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Neural Regeneration in *Caenorhabditis elegans*

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Abstract

Axon regeneration is a medically relevant process that can repair damaged neurons. This review describes current progress in understanding axon regeneration in the model organism *Caenorhabditis elegans*. Factors that regulate axon regeneration in *C. elegans* have broadly similar roles in vertebrate neurons. This means that using *C. elegans* as a tool to leverage discovery is a legitimate strategy for identifying conserved mechanisms of axon regeneration.

Keywords

axon; regeneration; *Caenorhabditis elegans*

INTRODUCTION

Caenorhabditis elegans is a comparatively new model for the study of axon regeneration. Here, we review progress in the field since its beginning in 2004 and discuss directions that future investigations might follow.

The Unrealized Potential of Axon Regeneration

Damaged axons can in some cases regenerate, restoring function to nervous systems after injury or disease. Regeneration of injured neurons is thought to be initiated by signals arising from the injury site (20, 66, 74). These injury signals promote remodeling of the cytoskeleton and plasma membrane at the site of injury, resulting in the generation of a new growth cone. Regulated changes in gene transcription and local and somatic protein synthesis are also required for successful axon extension during regeneration.

The response of a specific neuron to injury is the result of the interplay between various pathways that promote or suppress regeneration. In part, regeneration is regulated by signals from the neuronal environment. For example, myelin-derived signals [Nogo, MAG (myelin-associated glycoprotein), and OMgp (oligodendrocyte myelin glycoprotein)] and chondroitin-sulfate proteoglycans (CSPGs) inhibit regeneration in the vertebrate central nervous system (CNS) (97). However, regeneration is also regulated by intrinsic factors, and altering the intrinsic state of a neuron can improve regeneration, even in the presence of inhibitory factors (58, 59, 65). Thus, building a complete model of the mechanisms that

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regulate and mediate regeneration is critical to understanding how regeneration works and why it often fails.

***Caenorhabditis elegans* as a Regeneration Model**

The nematode worm *C. elegans* has recently emerged as a model for axon regeneration. Axons can be severed through the transparent skin of the worm using a pulsed laser, triggering regeneration (96). Regeneration can then be studied in vivo at single-axon resolution (Figure 1). When this technique is combined with the powerful genetics and other tools available in *C. elegans*, it becomes possible to investigate the complex genetics and cell biology of axon regeneration more rapidly and in greater molecular detail than previously possible (27–29, 36, 57, 92, 95). An important limitation of *C. elegans* for these studies is the relative simplicity of the extrinsic neuronal environment. Worms have no myelin and no invading macrophages. Thus, the worm is a poor model for certain questions regarding the interaction between an injured neuron and its environment. In many cases, however, the simple environment surrounding *C. elegans* neurons can accelerate discovery by removing confounding factors.

An essential question of any biological model is its relevance to human biology. A precise comparison between the mechanisms of axon regeneration in *C. elegans* and those in humans (or at least vertebrate models) is not yet possible because these mechanisms are incompletely understood. However, results so far are encouraging: (a) In *C. elegans*, as in vertebrates, cAMP (cyclic adenosine monophosphate) acts to promote axon regeneration after injury (8, 28, 58); (b) the DLK-1 mitogen-activated protein kinase kinase kinase (MAPKKK) promotes regeneration in both *C. elegans* and mice (36, 43, 95); and (c) axon guidance molecules are important for regeneration in worms and mice (27, 37, 92). Thus, many factors that regulate axon regeneration in *C. elegans* have broadly similar roles in vertebrate neurons. In this review, we describe how *C. elegans* is used to study regeneration and summarize recent work that has begun to describe the mechanisms that regulate and execute regeneration.

INJURY MODELS IN CAENORHABDITIS ELEGANS

A major challenge in studying regeneration is that regeneration only occurs only in response to nerve injury. In mammals, injury is delivered by surgically cutting or crushing a nerve or a section of the spinal cord. These techniques are not easily applied to *C. elegans* because of the small size of the worm. Thus, the study of regeneration in *C. elegans* has required development of two novel injury models: laser axotomy and the β -spectrin mutant background.

Laser Axotomy

In 2004, it was found that laser pulses can sever GABA (γ -aminobutyric acid) neurons in *C. elegans* in intact, living animals and that regeneration can be observed after surgery (96). The first laser used to cut axons in *C. elegans* was a sophisticated, amplified Ti-sapphire laser (96). It is now clear that many varieties of pulsed laser can cut axons and elicit regeneration, including unamplified Ti-sapphire lasers (92), coumarin dye lasers (92), Nd:YAG (neodymium-doped yttrium aluminum garnet) lasers (68), and solidstate lasers (89). In addition to the GABA motor neurons, laser axotomy has been used to sever many other neurons, including the PHA phasmid sensory neurons (92), the PLM and ALM touch neurons (92), the sensory neuron AFD (18), the AWB chemosensory neuron (92), the AVM mechanosensory neuron (27), the DA/DB motor neurons (27), and the HSN neuron (27). There are no examples of neurons that cannot be cut, suggesting that laser axotomy can be used on any neuron in the worm.

Pulsed lasers sever axons by generating free electrons, by causing the formation (and collapse) of nanoscale bubbles, and by generating free plasma that results in a cavitation bubble (10). The precision of laser axotomy enables very accurate and controlled injuries at the single-neuron level. For example, Wu and colleagues (92) found that the success of regeneration in the ALM and PLM neurons depends on how close to the cell body the injury occurs: Injury close to the cell body results in increased regeneration. Similarly, Gabel and colleagues (27) showed that in the DA/DB motor neurons cutting closer than 20 μm to the cell body elicits growth from the cell body rather than the axon stump. A potential limitation of laser axotomy is the requirement to individually target and sever each individual axon that is to be studied. However, diligent application of this technique has yielded nearly all current data, including one study that analyzed hundreds of genotypes (15). One way to increase the speed of laser axotomy is by using a microfluidic platform to immobilize worms, combined with an automated axotomy system (3, 32, 73). A microfluidic and automated axotomy system has been used to screen for chemical modifiers of regeneration (75). Such systems hold the promise of significantly increasing experimental throughput.

β -Spectrin Mutants

β -spectrin is a major and essential component of the membrane skeleton, a dense two-dimensional protein mesh that is tightly associated with the plasma membrane in most metazoan cells (6, 7). In *C. elegans*, β -spectrin is encoded by the *unc-70* gene (34). Neurons in animals lacking *unc-70*/ β -spectrin develop with normal morphology. However, these neurons are fragile, and their axons break spontaneously when the mutant animals hatch and start to crawl (35). Broken axons regenerate, so regeneration can be studied in *unc-70*/ β -spectrin mutant animals without the requirement for experimentally delivering nerve injuries.

Axon breaks in *unc-70*/ β -spectrin mutant animals occur stochastically and may even recur in the same axon (35). Thus, it is more difficult to collect quantitative single-neuron data on regeneration using the *unc-70*/ β -spectrin mutant background than using laser axotomy. However, because regeneration occurs spontaneously and constitutively, it is far easier to collect many animals that are regenerating. The *unc-70*/ β -spectrin mutant background therefore may have advantages over laser surgery for experiments that require very large numbers of animals, such as genetic screens and genomic and proteomic studies. For example, this sensitized mutant background was successfully used in an unbiased RNA interference (RNAi) screen for genes affecting regeneration (36).

Differences Between Injury Models

An important and unresolved question is the extent to which different injury models—the use of different types of lasers, the study of different neurons, or the spontaneous breaks in *unc-70*/ β -spectrin mutants—converge on common regeneration mechanisms. Detailed studies using femtosecond lasers to sever *C. elegans* axons show that dramatic changes in regeneration occur as a result of changing the number and energy of laser pulses, even though the total amount of energy applied is constant (10). Even the choice of fluorescent marker can influence regeneration (92). In some cases, such as the *dlk-1* mitogen-activated protein kinase (MAPK) pathway, regeneration mechanisms show consistent effects across multiple neuron types, markers, and even injury models (36, 95). In other cases, however, the extent to which mechanisms remain consistent across different cells, markers, and injury models remains to be determined.

INTRINSIC REGENERATION PATHWAYS

Regeneration is regulated by intrinsic factors, and altering the intrinsic state of a neuron can improve regeneration, even in the presence of an inhibitory environment (58, 59, 64). Intrinsic pathways can also inhibit regeneration (17, 63, 81, 84). Both pro- and antiregeneration pathways have been described in *C. elegans* and are reviewed in this section.

Calcium/cAMP/PKA

Calcium and cAMP (cyclic adenosine monophosphate) signaling promote axon regeneration in a variety of systems (4, 13, 102). Increased calcium influx as a result of injury can result from entry of extracellular calcium through the breached axonal membrane, activation of voltage-gated calcium channels in response to depolarization, and release from intracellular stores (47, 101). Although increased intracellular calcium can result in neuronal cell death in some cases, calcium signaling can promote membrane resealing and regeneration in surviving neurons (71, 93). Increased intracellular calcium triggers multiple signaling events, including an increase in cAMP through modulation of adenylyl cyclase activity (19).

Calcium and cAMP also promote axon regeneration in *C. elegans* (28). Axotomy of the PLM touch neuron axons results in an acute increase in intracellular calcium. This increase is at least partially dependent on the voltage-gated calcium channel subunit EGL-19 and the inositol triphosphate receptor ITR-1. Blocking the function of EGL-19 or ITR-1 reduces the acute increase in intracellular calcium and also reduces regeneration. Similarly, differences in the acute calcium increase in wild-type animals correlate with differences in the length of axon regeneration. Thus, intracellular calcium entry promotes regeneration in *C. elegans*. A major effector of calcium in regeneration is the adenylyl cyclase-cAMP-protein kinase A (PKA) signaling cassette. Interestingly, BAPTA (1,2-bis(o-aminophenoxy)ethane N, N, N', N' tetraacetic acid) treatment reduces the calcium baseline and reduces regeneration without affecting the size of the calcium transient. This suggests that total levels of calcium, rather than the relative size of the increase, are the final determinant of regeneration.

Channels, Transporters, and Neurotransmitters

Increased neuronal excitability results in more intracellular calcium, whereas decreased excitability results in less (reviewed above) (28). Consistent with the finding that calcium mediates regeneration, mutations that decrease excitability result in poor regrowth after axotomy, including the sodium pump *nkb-1*, the sodium channel *unc-8*, and the stomatins *unc-1* and *unc-24*. Similarly, mutations that disrupt the synthesis of acetylcholine (*cha-1*) or acetylcholine transport (*unc-17*), or that disrupt the *deg-3* subunit of the nicotinic acetylcholine receptor, result in decreased regeneration (15). Conversely, genetic lesions that make neurons more excitable, such as loss-of-function mutations affecting the *slo-1* K⁺ channel or the K⁺ channel regulator *mpps-1*, enhance regrowth (15).

MAP Kinase Signaling

The *dlk-1* MAPK regulates regeneration in both GABA motor neurons and sensory neurons (36, 95). An unbiased RNAi screen in *unc-70*β-spectrin mutants for genes affecting regeneration identified *dlk-1* as a candidate regeneration gene (36). *dlk-1* is a MAPKKK that functions in a pathway with the MAP kinase kinase (MAPKK) *mkk-4* and the p38 MAPK *pmk-3* (56) (Figure 2). A major target of *pmk-3* is the MAPKAPK (MAP kinase activated kinase) *mak-2*, which functions via the bZip-containing protein *cebp-1* (95). This entire signaling pathway is required for regeneration. Loss of any component reduces regeneration, whereas activation of the pathway (by overexpression of *dlk-1*) increases regeneration (36, 95).

The same RNAi screen in *unc-70* β -spectrin mutants identified a second MAPKKK, *mlk-1*, as also having a significant role in regeneration (36). *mlk-1* activates the MAPKK *mek-1*, which acts upstream of the JNK (c-Jun N-terminal kinase)-like kinase *kgb-1* (52) (Figure 2). Again, this entire signaling pathway functions in regeneration. Further, the *mlk-1* pathway exhibits significant cross talk with the *dlk-1* pathway. In addition, the *dlk-1* and *mlk-1* MAPK pathways share common negative regulators. *rpm-1*, an E3 ubiquitin ligase that regulates synapse morphology and axon termination (30, 77, 100), mediates the degradation of *mlk-1* as well as *dlk-1* (56, 60). Hyperactivation of one or both of these pathways in *rpm-1* mutants leads to increased regeneration (36). Both the *dlk-1* and *mlk-1* MAPK pathways are also regulated by the dual phosphatase *vhp-1* (60).

In addition to these common regulators, the *mlk-1* pathway is regulated by growth factor signaling. *svh-1* is homologous to hepatocyte growth factor, macrophage stimulating protein, and plasminogen. *svh-1* is a secreted factor that acts via the receptor-type tyrosine kinase *svh-2* to activate regeneration via activation of the *mlk-1* pathway. Although *svh-2* functions in the injured neuron, *svh-1* is expressed in the ADL neurons and functions as an extrinsic signal. Thus, the response of injured neurons is regulated in part by secreted factors deriving from other, uninjured neurons.

How does MAPK signaling facilitate regeneration? Activation of the *dlk-1* pathway results in the stabilization of the *cebp-1* messenger RNA (mRNA) through its 3' untranslated region. Some mammalian MAPKs (homologous to *mak-2*) can phosphorylate mRNA binding proteins, which may result in differential regulation of mRNA turnover (40). Thus, *mak-2* may mediate *cebp-1* mRNA stability by modifying RNA-binding proteins in response to axotomy. In the end, via the *dlk-1* and *mlk-1* pathways, axotomy results in localized translation of *cebp-1* and also activation of *kgb-1*. However, the downstream effectors of *cebp-1* are not known, nor are the effectors of *kgb-1*.

***efa-6* and Microtubule Dynamics**

Microtubules play a crucial role in axon regeneration. Destabilization of microtubules with nocodazole in regenerating rat sciatic nerve inhibits growth and transforms growth cones into retraction bulbs. Conversely, chemical stabilization of microtubules in the rat spinal cord by taxol treatment reduces the formation of retraction bulbs and enhances growth-cone formation. Retraction bulbs exhibit a disorganized microtubule structure and are static, whereas growth cones show a characteristic bundling of microtubules essential for growth-cone movement and axonal regrowth (22). Manipulation of microtubules results in enhanced regeneration in the mammalian CNS and may present novel therapeutic possibilities (39, 78, 82).

Microtubule dynamics are also important for *C. elegans* axon regeneration. *efa-6* mutations result in increased PLM regrowth after axotomy (15). EFA-6 contains a variable N-terminal domain, a pleckstrin homology (PH) domain, and a Sec7 domain, which has a guanine exchange factor activity for ADPribosylation factor GTPases (26, 61). However, neither the PH domain nor the Sec7 domain of EFA-6 is required for inhibition of regeneration. Rather, the N terminus of EFA-6 inhibits regeneration by regulating microtubule organization. Although loss of *efa-6* results in increased regeneration, overexpressing the N terminus of EFA-6 results in dysmorphic growth cones, less regeneration, and a reduction in dynamic microtubules (15). Stabilizing microtubules with taxol suppresses the inhibition of regeneration resulting from EFA-6 overexpression. This suggests that EFA-6 destabilizes microtubules in regenerating PLM axons and leads to failure of regrowth.

Notch Signaling

The *lin-12* Notch receptor is a potent inhibitor of regeneration in *C. elegans* motor neurons (21). *lin-12* Notch null animals regenerate significantly better than wild-type animals, whereas *lin-12* gain-of-function alleles regenerate worse. Notch functions in the injured neuron to inhibit regeneration. Further, Notch activation after injury is necessary for Notch to inhibit regeneration: Blocking Notch activation with the γ -secretase inhibitor DAPT (N-[(3, 5-difluorophenyl)acetyl]-L-alanyl-2-phenylglycine-1,1-dimethylethyl ester) increases regeneration, even when injury has already occurred.

Notch activation typically involves sequential cleavage of Notch, first by a metalloprotease (site 2 cleavage) and then by the γ -secretase complex (site 3 cleavage). These cleavages release the Notch intracellular domain (NICD) into the cytoplasm. Notch inhibition of regeneration proceeds by this same activation mechanism: Both the *sup-17*/ADAM10 metalloprotease and presenilin (the catalytic component of γ -secretase) are required for Notch to inhibit regeneration, and NICD overexpression is sufficient to inhibit regeneration.

The activation mechanism for Notch inhibition of regeneration is not known. DSL (Delta/Serrate/LAG-2)-family Notch ligands are single-pass transmembrane proteins (87, 98). In addition to transmembrane DSL ligands, *C. elegans* expresses multiple secreted ligands thought to activate the receptor in cooperation with transmembrane ligands (16, 25, 45). However, loss of any single ligand does not improve regeneration. One possibility is that multiple ligands can redundantly activate Notch to inhibit regeneration. Alternatively, Notch activation might be ligand independent. In *Drosophila*, endocytosis of the Notch receptor can be mediated through Deltex in a ligand-independent manner and stabilized by Hif- α (41, 42, 55, 88, 94). Once endocytosed, Notch is cleaved by presenilin, and the NICD can be released into the cytoplasm. Calcium-dependent, ligand-independent Notch activation has also been described (67).

Notch activation is required within a short time frame (2 h) of axotomy to inhibit regeneration. This limited window of activation suggests that Notch may be directly activated by injury and that it signals immediately to inhibit regeneration. This immediate activation would allow enough time for an inhibitor of regeneration to be transcribed and translated to inhibit growth-cone initiation. Alternatively, Notch may be active at all times but may need to interact with other pathways that respond directly to injury or may need to be active continuously because its effectors are short lived. How Notch signaling affects regeneration remains an open question. One possibility that is consistent with our findings is that Notch could act by activating transcription of a direct inhibitor of regeneration. Some targets of Notch signaling have been identified, but their role in regeneration has not been examined.

Synaptic Vesicle Recycling

Synaptic vesicle exocytosis at nerve terminals results in the transfer of vesicle proteins and lipids to the plasma membrane; these are retrieved by endocytosis. Endocytosis— but not exocytosis—is also required for regeneration (15). Mutations in any of three key endocytosis genes (*unc-26*/synaptojanin, *unc-57*/endophilin, and *unc-41*/stonin) result in decreased regeneration. The requirement for synaptic vesicle endocytosis (but not exocytosis) genes in regeneration suggests that regeneration does not depend on synaptic vesicle cycling but on different cellular process. One potential mechanism is MAPK signaling, as the requirement for *unc-57*/endophilin during PLM regeneration can be bypassed by increased DLK-1 expression (15).

An Emerging Role for MicroRNAs

MicroRNAs function in many aspects of neural development and disease (46). Studies in cultured neurons, mice, and zebrafish suggest a role for microRNAs in central and peripheral regeneration (83, 91, 99). Because microRNAs are generally involved in the downregulation of their target genes, a regeneration-promoting role of microRNAs suggests an important role for the negative regulation of inhibitors of regeneration. In *C. elegans*, the Argonaute homolog *alg-1* is required for PLM axon regeneration (15). *alg-1* functions in the biogenesis of microRNAs (31), suggesting a role for microRNAs in regulating regeneration in *C. elegans*. Although a general regeneration-promoting role for microRNAs has recently emerged, further study of the roles of individual microRNAs may reveal specific roles.

EXTRINSIC REGULATION OF REGENERATION IN *CAENORHABDITIS ELEGANS*

Regenerating axons navigate a very different environment than the one they encounter during development. Although some developmental growth and guidance cues may be maintained in adults, others are lost or replaced by inhibitory factors. Numerous studies in mammalian systems show that extrinsic factors from the glial scar and myelin act as potent inhibitors of regeneration in the CNS (reviewed in 11, 24, 50, 79, 97). *C. elegans* axons are not myelinated and navigate a very different extracellular environment than mammalian neurons. However, recent evidence suggests that axon guidance factors and elements of the *C. elegans* extracellular matrix (ECM) also play a role in the regulation of axon regeneration in worms.

Axon Guidance Factors

Regeneration in *C. elegans* is often characterized by guidance errors, premature termination, or branching (27, 92). By contrast, development of the nervous system is largely invariant. One mechanistic difference between regeneration and development is the function of axon guidance pathways. During development, ventral migration of the ALM mechanosensory neuron depends on the netrin and the Slit/Robo guidance pathways (27). These pathways depend on extracellular ligands (netrin and Slit) and their neuronal receptors (the netrin receptors *unc-5* and *unc-40* and the Slit receptor Robo). During regeneration, netrin and Slit also help to guide ALM ventral migration: Migration is defective in mutants for these genes. However, the neuronal receptors are dispensable for this process (27). Thus, axon guidance signals function differently in ALM during regeneration than during development.

A change in response to axon guidance molecules is also found in the PLM neurons. In these neurons, Slit/Robo signaling promotes outgrowth (49). During regeneration, however, Slit/Robo signaling inhibits axon extension (15). Further, accurate regeneration in the PLM neuron in adult animals is inhibited by the *vab-1* Eph receptor (92). However, this function is independent of the kinase activity of *vab-1* (92), whereas kinase activity is required for *vab-1*'s developmental function (53). Thus, differences in how axon guidance pathways function may account for some of the pathfinding errors during regeneration.

Extracellular Matrix

Many axons in *C. elegans* migrate along the basement membrane that runs between the body wall muscles and the epidermis. This basement membrane is required to maintain attachment of the muscle to the epidermis. Three genes—*pxn-2*, *spon-1*, and *vab-19*—that inhibit regeneration encode proteins that function to maintain muscle attachment. PXN-2 is a peroxidase, which is a secreted protein that contains a catalytic peroxidase domain, immunoglobulin domains, and leucine-rich repeats. Peroxidases can inactivate metalloproteases, cross-link ECM components, and mediate cell-ECM adhesion (38, 70, 86).

SPON-1 is spondin, a component of the ECM (44). VAB-19 is an ankyrin repeat protein, homologous to the human tumor suppressor Kank (76). VAB-19 is a cytoplasmic protein that localizes with components of epidermal attachment structures. Loss of any of these three genes causes defects in attachment of muscle to epidermis (29, 76, 90) and also results in increased regeneration (15, 29). These data suggest that muscle-epidermis attachment via the basement membrane inhibits regeneration. ECM proteins could act as a physical barrier to regeneration or could sequester essential soluble factors, rendering them unavailable to regrowing neurons. Alternatively, there may be receptor-mediated inhibitory mechanisms in *C. elegans* that are similar to inhibition of regeneration in vertebrates by CSPGs.

FUSION AND FUNCTIONAL REGENERATION

C. elegans neurons are capable of reconnecting directly to the severed distal fragment (14, 28, 57). Fusion restores membrane and cytoplasmic continuity between the two halves of the axon (28, 57). Fusion can also prevent degeneration of the distal fragment but only if it occurs relatively quickly after axotomy (57). If regenerating axons are given a choice of two distal fragments, one from the regenerating axon and one from another neuron, fusion occurs nearly exclusively with the correct fragment (57). Thus, reconnection after fusion can restore neuronal circuitry, at least in terms of morphology.

Fusion after axotomy requires the fusogen *eff-1* (28). *eff-1* also functions in homotypic fusions during normal development of epithelia, muscles, and sensory dendrites (54, 62). Fusion is also promoted by calcium and cAMP via PKA (28). Thus, fusion after axotomy is mediated by some of the same mechanisms as axon regeneration.

What of functional recovery? Regenerating axons in vertebrates can form de novo synapses and restore function providing they reach their target (1, 80). In *C. elegans*, loss of GABA neuron function (for example, in mutants that cannot synthesize GABA) leads to a characteristic shrinker behavioral phenotype (51). Severing all GABA commissures also results in the shrinker phenotype, and regeneration can restore GABA-mediated behavior (21, 96). Further, reduction of GABA regeneration (by increasing Notch signaling) also reduces behavioral recovery (21). Thus, regeneration can restore function to damaged neural circuits.

PERSPECTIVES

In a relatively short time, *C. elegans* has emerged as a highly tractable model to study axon regeneration. Current data suggest that many genes function similarly in worm and mammalian regenerating axons (see Introduction above). *C. elegans* offers the axon regeneration field a simple, highly invariant nervous system, genetic tractability, and the ability to study individual neurons in vivo. Although many novel signaling pathways that function in regeneration have been identified, many questions remain.

What Activates Regeneration Pathways?

MAPK pathways (48, 95) and probably also Notch signaling (21) are activated by injury. The cellular mechanism that links injury to activation is currently unknown. A variety of injury signaling mechanisms have been proposed, including calcium entry, electrical signals, and changes in trafficking of a regeneration factor. Of these, only calcium is known to function in *C. elegans* regeneration (28). One possibility is that calcium signaling is the single cue that activates all of the acute responses to injury. For example, modulation of calcium in vivo and at physiological conditions can activate Notch (69). Alternatively, other injury signals may exist. For example, disruption of the microtubule cytoskeleton in nerve injury may be an injury signal. In support of this idea, microtubule depolymerization can

activate *dlk-1* signaling (9). However, it remains to be determined whether this activation mechanism occurs in regenerating axons.

Answers to these questions and others may come via genetic screens. So far, two screens (an unbiased RNAi screen in *unc-70*β-spectrin mutants and systematic screening of existing mutant alleles using laser axotomy) identified many genes involved in regeneration (15, 36). Additional genetic screens are likely to provide more details on the function of known pathways and also to identify additional factors that mediate regeneration.

What Are the Effectors of Regeneration Pathways?

In the end, regeneration signaling must converge on growth mechanisms that enable the injured neuron to generate a new growth cone, maneuver the growth cone to its target, and reconnect. These growth mechanisms are for the most part not understood in *C. elegans*. However, the recent description of microtubule dynamics in regenerating axons and the discovery of a novel regulator of these dynamics (see section on *efa-6* and Microtubule Dynamics) suggest that it will be possible to analyze regeneration at the cell-biological level. Further application of cell-biological techniques, such as electron microscopy and super-resolution imaging, in combination with genetic analysis, should result in a better understanding of the growth mechanisms that mediate regeneration.

A second approach to cell biology that also has potential translational applications is the use of *C. elegans* as a platform to screen for drugs that are effective in improving regeneration after injury. A high-throughput screen using microfluidics identified a chemical enhancer of regeneration (75). Further, microinjecting compounds directly into the pseudocoelom of animals immediately after axotomy can enhance regeneration (15, 21). This technique may be used in the future to assess limited numbers of compounds for effects on regeneration. Such compounds may identify particular cell-biological processes that are important for regeneration.

How Good is Regeneration?

Most regeneration studies in *C. elegans* have used the morphology of the regenerating neuron as a measure of regenerative success. To date, two studies in the GABA motor neurons have shown that regeneration is accompanied by functional recovery at the level of whole-animal behavior (see section on Fusion and Functional Regeneration). However, such experiments are limited in their ability to accurately assess the function of individual regenerated neurons compared with their uninjured counterparts. *C. elegans* is a tractable model for a more detailed study of neuronal function using electrophysiological and optogenetic techniques (23, 33, 72). The future application of these techniques to the study of regeneration will yield information about how effective functional recovery is and may identify new pathways that are required for functional, rather than merely morphological, regeneration.

Why Inhibit Regeneration?

Axon regeneration can restore function, so why inhibit it? In the vertebrate CNS, pathways that inhibit regeneration also function in uninjured nervous systems and help maintain a stable, functional system by inhibiting aberrant growth and plasticity (2, 97). For example, CSPGs inhibit regeneration after injury in the CNS (11, 24, 50, 79, 97). In uninjured animals, enzymatic degradation of CSPGs results in ectopic growth and sprouting (5, 12, 20). These data suggest that inhibition of the injury response in the CNS is part of a broader program to limit plasticity. Similarly, it is possible that the *C. elegans* pathways that inhibit regeneration also function to stabilize the mature, uninjured nervous system.

Neurons in old *C. elegans* animals show a loss of stability and accumulate ectopic branches (85). Loss of *jnk-1* causes an increase in ectopic neuronal branching in old animals, suggesting that *jnk-1* contributes to nervous system stability (85). Consistent with the idea that stability pathways can also inhibit regeneration, *jnk-1* mutants also display increased regeneration after nerve injury (36). In other ways, however, stability and regeneration seem to be separate processes. The *dlk-1* pathway is required for regeneration but not for spontaneous branching in old animals. Loss of *mlk-1* increases the incidence of age-dependent branching but decreases regeneration (36, 60, 85, 95). This suggests that in *C. elegans*, regeneration and spontaneous branching are promoted by different mechanisms. It remains to be seen whether other inhibitors of regeneration, such as Notch (21) and *efa-6* (15), affect spontaneous branching and the overall stability of the nervous system.

Why Can Worms Regenerate at All?

Neurons in *C. elegans* can regenerate in response to injury. Is this ability due to a specialized regeneration mechanism that has evolved specifically to respond to nerve injury? Or is the response to injury part of a more general mechanism, such as homeostasis or stress response? Or is it even a pathological and undirected response to trauma?

During their life span in the wild, worms are subjected to desiccation, mechanical trauma, and predators. Thus, the ability to regenerate neurons quickly after they have been severed and to regain movement may pose a significant selective advantage. Alternatively, neuronal regeneration may be a particular manifestation of some broader biological process. A more complete understanding of the mechanisms that mediate regeneration might eventually help answer this question.

Conclusion

C. elegans has advanced the field of axon regeneration by providing a genetic system that is compatible with both high-throughput screening and single-neuron analysis. Additional forward genetics, a wider use of high-throughput techniques, functional regeneration assays, and drug validation will make *C. elegans* an even more robust model and further advance the study of axon regeneration.

Glossary

MAG	myelin-associated glycoprotein
Omgp	oligodendrocyte myelin glycoprotein
CSPG	chondroitin-sulfate proteoglycan
CNS	central nervous system
cAMP	cyclic adenosine monophosphate
MAPKKK	mitogen-activated protein kinase kinase kinase
GABA	γ -aminobutyric acid
RNAi	RNA interference
MAPK	mitogen-activated protein kinase
PKA	protein kinase A
BAPTA	1,2-bis(o-aminophenoxy)ethane N, N, N', N', tetraacetic acid
MAPKK	mitogen-activated protein kinase kinase

N-[(3, 5-difluorophenyl)-acetyl]-L-alanyl-2-phenylglycine-1,1-dimethylethyl ester small (DAPT)	molecule inhibitor of γ -secretase
NICD	Notch intracellular domain
DSL	Delta/Serrate/LAG-2
ECM	extracellular matrix

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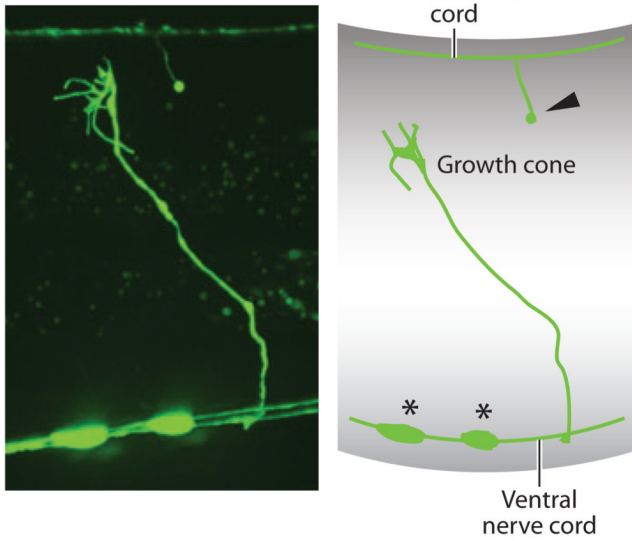
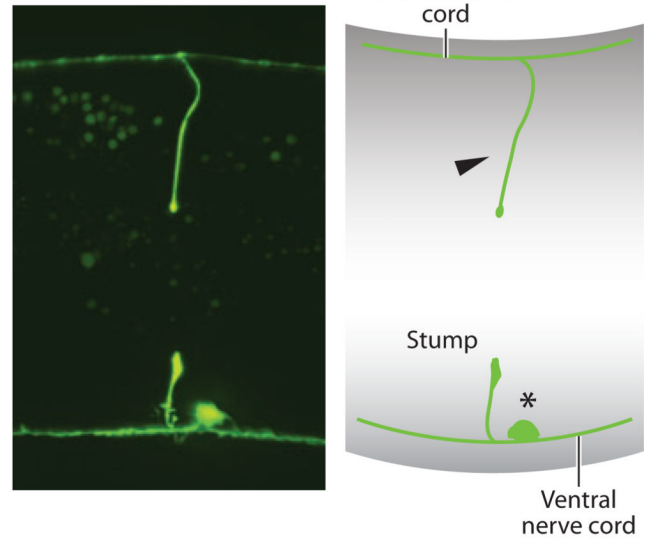
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a Regeneration**b** Failed regeneration**Figure 1.**

(a) A regenerating *Caenorhabditis elegans* GABA (γ -aminobutyric acid) neuron after laser surgery, and (b) a neuron that has failed to respond to injury (21).

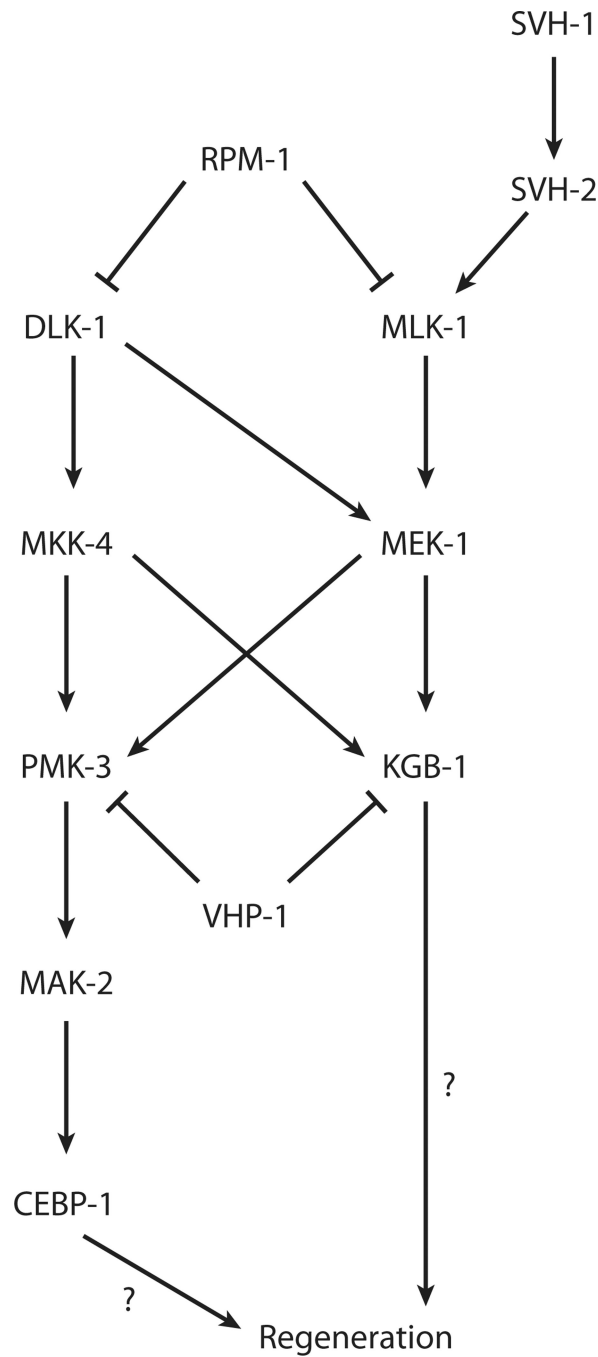


Figure 2. Diagram of mitogen-activated protein kinase signaling during regeneration in *Caenorhabditis elegans*. Adapted from 48, 60.