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Genetic dissection of axon regeneration

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Abstract

Axon regeneration has long been studied in vertebrate model organisms and neuronal cultures. Recent development of axon regeneration paradigms in genetic model organisms, such as *C. elegans, Drosophila* and zebrafish, has opened an exciting field for *in vivo* functional dissection of regeneration pathways. Studies in these organisms have discovered essential genes and pathways for axon regrowth. The conservation of these genes crossing animal phyla suggests mechanistic relevance to higher organisms. The power of genetic approaches in these organisms makes largescale genetic and pharmacological screens feasible and can greatly accelerate the mechanistic understanding of axon regeneration.

Keywords

Axon regeneration; genetic models; laser axotomy; DLK MAPKKK; MAPK signaling; mTORdependent protein synthesis

Introduction

The limited abilities of mature neurons in adult nervous systems to regenerate or repair damages remain one of the poorly understood phenomena in neuroscience and have been intense subjects of investigation for over a century. Decades of research primarily from vertebrate models, such as rodents and fish, has established that both intrinsic properties of injured neurons and extrinsic environments play intricate roles in the extent and accuracy of regenerating axons. Recent progress in the identification of extrinsic factors, particularly myelin-associated inhibitors, and of intrinsic regeneration-promoting pathways has rapidly moved the field into the molecular era. Studies from *in vitro* cultured neuronal models and from lower vertebrates and invertebrates such as Aplysia have also greatly enhanced our knowledge of cellular dynamics, particularly in the early phase of injury responses [1]. An exciting development in the last few years is the establishment of axon regeneration models in genetic organisms including the nematode *Caenorhabditis elegans*, the fruitfly *Drosophila melanogaster* and the zebrafish *Danio rerio*. Emerging findings from these model organisms have shown great promises of using forward and reverse genetic manipulations to discover key players of axon regeneration. In this minireview, we will first

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Newly developed axon regeneration paradigms in genetic organisms

The small size of neurons in *C. elegans*, Drosophila and lower vertebrates presented technical challenges for nerve injury studies. In 2004, Yanik *et al.* developed a femtosecond laser surgery protocol to perform acute *in vivo* axotomy on transgenically labeled fluorescent axons in live *C. elegans* animals. They found that the GABAergic motor neuron axons are capable of functional regeneration post laser axotomy [2]. Follow-up studies by Wu *et al.* and Gabel *et al.* also demonstrated that neurites of various neurons in *C. elegans*, including mechanosensory neurons, chemosensory neurons and other motoneurons, can regenerate with different degrees of regrowth ability [3–5] (Table 1). More recently, customized microfluidic devices have been developed to facilitate laser axotomy and live imaging of regenerating axons in awake animals, and to enable high-throughput genetic and pharmacological screenings [6]. Similar laser axotomy techniques have also been developed in zebrafish [7–9] and *Drosophila* [10] (Table 1). Laser-severed trigeminal sensory neurons in zebrafish larvae can regenerate into skin, but their ability to innervate originally denervated skin territory depends on developmental stages [8]. A high-throughput screen method using laser axotomy in zebrafish larvae has also been established very lately [7]. Laser transection of axons or dendrites of Class I dendritic arborization (da) neurons in the fly brain triggers distinct patterns of cytoskeletal rearrangement and neurite regrowth [10]. These efforts have begun to push forward the understanding of genetic pathways regulating axon regeneration (Table 1). They have also prompted laser axotomy as a tool to study regeneration-associated events in mouse hippocampal neurons [11]. Importantly, as described below, the cellular responses and self-repair signalings triggered by laser axotomy exhibit fundamental similarities to those induced by mechanical surgery.

The roles of mitogen-activated protein kinase (MAPK) pathway in axon regeneration

Two parallel studies published in 2009 have uncovered a critical role of a conserved MAP kinase in axon regeneration in *C. elegans* [12,13] (Figure 1A). In a mutant strain that is defective in β-spectrin, a component of the cortical cytoskeleton, axons of GABAergic motor neurons are fragile and undergo progressive breakage caused by locomotion. These broken axons can constantly regenerate toward the dorsal segments in an attempt to re-build their connections [14]. Hammarlund and his colleagues performed a large-scale RNAi screen in this strain in order to isolate mutants showing diminished spontaneous axon regrowth. The DLK-1 (Dual Leucine zipper-bearing Kinase 1), a member of an evolutionarily conserved MAP kinase kinase kinase family, was identified as a candidate with the strongest effect [12]. In a separate study, Yan *et al.* found that laser-severed PLM mechanosensory neurons also fails to regenerate in *dlk-1* loss-of-function *(lf)* mutant animals [13], whereas overexpressing DLK-1 in these neurons accelerates regrowth following laser axotomy. In both type of neurons, DLK-1 functions cell-autonomously for growth cone formation and initiation of regrowth. It acts through two downstream kinases, the MKK-4 MAP kinase kinase and the PMK-3 p38 MAP kinase [12,13]. This kinase cascade has been previously shown to function in the developing nervous system to regulate synapse formation [15]. Experiments using acutely induced rescue of DLK-1 expression in *dlk-1(lf)* mutant animals show that DLK-1 is required within a short temporal window around the time of injury, suggesting that its activation needs coincide with injury signals to initiate regrowth [12] (Figure 1A).

In the nervous systems of *Drosophila* and mouse, the DLK MAPK pathway has been shown to regulate Wallerian-type axon degeneration, a process usually preceding or paralleling axon regeneration in which axons detached from injured neurons undergo self-destruction [16,17]. Miller *et al.* examined a degeneration paradigm in the olfactory receptor neurons (ORN) in adult flies, and reported that loss of function in DLK (known as Wallenda in *Drosophila*) slows the Wallerian degeneration [17] (Figure 1B). The degenerationpromoting effect of DLK is also observed in the distal segments of peripherally injured mouse dorsal root ganglion (DRG) neurons [17]. In another parallel study, Massaro *et al.* showed that in *Drosophila* larvae disassembly of the neuromuscular junction (NMJ) caused by loss of α -spectrin in presynaptic motor neurons surprisingly shares mechanistic similarities with Wallerian degeneration [16]. However, elevation of the DLK activity, resulted from loss of the DLK -degrading E3 ligase Highwire (Hiw, *Drosophila* homolog of *C. elegans* RPM-1), suppresses NMJ retraction in the spectrin mutant [16] (Figure 1C). This finding is consistent with the growth-promoting function of DLK in *C. elegans* neurons but apparently contrary to Miller *et al.*'s finding. The differential effects of DLK in these studies could reflect the context-dependent and compartment-specific (proximal *vs* distal axon fragments) functions of this injury response pathway. In both studies, DLK acts through the Jun N-terminal kinase (JNK) MAP kinase. JNK has also been reported to promote axon regeneration in the small lateral neurons ventral (sLNv) injured by microdissection in the whole explanted *Drosophila* brain [18], although the role of DLK/Wallenda is yet to be examined (Figure 1B).

Take together, the above findings not only demonstrate the important function of the DLK pathway in adult axon injury responses, but also reveal the complexity underlying the context-dependent action of these conserved kinases. DLK can stabilize injured axons and promote axon regrowth from proximal axon ends, mostly likely by acting as the immediate injury sensor. It can also protect disrupted NMJ synapses from disassembly. Meanwhile, it can accelerate degeneration of distal axon fragments, which may facilitate regrowth by removing biochemical and physical barriers presented by these fragments. It is important to note that in the developing nervous system DLK kinases are critical regulators of synapse formation and growth [15,19]. These studies bring about a surprising convergence between disparate cellular processes in the developing and mature nervous systems. Many questions remain to be addressed regarding the DLK signaling in axon regeneration. For example, how is the DLK MAPK cascade activated by injury signals in the first place? How does the DLK pathway regulate axon regeneration? Microtubule dynamics and global gene transcription have been suggested to mediate the effects of the DLK signaling [20–22]. Moreover, multiple MAPK pathways exist in *C. elegans, Drosophila* and vertebrates. Do other uncharacterized MAPK pathways play similar or divergent roles in axon regeneration? Further investigations will be directed to fully understand both the general signaling mechanisms and also the neuron- or axon-type specific effects of the DLK cascade in axon regeneration and degeneration.

The role of second messengers and cytoskeletal rearrangement in injury responses

Nerve injury-triggered signals lead to a series of changes, including rearrangement of cytoskeletons, retrograde transport of injury-responding factors, and changes in gene expression, all of which contribute to the repair and regeneration processes [1,21,23,24]. Axonal injury in cultured neurons can cause elevation of intracellular Ca^{2+} and cAMP, which serve as initial injury messengers [25–27]. However, it was poorly understood whether these initial signals promote axon regeneration *in vivo* and what their downstream effectors are. Recently, Ghosh-Roy *et al.* found that in *C. elegans* PLM mechanosensory neurons, genetic elevation of Ca^{2+} and cAMP can accelerate formation of regenerating

growth cones, facilitate fusion of damaged axons, and promote formation of synaptic branches [28]. Furthermore, the regrowth-promoting effects of Ca^{2+} and cAMP rely on protein kinase A (PKA) and the DLK-1 MAPK pathway (Figure 1A). Pharmacological inhibition of PKA activity reduces axon regrowth while genetic enhancement of PKA activity causes an opposite effect [28]. A similar regrowth-promoting effect of PKA has also been observed in lateral neurons ventral (LNv) in fruitfly brain [18] (Figure 1B).

Axon injury also triggers dynamic rearrangements of actin filaments and microtubules, which contribute to the initial injury signal propagation and subsequent axon extension [29,30]. Reggie-1 and Reggie-2, two conserved homologous proteins originally identified as plasma-membrane associated scaffolding proteins, are up-regulated in retinal ganglion cells (RGCs) after optic nerve section in goldfish and rats [31,32]. Recent studies have revealed that they function in axon regeneration via regulating F-actin dynamics [33,34]. Knocking down Reggies in zebrafish RGCs impedes axon regeneration *in vivo*, likely by affecting the activation status of Rho GTPases, the N-WASP complex stability, and activation of p38 MAPK and focal adhesion kinase (FAK) [35]. In mice, many environmental inhibitors such as myelin-associated inhibitors act through the actin regulator RhoA and its downstream Rho-associated kinase II (ROCKII) to inhibit axon regrowth [23,36]. Deletion of ROCKII makes neurons less sensitive to inhibition by Nogo or CSPG and improves axonal regrowth after trauma in the adult spinal cord [37].

Isoform-specific regulation of tubulins has long been observed in various injury models. α1 tubulin isoform Tuba1a is up-regulated in zebrafish RGCs after optic nerve injury [38]. Knockdown of Tuba1a, but not the closely related Tuba1b, dramatically hinders RGC axon regeneration [39]. In *Drosophila*, a global up-regulation of microtubule dynamics occurs when an axon of a neuron is surgically removed. A regenerating axon will be generated from an existing dendrite by the reversal of microtubule polarity in this neurite. Such drastic microtubule rearrangements require the JNK signaling [10]. Similar dendrite-to-axon conversion post axotomy is also observed in cultured mature mouse hippocampal neurons [11]. Conceivably, injury-induced kinase activity can modify the property of cytoskeletal components or the activity of cytoskeleton-binding proteins. Such changes may create new sites of cytoskeleton dynamics and/or facilitate transduction of injury signals mediated by motors and other proteins. Although the regulation of regenerative cytoskeleton dynamics is not fully understood, recent data have suggested that it can be different from that during development. In *C. elegans*, homologs of Rac, lamellopodin, and Enabled/VASP are required for regenerative growth cone dynamics but not indispensable for initial developmental growth [3]. These studies underscore the importance of further investigation of the mechanisms regulating cytoskeleton dynamics in axon regeneration.

The roles of injury-triggered gene expression in axon regeneration

The intrinsic regenerating ability of injured axons heavily relies on gene transcription and protein translation [24,40,41]. Recent studies have started to reveal key players in these processes. In axon regeneration of worm mechanosensory neurons, a key downstream effector of DLK-1 is a basic leucine zipper (bZip) transcription factor CEBP-1 (the CCAAT/ enhancer-binding protein 1) [13]. CEBP-1 mRNAs can be detected in axons, and axotomy induces local translation of axonal CEBP-1 mRNAs in a DLK-1-dependent manner. It remains unknown whether these locally translated CEBP-1 proteins may function retrogradely in the nucleus as a transcription factor or have other roles in the axon [13]. In *Drosophila*, the JNK signaling stabilizes transiently disrupted NMJ through the bZip transcription factor Fos in the nucleus [16]. In mice, C-Jun, another downstream target of JNK, is involved in facial nerve regeneration [42]. In *C. elegans*, the Jun homolog JUN-1 is also required for axon regrowth of mechanosensory axons [28]. In contrast, the cAMP

response element-binding protein CRH-1/CREB is not required for regrowth, but instead promotes the ventral branching in the same type of neurons [28]. These studies suggest diverse roles of the bZip family of transcriptional factors in axon regeneration and degeneration.

Another family of transcription factors recently emerged from axon regeneration studies is the Zinc finger Kruppel-like transcription factors (KLFs). In zebrafish RGCs, KLF6a and KLF7a and their transcriptional target α 1 tubulin isoform Tuba1a are highly induced after injury and promote RGC axon regrowth in explant culture [38,39]. In mouse RGCs, the KLF family is also suggested to play important roles in neurite outgrowth and axon regeneration [43]. KLF4 was identified as a strong suppressor of hippocampal neurite outgrowth. KLF4 knockout mice showed significantly decreased regeneration of adult RGCs *in vivo*. Moreover, outgrowth-enhancing KLFs (KLF6 and KLF7) are often downregulated while outgrowth-inhibiting KLFs (KLF4 and KLF9) are upregulated in postnatal RGCs [43]. These findings are consistent with the notion that the intrinsic growth capability of neurons is programmed to be attenuated in mammalian adults. It also partially explains why CNS neurons can regenerate robustly in adult Zebrafish in contrast to the poor regenerating performance in mammalian CNS. Additionally, the transcription of some genes, such as GAP-43 and α 1 tubulin, can be differentially regulated during development and in regeneration [44–46]. A recent study has discovered that the fish *gap43* promoter contains fragments specific for regeneration-associated expression [47]. Whether such regenerationspecific regulation of gene expression is a general strategy used by regenerating neurons remains to be examined.

In addition to gene transcription, a few recent studies have elegantly demonstrated an essential role of protein translation mediated by the PI3K/mTOR (Phosphoinositide 3 kinase/mammalian target of rapamycin) pathway in regenerating mouse RGCs (Figure 1D). The PI3K/mTOR pathway targets on subunits of the protein translation machinery and is important for cell growth and survival [48]. In neurons, this pathway is necessary for efficient growth cone regeneration [40]. The mTOR pathway is gradually suppressed during development and completely silenced in injured adult RGCs. PTEN (phosphatase and tensin homolog) and TSC complex (tuberous sclerosis complex consisting of TSC1 and TSC2) are negative regulators of the mTOR pathway. It is suggested that removal of PTEN and TSC1 re-activates mTOR-dependent protein translation [49]. In deed, deletion of PTEN or TSC1 from RGCs in mice leads to enhanced cell survival rate and axon regeneration after optic nerve injury *in vivo* [49]. One of the upstream signaling pathways regulating mTOR in regeneration involves SOCS3 (suppressor of cytokine signaling 3) [50]. SOCS3 deletion releases its suppression on gp130 cytokine receptor which activates the mTOR pathway. Intravitreal application of ciliary neurotrophic factor (CNTF) further enhances axon regeneration in SOCS3-deleted RGCs [50]. It will be intriguing to examine whether the PTEN/mTOR pathway is universally involved in regeneration of PNS and CNS neurons.

The roles of guidance cues and extrinsic inhibitory factors in axon regeneration

Axon guidance cues have long been major interests of regeneration in mature neurons [51]. Correlated expression studies have shown that multiple well-characterized developmental guidance molecules are up- or down- regulated after nerve injury [52,53]. Emerging studies suggest that the effects of the same axon guidance cues may differ during development and in regeneration, possibly through differential use of receptors or activation of redundant pathways. The Slit and Netrin guidance cues are known to primarily act as chemorepellants and chemoattractants respectively to guide the pathfinding of axons during development in various organisms [54]. In adult *C elegans*, AVM mechanosensory neurons in *slt-1*/Slit(lf)

and *unc-6*/Netrin(lf) mutants show better regrowth than controls but with lower precision. In contrast, neither the Slit receptor SAX-3/Robo nor the Netrin receptors UNC-40/DCC and UNC-5/Unc5 exhibit a critical requirement as their ligands in AVM regeneration [3]. Similarly, in rodents, Netrin-1 acts as a chemoattractant during development. However, in axon regeneration, it functions as an oligodendrocyte-associated inhibitor and impedes axon regrowth after adult spinal cord injury [55]. The repulsion-mediating netrin-1 receptor UNC5 seems to be involved in this adult axon function. The Ephrin/Eph signaling has also been shown to regulate regrowth and guidance of regrowing axons in *C. elegans* [4] and in vertebrates [56,57]. The sole Eph receptor VAB-1 in *C. elegans* can facilitate axon regeneration of mechanosensory neurons [4]. In mice, EphB3 supports adult RGC axon outgrowth [57], while Ephrin-B3 acts as a myelin-based inhibitor of axon regrowth in the spinal cord [56]. These findings illustrate the complexity of axon pathfinding in the mature nervous system. It would be of future interests to define the effective phase in which the axon guidance cues act and the downstream effectors they utilize.

It has long been known that myelin-derived inhibitors represent major obstacles in regenerative growth in mammalian CNS post injury. Among them best characterized are the Nogo, myelinassociated glycoprotein (MAG), oligodendrocyte-myelin glycoprotein (OMgp) [23,36,58], and readers are referred to the studies of these inhibitors in mouse models summarized in recent papers [59–62]. Another major class of extracellular inhibitors are chondroitin sulfate proteoglycans (CSPGs). Expression of these CSPGs is dramatically upregulated within the extracellular matrix of scar tissue in the brain and spinal cord of mature animals [23,58]. A breakthrough in the understanding of CSPG action is the recent identification of a transmembrane protein tyrosine phosphatase, PTPσ, as the receptor of neural CSPGs in mice [63]. PTPσ-deleted sensory neurons show reduced sensitivity to CSPGs i*n vitro* and can extend further after a spinal cord lesion *in vivo* [63]. Together with traditional regeneration strategies, genetic approaches will unveil more signaling molecules mediating the environmental guidance and inhibition during axon regeneration in the future.

Perspectives

Genetic model organisms, such as *C. elegans* and *Drosophila*, have been rapidly established as tractable models for dissecting the genetic basis of axon regeneration. Encouragingly, new regeneration-associated pathways identified in invertebrates have shown functional conservation and relevance to those in vertebrates, although the effects of a specific pathway on regeneration may display variations depending on cell types and environments. In addition, increased redundancy and complexity of regeneration-regulating pathways in the vertebrate nervous systems present tremendous challenges to current studies. Manipulating one pathway often could not generate notable impacts on functional recovery of injured neurons, while combinatory treatments in mouse spinal cord injury models have led to cheerful results [64,65]. Thus, investigating synergistic effects of multiple genetic mutations can be a future focus of regeneration research in genetic model organisms. Rich sources of genetic mutants and effective methods of manipulating signaling pathways *in vivo* will no doubt favor such investigations. With elegantly designed large-scale genetic and pharmacological screens, these low-cost genetic model organisms well suit the urgent need of identifying new targets and new chemicals for axon regeneration therapeutics.

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Figure 1.

Representative axon regeneration-regulating pathways discovered in genetic model organisms. (a) In *C. elegans*, axon injury triggers elevation of intracellular Ca^{2+} and cAMP, and activation of the DLK-1 MAPK cascade and PKA kinase. The DLK-1 level is negatively regulated by the E3 ligase RPM-1. The transcriptional factors CEBP-1 and CRH-1/CREB mediates the effects of these kinases respectively and regulate different aspects in axon regeneration. (b) In *Drosophila* adult CNS, the DLK/Wallenda MAPKKK promotes Wallerian degeneration through the JNK signaling. Together with PKA, JNK also promotes axon regrowth in sLNv neurons in the fly brain. (c) In *Drosophila* larvae NMJ, the DLK pathway stabilizes synapses after transient cytoskeletal disruption. In the presence of persistent cellular stress, the JNK phosphatase Puckered (Puc) is also activated and leads to synapse disassembly by antagonizing the JNK signaling. (d) In mouse RGCs, the mTORdependent protein synthesis is essential for neuron survival and axon regrowth after optic nerve injury. Releasing suppression of SOCS3, PTEN, or TSC1/2 on this pathway can reactivate the mTOR signaling and promote axon regeneration.

Table 1

New laser axotomy and regeneration paradigms in *C. elegans*, *Drosophila* and zebrafish

*** Note: An Axon or dendrite is severed at a distance from its soma by a two-photon laser, except in Class I da neurons the axon is completely removed by a UV laser.