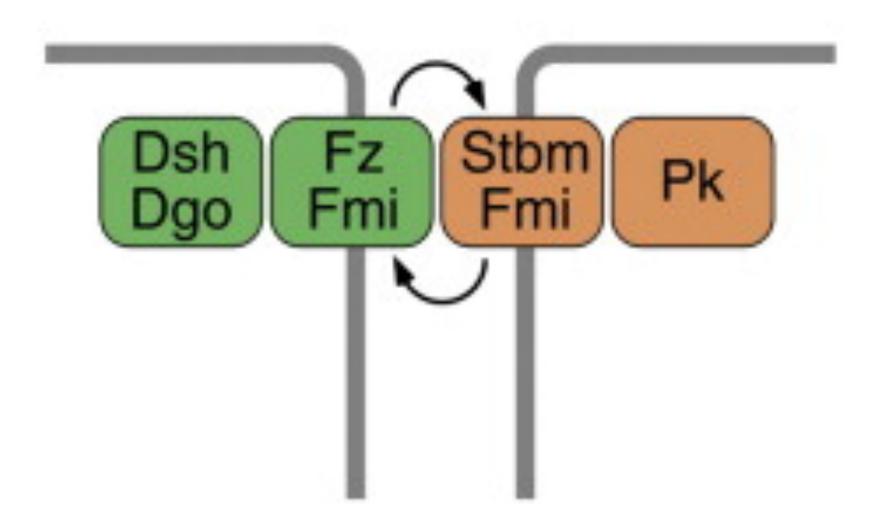


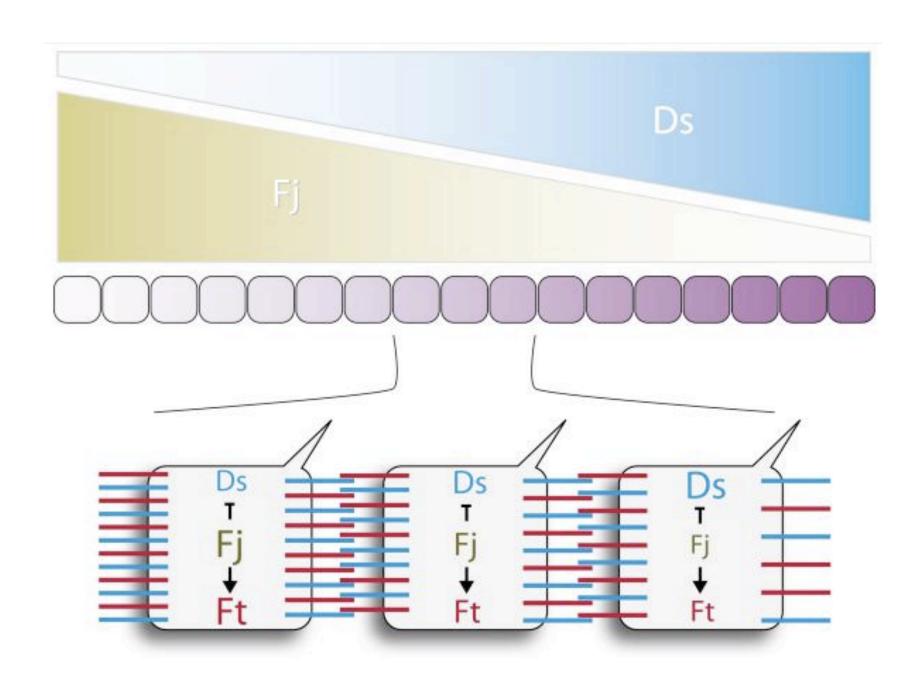


Dynamin-dependent endocytosis was required for spreading of Dpp, but not Wg.

White et.al

RA promotes and Fgf suppresses RA degradation, thereby linking the shapes of RA and Fgf gradients in zebrafish. Computational models suggest that this linkage helps make RA-mediated patterning robust.

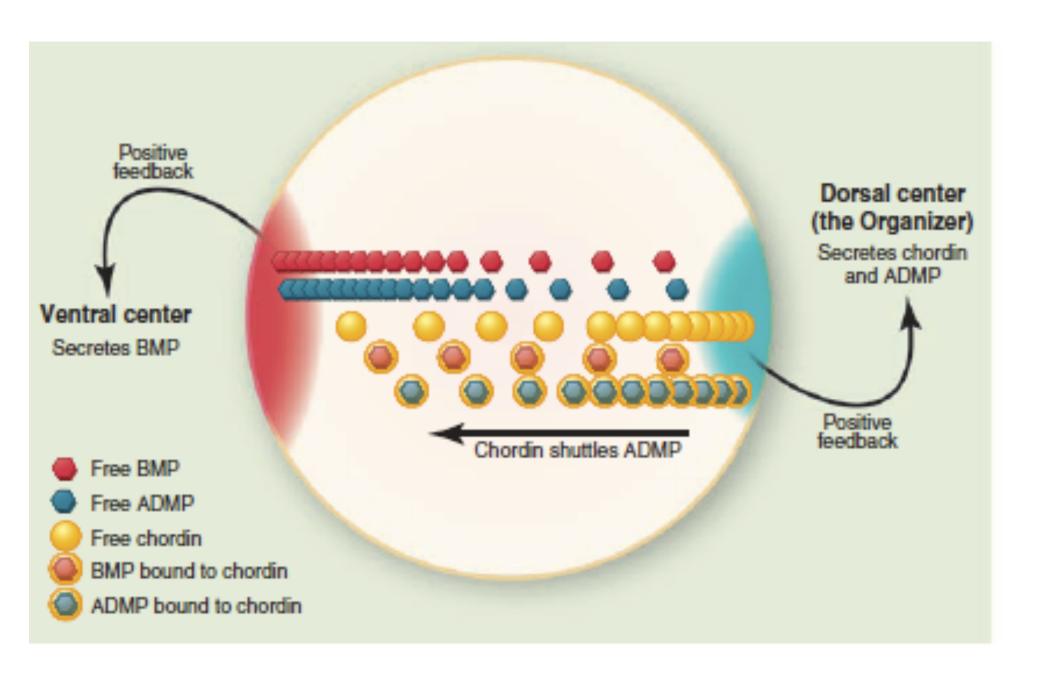


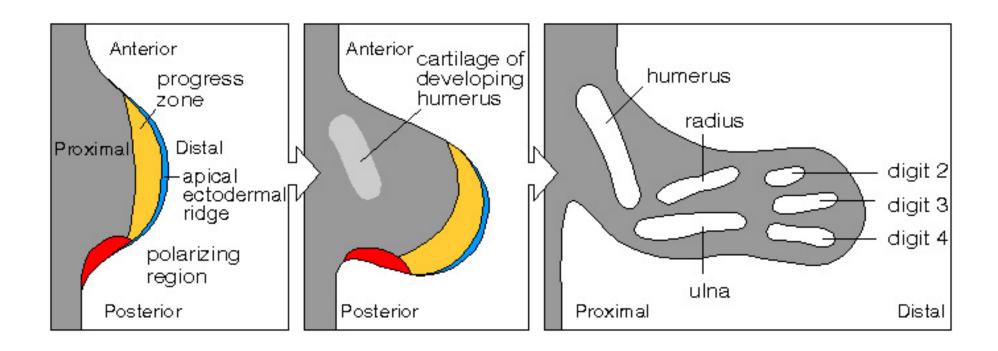


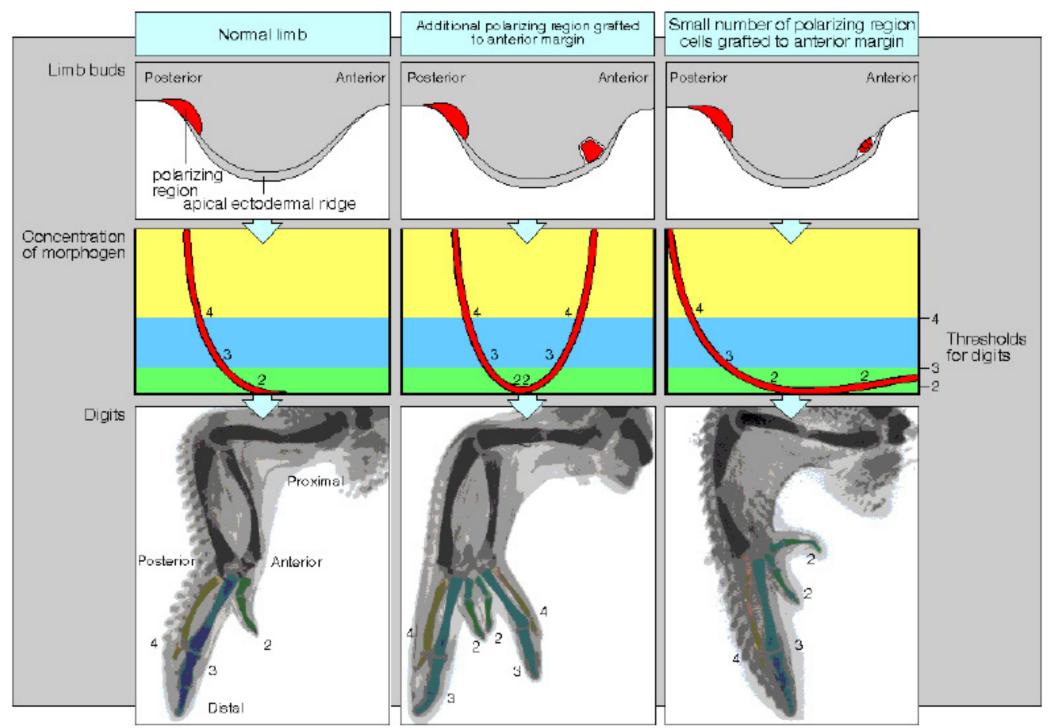
Monk

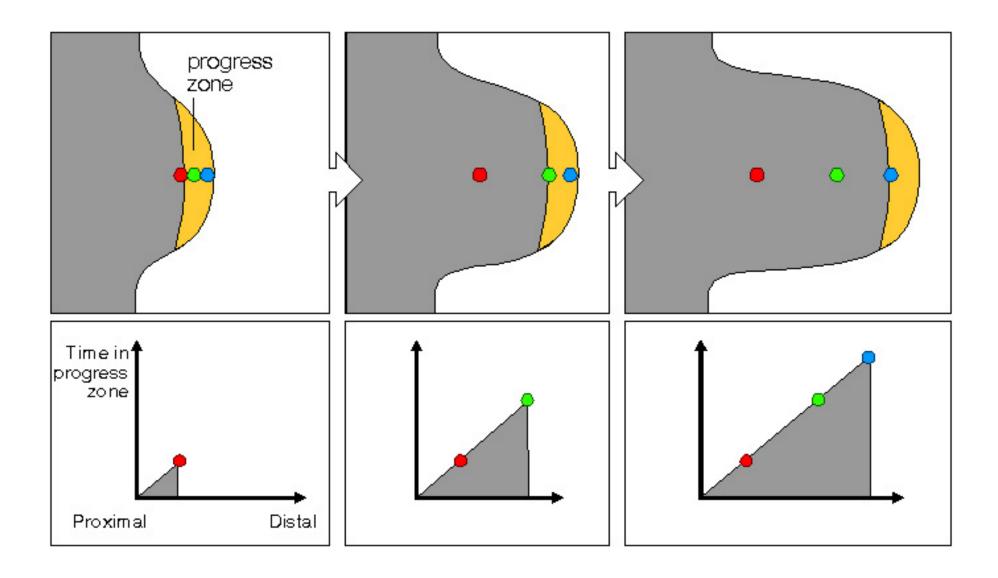
Our results predict that the regulation of Hedgehog transport and stability by glypicans, as well as multiple overlapping feedbacks in the Hedgehog response network, can combine to enhance the robustness of positional specification against variability in Hedgehog levels.

There is a scaling effect which most models of morphogens do not take into account. There are several proposed mechanisms such as two diffusible molecules that emanate from opposing poles and define the activation profile through their ratio. A model based on expansion-repression by Ben-Zvi and Barkai (2010) suggests that scaling emerges as a natural consequence of a feedback topology.

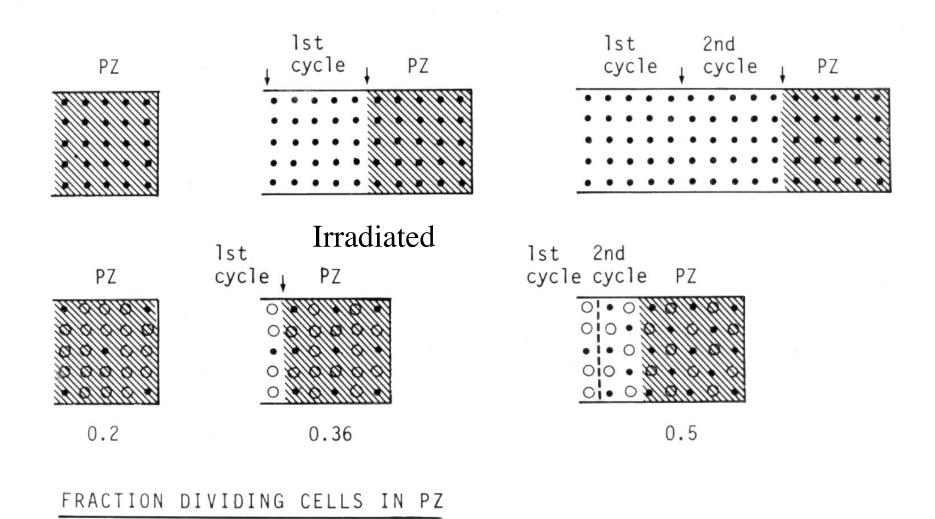




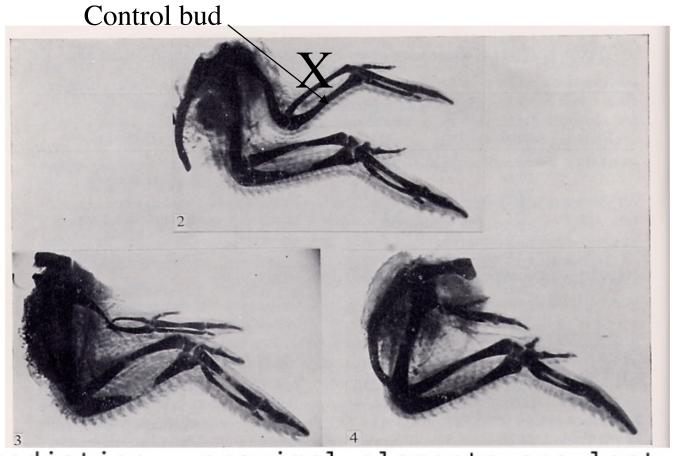




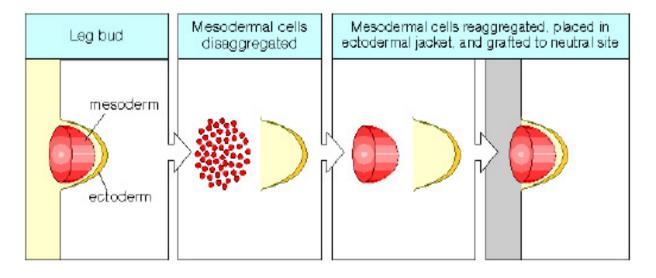
Death in the progress zone leads to loss of proximal elements

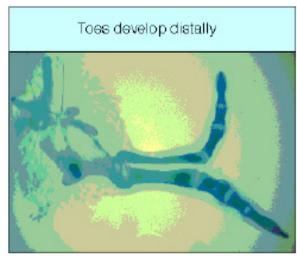


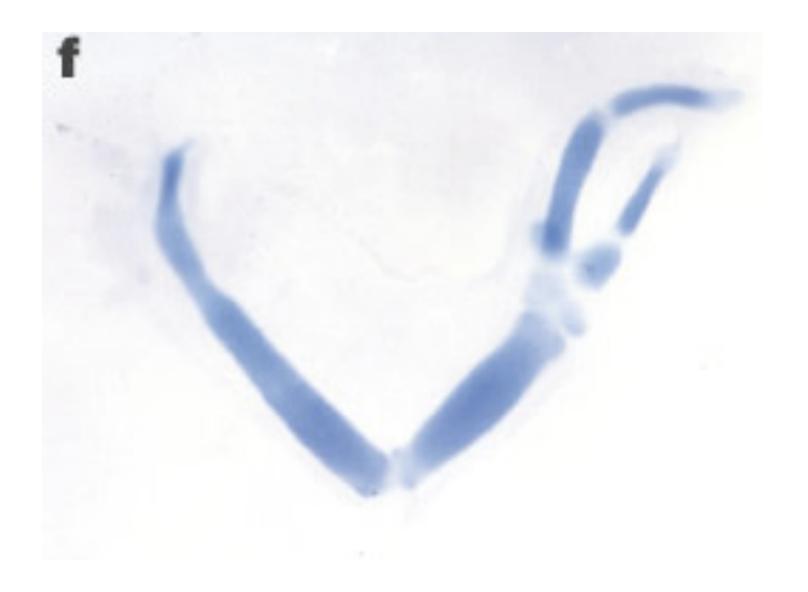
Death in the progress zone - loss proximal elements



Irradiation - proximal elements are lost







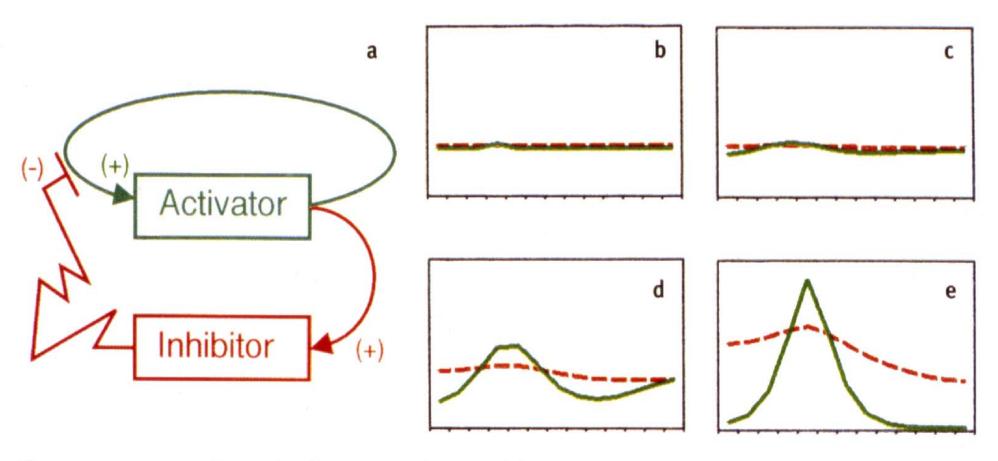
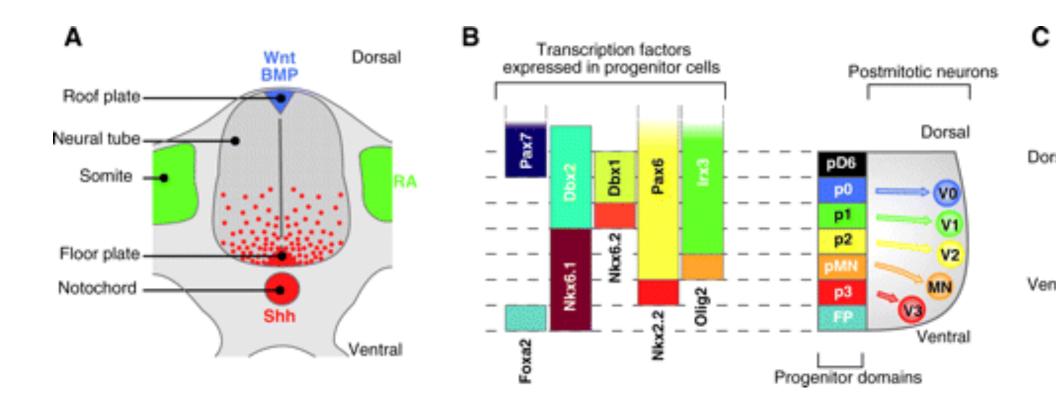


Figure 2.2. Pattern formation by autocatalysis and long-range inhibition. (a) Reaction scheme. An activator catalyses its own production and that of its highly diffusing antagonist, the inhibitor. (b-e) Stages in pattern formation after a local perturbation. Computer simulation in a linear array of cells. A homogeneous distribution of both substances is unstable. A minute local increase of the activator (—) grows further until a steady state is reached in which self-activation and the surrounding cloud of inhibitor (- - - -) are balanced [S22].

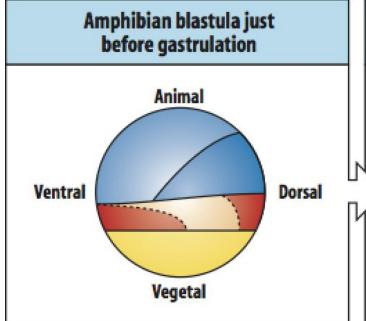
The way in which a signalling gradient regulates differential gene expression in a concentration-dependent manner raises several mechanistic issues.

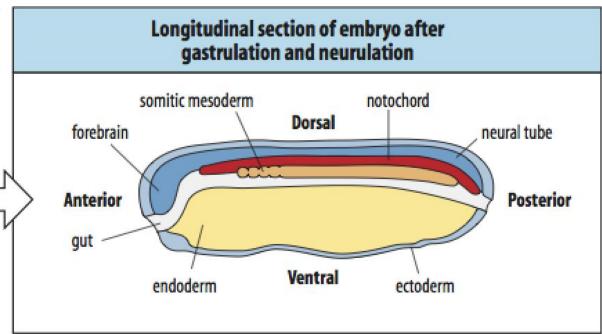
The most detailed model of how gradients can be interpreted comes from how Shh signalling from the notochord assigns the positional identities of distinct neuronal subtype progenitors throughout the ventral neural tube.

The duration of signalling is important not only for the assignment but also for the refinement and maintenance of positional identity.

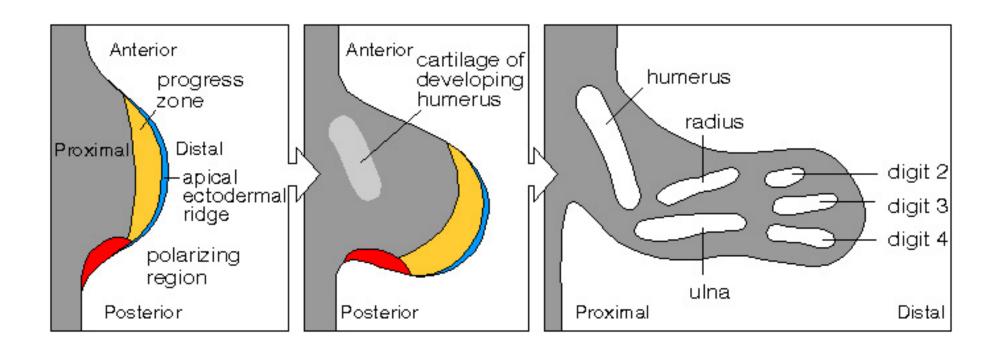


Chordin, a secreted BMP antagonist, is expressed in the dorsal region while on the opposite side BMPs are expressed at high levels. Wnt signals are strong in the posterior region and weaken anteriorly. Cells at different positions within these two Cartesian axes may read these morphogen gradients.





There is however, no good evidence for the quantitative aspects of any of the gradients just described, or details how they are set up.



Growth limb elements

