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Using Xenopus laevis retinal and spinal neurons to study mechanisms of axon guidance in vivo and in vitro

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Abstract

The intricate and precise establishment of neuronal connections in the developing nervous system relies on accurate navigation of growing axons. Since Ramón y Cajal's discovery of the growth cone, the phenomenon of axon guidance has been revealed as a coordinated operation of guidance molecules, receptors, secondary messengers, and responses driven by the dynamic cytoskeleton within the growth cone. With the advent of new and accelerating techniques, *Xenopus laevis* emerged as a robust model to investigate neuronal circuit formation during development. We present here the advantages of the *Xenopus* nervous system to our growing understanding of axon guidance.

Graphical Abstract

Keywords

Xenopus laevis; retinal ganglion neurons; spinal neurons; axon guidance; growth cone

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1. Advantages of Xenopus laevis as a model organism for axon guidance

The complexity of neuronal networks has been a long-standing puzzle that has challenged scientists for centuries. Unveiling how this complex wiring is established in the mammalian brain has, in large part, relied on examination of simpler organisms with comparatively less intricate networks. For example, Ramón y Cajal's work on the chick brain produced the first description of the growth cone [1], [2], and Harrison's work with frogs established the first neuronal culture system [3]. Furthermore, Sperry's pivotal experiment on frog retinal neuron regeneration [4] explained the chemospecificity of connections [5], which has been refined by further studies in systems such as *Xenopus* [6].

Xenopus, as a whole, offers an advantageous complementary vertebrate model, with a multitude of benefits. First of all, recently sequenced genomic data from *Xenopus* shows high similarity with the human genome [7]. There are several species of the *Xenopus* genus, but two have become increasingly popular in research. The diploid western-clawed *Xenopus tropicalis* offers advantages in genomic studies due to its smaller genome. On the other hand, despite its large allotetraploid genome and longer maturation time, the African clawed frog *Xenopus laevis* provides numerous advantages which make it a gold standard for studying axon guidance in development (Figure 1).

The use of *Xenopus laevis* in axon guidance research is advantageous for multiple reasons. Frog husbandry is relatively straightforward, and female frogs can be easily stimulated to produce eggs by simply injecting chorionic gonadotropin hormone. Eggs are comparatively large in diameter, 1-2 mm, and are produced in large quantities. Fertilization occurs *ex utero* and provides the opportunity to track and manipulate embryonic development at desired stages (Figure 2A). Furthermore, embryos can tolerate extensive surgical manipulations varying from microinjection to cell and tissue transplantation. Based upon the given developmental stage and the known fate map of *Xenopus laevis,* it is possible to target specific cell types. For instance, microinjecting mRNA at stages as early as the 1-4 cell stage results in global alteration of gene levels (Figure 2B). Alternatively, injecting embryos at later stages, for example 16-64 cell, allows the restriction of gene manipulations to a more specific tissue [9], [10] (Figure 2C).

Compared to other systems, *Xenopus laevis* neurons can be simply isolated and maintained at room temperature, permitting easy manipulation of live neurons as high resolution images are acquired, forgoing the need for strict incubation conditions such as those provided by CO2 imaging chambers [11]. The primary benefit of *Xenopus laevis* for these studies, however, is its large growth cones, which can be up to 10 to 30 microns in diameter and are perfect for clear and detailed analysis of subcellular cytoskeletal structures and dynamics. There may be no other vertebrate model system with growth cones as large and as easy to culture, manipulate, and image as *Xenopus laevis*.

2. Manipulation of the Xenopus laevis molecular arsenal

Delivery of molecules such as DNA, mRNA, antibodies, or fluorescent dextrans to modify expression of a particular gene or label a specific tissue is available via approaches such as microinjection or electroporation [12]–[15]. Genetic knockdown can be achieved via a

variety of methods, the most common of which has been microinjection of antisense morpholino oligonucleotides (MOs). With the use of standard controls [16], MOs are advantageous tools to manipulate gene products [17], and they have been widely used for *Xenopus* gene knockdown since 2000 [18]. While the effects of MOs last for only a few days, however, the ability to achieve prolonged and heritable gene modifications is now possible with recently developed gene-editing nuclease systems. Transcription activator-like effector nucleases (TALENs), able to deliver high efficiency genetic knockout, have been used in *laevis* for multiple genes [19]–[21]. CRISPR-Cas9, for which it is much easier to produce guide RNA, and which displays even less off-target effects than TALENs [22], has been shown to be effective at disrupting pancreatic genes and pigment genes in *laevis* [22], and this technique will likely useful for investigations into neuronal genes as well. Together, as long as the proper control experiments are conducted, traditional MO approaches and/or newer CRISPR-Cas9 and TALEN systems of genetic manipulation provide complementary tools for efficient alteration of *Xenopus laevis* proteins for axon guidance studies.

In addition to microinjection, electroporation allows manipulation of genes in later stage tissue and can be advantageous over other delivery methods. For instance, if the molecule of interest takes part in neuronal development as well as earlier stages of embryonic development, manipulation of its levels at blastomeric stages may result in lethality or embryonic abnormalities. Therefore, the cell autonomous role of a particular protein during axon guidance is better examined if its function or level is manipulated at stages 20-40 when axonogenesis and brain wiring are still in progress [24] (Figure 2D).

With respect to axon guidance studies, the developing *Xenopus* nervous system provides a valuable working space. In particular, retinal and spinal neurons have been extensively used to decipher aspects of axon guidance machinery. Modern approaches have been developed for both systems to monitor mechanisms of axon guidance *in vitro* and *in vivo*, and this review aims to characterize some of these important examples.

3. Use of the Xenopus laevis retinotectal pathway to study axon guidance in vivo and in vitro

Since studies beginning in 1959 with Gaze [25] and others [6], the visual pathway formation of *Xenopus* has been used as a model for axon pathfinding. The retinotectal neural pathway lies close to the surface, allowing for easy accessibility to neurons for gene manipulation and imaging. Moreover, the pathway is solely formed by the projection of retinal ganglion cells (RGCs), providing a homogenous population of axons that protrude in a stereotypical manner. Most importantly, the basic structure of the pathway as well as the molecules that sculpt this structure are shared between *Xenopus* and mammals [26] (Figure 3).

Christine Holt and colleagues have made significant contributions to the field by developing key techniques to manipulate and image live retinal axons at the cellular level. Seminal studies from the Holt lab have documented the spatiotemporal map and morphology of RGC axons along the visual pathway [27], [28]. Following up the initial observations of growth cones in fixed tissue, it has become possible to trace living growth cones in action. In addition to the Holt Lab, the Cohen-Cory and Cline labs have also made significant

contributions by characterizing the dynamics of retinal axon innervation and synapse formation in the tectum of *Xenopus* tadpoles *in vivo* [29]-[32]. Harris *et al* recorded the motility of growth cones along the optic tract in the living brain and characterized initial and final growth cone advance rates, branching, pausing and retracting behaviors [33]. One of their most striking observations was that the growth cones of retinal axons isolated from their cell bodies are still able to navigate correctly in the tectum. This suggested that the growth cone has the necessary machinery to recognize the guidance cues and determine whether or not to advance, pause, or retract, as it navigates retinal axons.

3.1 Exposed brain preparations for live imaging of retinal growth cones in vivo

The technique that Harris *et al* introduced to examine living growth cones was called the "exposed brain preparation," in which the developing optic tract is exposed in an intact *Xenopus* embryo [33]. With this technique, bath-application of reagents can easily penetrate through the exposed tissue, allowing pharmacological manipulations. The paths of living or fixed growth cones labeled with dyes such as DiI or Horseradish Peroxidase (HRP), respectively, can then be easily traced. By employing these advantages, Chien *et al* tested whether or not filopodia (finger-like, actin-rich protrusions that act as sensory and motor structures) are necessary for growth cone navigation [34], [35], [36]. They demonstrated that bath-applied cytochalasin, a drug that can disrupt actin networks, reduces the number of filopodia and results in slower growth cone advance rates. Moreover, fixed and HRP-filled optic-tracts of *Xenopus* embryos show aberrant pathfinding and retinal axons fail to make caudal turns at the mid-diencephalon [34]. One limitation of this study is that bathapplication of a drug might have a global effect on both the retinal axons addressed and the surrounding neuroepithelial tissue along which axons travel. Targeting solely filopodia may give rise to different results; as shown in later reports, abolishing filopodia through depletion of Xena/XVASP does not cause navigational errors [37]. This example demonstrates how cell-autonomous strategies may be necessary to mitigate certain non-cell-autonomous interferences.

3.2 Electroporation of retinal neurons for targeted gene delivery

Current techniques in *Xenopus* offer ways to overcome non-cell-autonomous impacts by enabling tissue-targeted manipulations via microinjection approaches or electroporation. Electroporation emerged as a successful approach to mediate gene transfer into the intact *Xenopus* nervous system at the single cell level or in bulk during their development (Figure 4A) [13], [14], [15]. While microinjection of embryos sometimes causes early death or developmental abnormalities, electroporation is more efficient and can produce cell or tissue cultures that are in many ways more reliable [24]. Falk *et al* describe a detailed electroporation protocol to specifically study the wiring of *Xenopus* embryonic brains between stages 21-40. Depending upon the region of interest, commonly either brain or eye, embryos are positioned in electroporation chambers tailored to fit their developmental stage. For eye-targeted delivery, a solution containing the molecule of interest is injected into the lumen of the eye of embryos ranging from stage 22 to stage 35/36. Injection is immediately followed by a voltage application with the appropriate frequency and duration. GFP expression can be readily observed 12h after electroporation. Importantly, electroporation of different regions of the same embryo is possible [24]. This dual-electroporation technique is

highly exciting because it allows, for example, simultaneous manipulation of receptors on the growth cone and guidance cues on its terrain [38], making it possible to study axonpathway interactions during pathfinding.

In addition to DNA electroporation, MO electroporation is described to knockdown gene expression in *Xenopus* embryos [24] and tadpoles [39]. At early-stages (2 to 64 cell), MO can be directly injected into the cell; however, beyond these cell-stages, the neutral MO is confronted by the negatively charged cell membrane. This is why conventional MOs cannot be used for electroporation; instead, tagging MOs with a negatively charged fluorescent reporter (such as Lissamine™) ensures both tracing and delivery. MO can interfere with protein expression as soon as 1h after electroporation and can persist up to 48h after electroporation. Co-electroporation coupling DNA/DNA or DNA/MO also yields efficient uptake with this technique [24], [39]. For example, Bestman *et al* used this approach and coelectroporated MO along with a plasmid that drives GFP expression under the control of cell specific promoter to identify genes that contribute to neurogenesis of the optic tectum in *Xenopus* tadpoles [40].

Coupling the aforementioned gene manipulation approaches with *in vivo* imaging techniques allows for the deciphering of guidance machinery components while the growth cone continuously makes decisions along its terrain [41]. A study by Leung *et al* beautifully exemplifies this. The group spatially modulated the function of NF-protocadherin (NFPC) via targeted electroporation of either dominant negative NFPC E-myc or NFPC-MO in retinal axons or along their substrate along the optic tract in brain. DiI staining and live imaged then revealed navigational errors along the mid-optic tract, suggesting a requirement for homophilic interactions between the axon and its substrate at this region [38] (Figure 4A-C).

3.3 Whole pathway explant preparation to assess retinal growth cone response in vitro

While these studies enable us to understand the topography of axon pathfinding in an intact organism, it still remains to be understood how an individual growth cone is modulated during guidance decisions. Thus, some current investigations have shifted toward understanding the intracellular machinery. In this regard, Lohof *et al* developed an approach to assess the behavior of individual cultured *Xenopus laevis* growth cones in response to a gradient of signals repetitively released by micropipette [42]. Taking advantage of this technique, de la Torre *et al* demonstrated that the growth cone turning of RGCs induced by netrin-1 involves the DCC receptor to mediate the chemotropic action of netrin-1, as antibodies to DCC interfere with the chemotropic response to netrin-1 [43]. This experimental system has also been extensively used with *Xenopus* spinal neurons (discussed in section 4.1).

RGC axons first leave the retina at stage 28 [24]; therefore, any attempt to culture the retina beyond that stage will sever axons and represent a regeneration study rather than a study of *de novo* axon outgrowth. Shewan *et al*, however, developed an approach they term "whole pathway explant preparation," used for testing the responsiveness of *de novo* growing RGC axons to a netrin-1 signal at stages ranging from 24 to 40 during visual pathway formation [44]. Based upon the visual pathway map [27], cuts can be made ahead of the growth cones

at various stages along the visual pathway. Cultured retinal axons extending from the cuts onto the substrate can then be tested for their response to a specific chemotropic gradient. The purpose of whole pathway explant testing is to allow axons to experience their native environment up until the point where their response is examined *in vitro*. Using this method, Shewan *et al* found that the response of RGCs to netrin-1 gradients depends on the developmental stage, as young retinal axons (pre-optic nerve head) show attraction toward gradients of netrin-1, whereas older axons (post-chiasm) are repelled [44]. This study revealed that growth cones aged in culture without experiencing a native pathway also switch their response in parallel with the response of axons *in vivo*, suggesting the presence of an intrinsic molecular program within the growth cone for response generation. The intrinsic molecular program proposed by Shewan *et al* supports the findings of Harris *et al* on growth cones' ability to navigate through the optic tectum *in vivo* even when separated from their cell bodies [45].

3.4 Photoconvertible protein recovery to investigate local mRNA translation

Campbell and Holt, using the aforementioned chemotropic assay, demonstrated for the first time that the attractive and repulsive responses of cultured *Xenopus* retinal growth cones to netrin-1 and Sema3A are abolished in the growth cone upon inhibition of local protein synthesis with anisomycin and cyclohexamide [46], suggesting a translation-dependent chemotropic response in growth cones. The role of translation also explains the reduction in immunoreactivity to DCC in cultures of old axons compared to young axons concomitant with reduced attraction response to netrin-1 that Shewan *et al* observed [41].

To further investigate the dynamics of protein translation in response to gradients of guidance signals, a photoconvertible protein called Kaede can be linked to a protein of interest [46]. Neuronal cultures can be established from *Xenopus* embryos injected with mRNA for this construct. The initial green fluorescence of Kaede can then be converted into red light upon UV irradiation. With *de novo* protein synthesis of the injected mRNA, green signal is recovered, allowing for an assay that can help determine the rate at which the protein of interest is translated under certain conditions [47]. Using this technique, Leung *et al* showed that *Xenopus* retinal growth cones introduced to gradients of netrin-1 initiate a recovery of green signal near the netrin-1 applied site when Kaede is linked to beta-actin, implying netrin-1-induced asymmetric beta-actin translation. This recovery occurs in isolated growth cones as well, suggesting an increase in translation controlled by local machinery [48]. The advent of photoconvertible molecules shows promise for furthering our understanding of protein turnover in cells, especially in the area of the tightly regulated local translation of the growth cone.

4. Xenopus laevis spinal neurons to study axon guidance in vivo and in

vitro

Another well-studied neuronal system in regard to axon guidance is that of *Xenopus* spinal neurons. Previously described techniques used for manipulation of gene expression in RGC neurons are applicable to spinal neurons as well [49]. *Xenopus* spinal neurons have been central to the examination of the interaction between extracellular guidance cues [50], [51],

their receptors [51], [52] and cytoplasmic secondary signals [53]-[55] in growth cone turning events. Axon guidance mechanisms that operate during development can be studied *in vitro* with cultured spinal neurons prepared at the time (stage 20) that the neural tube closes and axon outgrowth just begins. Isolation or manipulation of neurons beyond this stage can be used to study the dynamics of axon regeneration [56], [57]. In this section, we explore some of the ways in which *Xenopus* spinal neuron techniques have been used to help develop a further understanding of axon guidance.

4.1 Modulation of intracellular signals in spinal neuron growth cones

Asymmetric application of molecular gradients to cultured *Xenopus* spinal growth cones and examination of subsequent growth cone turning response elucidate the link between the extracellular cues cytoplasmic signals in growth cone turning events [42] (Figure 5). As well as testing the impact of a single type of molecule on growth cone turning, homologous and heterologous modulations can be tested by bath applying molecules and simultaneously administering them in gradients. This approach allows for characterization of adaptation phenomena in growth cones that undergo consecutive desensitization and re-sensitization cycles, such as in response to gradients of netrin-1 when netrin-1 is also bath applied in culture. This adaptation effect can also be observed heterologously between two guidance cues that share common cytosolic transduction pathways, such as with netrin-1 and BDNF [50].

In addition to guidance signals, secondary messengers like cAMP, cGMP or Ca^{++} and their inhibitors, can also be applied in gradients and/or via bath application. Studies have identified that when cAMP is uniformly applied to *Xenopus* spinal neuron cultures, the attractant turning response toward BDNF [53] and netrin-1 [54] was inhibited and instead induced repulsive turning. Similarly, a repulsive response to Sema3D was found to be converted to attraction when a cGMP agonist or cAMP antagonist was applied to the culture [55].

Moreover, interaction between receptors and their associated guidance signals can be tested by bath applying antibodies that occupy receptors and interfere with signal transduction. When an antibody against the extracellular domain of DCC, a netrin-1 receptor, was introduced to the *Xenopus* spinal neuron cultures, netrin-1-induced attractive turning was abolished [54]. Alternatively, spinal neurons expressing wild-type or chimeric receptors isolated from *Xenopus* embryos microinjected with mRNA encoding receptors can be tested for receptor engagements. Using this approach, Hong *et al* identified that interaction between the cytoplasmic domains of the UNC-5 and DCC receptors is required to generate a repulsive response to netrin-1 gradients [52]. Interactions between receptors of opposing signals, like netrin-1 and slit, have also been reported in *Xenopus* spinal neurons, underlying the hierarchical response to attractive and repellent signals [53] during midline crossing.

Techniques used *in vitro* to investigate signal transduction mechanisms within the growth cone can be adapted to be used *in vivo*, as well. Bath application of guidance molecules, pharmacological agents, antibodies, or fluorescent dyes can be taken up by an exposed *Xenopus* spinal cord. Moreover, asymmetric elevation or inhibition of molecules can be achieved by photoactivatable caged molecules or optogenetic approaches [58] (Figure 5B-

C). For example, Gomez and Spitzer used these approaches to investigate Ca^{++} dynamics, which have been shown to be responsible for growth cone turning events downstream of neurotransmitters and guidance signals [59] – [62]. They exposed the anterior half of the *Xenopus* spinal cord and labeled neurons with the fluorescent calcium indicator Fluo-3 AM. Subsequently, exposing the posterior half of the spinal cord enabled them to trace labeled growth cones and measure the frequency of endogenous $Ca⁺⁺$ transients as axons elongated into the unlabeled region. Additionally, by loading neurons with photoactivatable caged diazo-2 AM or NP-EGTA AM, the group performed targeted reduction or elevation of Ca^{++} levels, respectively. Their results demonstrated that Ca^{++} transients are regulated in a region-specific manner, and that higher frequencies of Ca^{++} transients reduce the axon elongation rate and increase growth cone stalling [63].

4.2 Extracellular matrix and adhesion dynamics in Xenopus laevis spinal neuron guidance in vivo

In addition to guidance signals, growth cone pathfinding is influenced by components of the ECM [64]. *Xenopus* spinal axons travel through environments presenting different ECM makeups, and engage with them via integrin receptors at adhesion sites [64]. Adhesions formed between the growth cone and the ECM are referred to as "point contacts" [65], [66]. Formation, maintenance, and turnover of these contact sites determined by intracellular signaling events are important determinants of growth cone morphology and motility, and hence pathfinding behavior [67], [68].

Examination of these sites mostly relies on immunofluorescence labeling of cell adhesion signaling proteins in fixed and living growth cones, followed by high resolution microscopy imaging [69]. Labeling or modulation of protein levels can be achieved by microinjecting mRNA of fluorescently tagged phosphotyrosine reporter constructs [69], adhesion markers, or their corresponding MOs. Antibodies are also available for targeting *Xenopus* scaffolding proteins involved in focal adhesion [69], allowing for assessment of their spatial distributions within lamellipodia or filopodia in fixed growth cones [70] via immunocytochemistry analysis. Live cell imaging is also possible: this permits monitoring of the changes in protein localization, modification, or turnover as the growth cone moves along different ECM substrates and makes navigational decisions in response to cues. Examples of such investigations come from the work of the Gomez Lab; Robles and Gomez characterized focal adhesion kinase (FAK) activity in *Xenopus* spinal growth cones and showed that tyrosine-phosphorylated protein accumulation depends on the formation of integrin-dependent adhesive contacts with the ECM, which ultimately impact growth cone motility and outgrowth [71]. Another work by the Gomez group examines adhesion dynamics upon guidance cue stimulation. They show that BDNF and netrin-1 gradients increase phospho-tyrosine-positive filopodial tips through activation of FAK, and hence Src, coinciding with enhanced filopodial motility and suggesting an asymmetric distribution of motile filopodia in growth cone steering [70], [72].

4.3 Xenopus laevis growth cones to assess cytoskeletal rearrangements during guidance

The above-mentioned *Xenopus* studies are examples of how interactions with guidance signals and ECM components give instructions through signaling molecules to affect growth

cone protrusion and steering during pathfinding. Downstream of these, the reorganization of the cytoskeletal polymers, actin filaments and microtubules modulate growth cone steering [73], [74]. Although neurofilaments are less well-studied, this third member of the axonal cytoskeletal polymer family also plays a crucial role, and *Xenopus* has been a particularly useful model to examine the role of neurofilaments as a modulator of axon growth during development and regeneration [75] - [78].

Examination of the growth cone in detail is difficult in an intact tissue, as it travels in a 3D environment, and this makes it challenging to track the moving growth cone without losing focus of its cytoskeletal features. Although recently developed imaging approaches have begun to make *in vivo* growth cone cytoskeleton examination possible [79], most of our knowledge on the structural organization of the growth cone comes from *in vitro* studies [80], [81]. In this regard, with their large growth cones, *Xenopus laevis* neurons provide an ideal system to study cytoskeletal dynamics. With the advance of protein labeling and highresolution imaging techniques, it is possible to track fluorescently tagged proteins to gain insights into the motility dynamics of the growth cone during axon growth and guidance.

The leading edge of the growth cone, where initial contacts with the adhesive substrate are made, is assembled by a filamentous actin network. Dynamic remodeling of the actin network in this region allows the growth cone to explore and interact with the molecules in the extracellular milieu [82]. Tracing both actin and its interacting partners in living growth cones allows for a better understanding of the mechanism by which actin is remodeled downstream of guidance signals [83], [84]. Actin labeling is mostly achieved by expressing fluorescently tagged actin or actin binding peptide markers such as F-lifeact- GFP [85] or utrophin [86], which bind along the F-actin lattice, or by loading growth cone cultures with cell-permeable kabiramide C conjugated to tetramethylrhodamine (TMR-KabC) [87], which binds the growing ends of F-actin and allows for visualization of actin dynamics such as growth, depolymerization, or retrograde flow. Alternatively, immunolabeling against phalloidin, an F-actin binding protein, can be used to monitor actin localization in fixed growth cones [79].

Expressing rhodamine-labeled tubulin emerged as a technique to track singular microtubules in cultured living *Xenopus* growth cones [85]. Tracking labeled microtubules revealed, for example, that spatial stabilization of microtubules via FLIP-released taxol can mediate growth cone turning [88]. Thus, mechanisms that promote microtubule growth have gained further attention. Microtubule binding proteins, particularly plus-end tracking proteins (+TIPs), have become a focus of study as they directly regulate microtubule plus-end dynamics [89], and tracking and quantitative analysis of fluorescently-tagged +TIPs in cultured *Xenopus* growth cones has provided new insights into how microtubule dynamics are regulated in the growth cone [90], [91]. Lee *et al* unveiled a mechanism for the +TIP CLASP and proposed that growth cones exposed to asymmetrically distributed repellent signal Slit triggers spatially restricted Abl kinase activation in the growth cone, which then impedes microtubule growth by targeting CLASP phosphorylation [92], [93]. In addition to CLASP, multiple other +TIPs have been shown to be potentially involved in the regulation of microtubule dynamics during axon outgrowth and steering events [94].

Furthermore, acquisition of high-resolution live images of fluorescently-labeled known +TIPs, such as EB1, in cultured *Xenopus* growth cones can be used to measure parameters of microtubule growth dynamics *ex vivo* [95]. There is open-source software, such as plusTipTracker, available for automated detection and analysis of these tagged +TIPs [96], [97]. Additionally, tools such as quantitative fluorescent speckle microscopy (QFSM) or kymography are also commonly used to extract quantitative information regarding cytoskeletal dynamics from high-resolution microscopy images of growth cones [90].

5. Conclusions

Since the first speculations on the phenomenon of axon guidance, studies using the *Xenopus laevis* nervous system have made significant contributions toward our understanding of this crucial developmental orchestration. Developing *Xenopus* neurons have been a key subject from the characterization of nerve patterns of the visual system to the identification and creation of novel mechanisms and techniques.

Although many of the techniques described in this review can be applied to higher vertebrate models such as mouse [98]–[101], rat [102], or chick [103], [104], the affordability and practicality of *Xenopus*, coupled with its large and easy-to-culture growth cones, makes it a highly attractive model system for deciphering the molecular mechanisms of axon guidance. Its phylogenetic position also makes it a preferable organism in comparison to other experimentally-utilized lower vertebrates or invertebrates.

In conclusion, *Xenopus laevis*, with existing and newly emerging techniques, will persist as an excellent model to study elusive aspects of the axon guidance mechanism both *in vivo* and *in vitro*, from network organization, to the single neuron or growth cone, and all the way down to the single molecule level.

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Highlights

- **•** *X. laevis* is an affordable and practical vertebrate model for *in vitro* and *in vivo* studies of axon guidance.
- **•** Gene manipulation techniques are well established and can be applied at any developmental stage.
- **•** Retinal and spinal axonal tracts are accessible and can be live imaged for guidance decisions *in vivo*.
- **•** Neurons from *X. laevis* retina and spinal cord are easy to isolate, maintain and manipulate *in vitro.*
- **•** *X. laevis* growth cones are large enough for high-resolution live imaging.

Figure 1. Timeline of *Xenopus laevis* **in axon guidance**

Key findings and important advances in *Xenopus laevis* (shown in green boxes) as a model organism during the development of the axon guidance field.

Figure 2. Gene manipulations at various stages of *Xenopus* **embryonic development**

(A) Female frogs are injected with chorionic gonadotropin (CG) 12-18 h before egg collection. Eggs are collected and maintained in a salt solution and fertilized with minced testes. Owing to its *ex utero* development, Nieuwkoop and Faber (NF) based developmental stages can be tracked as the development proceeds at room temperature and manipulations can be performed at desired stages. **(B)** Injection at early stages (2-8 cell) will target larger tissues, however targeting lighter blastomeres which form the dorsal tissues will target neural plate or notochord. **(C)** Injection at later stages will target more specific tissues. Based on the *Xenopus* fate map, retinal cells derive primarily from blastomeres D1.1.1 and D1.2.1 in a 32-cell stage embryo. The neural tube receives its major contribution from D1.1 and D1.2. **(D)** Electroporation for spatiotemporal targeting of the nervous system is possible particularly between stages 21-40, when retinal axonal tracts are first forming.

Figure 3. Diagram of embryonic *Xenopus* **retinal pathway map**

The spatiotemporal position of axons and guidance signals effective along the visual pathway during embryonic development. RGC retinal ganglion cell, ONH optic nerve head.

Figure 4. *Xenopus* **visual pathway as a model of axon pathfinding**

Manipulation of genes in retinal axons and axonal pathways in the same embryo is possible via eye and brain targeted electroporation respectively. **(A)** Retinal axons and the neuroepithelium of the optic tract can be electroporated between stage 27 after axon initiation begins and stage 32 when the first retinal axons pass the optic chiasm and enter the optic tract in the brain. **(B)** For complete pathway analysis, embryos can be fixed at stage 40 and labeled with DiI or HRP. Labeled retinal pathway can be exposed via open brain preparation and pathfinding behaviors of axons can be analyzed. **(C)** Open brain preparations prepared in living embryos after stage 32 allows live imaging of retinal axons as they navigate through the optic tract in the brain. Time-lapse movies of axons can be recorded for 24h with 3min intervals.

Figure 5. Assessing the response of cultured growth cones to guidance signals and secondary messengers

(A) Repetitive pulse application method to assess growth cone response to chemical gradients released via micropipette positioned 100 microns away from the center of the growth cone at an angle of 45°. **(B-C)** Growth cone response to asymmetrical alterations of intracellular secondary molecules Ca^{++} and cAMP. **(B)** Focal laser induced photolysis of caged Ca++ loaded (NP-EGTA) growth cones [60]. **(C)** Optogenetically generated cAMP transients in growth cone expressing photoactivated adenyl cyclase (PAC) [55].