

Axonal Degeneration Is Mediated by Necroptosis Activation

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Axonal degeneration, which contributes to functional impairment in several disorders of the nervous system, is an important target for neuroprotection. Several individual factors and subcellular events have been implicated in axonal degeneration, but researchers have so far been unable to identify an integrative signaling pathway activating this self-destructive process. Through pharmacological and genetic approaches, we tested whether necroptosis, a regulated cell-death mechanism implicated in the pathogenesis of several neurodegenerative diseases, is involved in axonal degeneration. Pharmacological inhibition of the necroptotic kinase RIPK1 using necrostatin-1 strongly delayed axonal degeneration in the peripheral nervous system and CNS of wild-type mice of either sex and protected *in vitro* sensory axons from degeneration after mechanical and toxic insults. These effects were also observed after genetic knock-down of *RIPK3*, a second key regulator of necroptosis, and the downstream effector *MLKL* (*Mixed Lineage Kinase Domain-Like*). RIPK1 inhibition prevented mitochondrial fragmentation *in vitro* and *in vivo*, a typical feature of necrotic death, and inhibition of mitochondrial fission by Mdivi also resulted in reduced axonal loss in damaged nerves. Furthermore, electrophysiological analysis demonstrated that inhibition of necroptosis delays not only the morphological degeneration of axons, but also the loss of their electrophysiological function after nerve injury. Activation of the necroptotic pathway early during injury-induced axonal degeneration was made evident by increased phosphorylation of the downstream effector MLKL. Our results demonstrate that axonal degeneration proceeds by necroptosis, thus defining a novel mechanistic framework in the axonal degenerative cascade for therapeutic interventions in a wide variety of conditions that lead to neuronal loss and functional impairment.

Key words: axonal degeneration; mitochondrial fragmentation; MLKL; necroptosis; neurodegeneration; RIP kinase

Significance Statement

We show that axonal degeneration triggered by diverse stimuli is mediated by the activation of the necroptotic programmed cell-death program by a cell-autonomous mechanism. This work represents a critical advance for the field since it identifies a defined degenerative pathway involved in axonal degeneration in both the peripheral nervous system and the CNS, a process that has been proposed as an early event in several neurodegenerative conditions and a major contributor to neuronal death. The identification of necroptosis as a key mechanism for axonal degeneration is an important step toward the development of novel therapeutic strategies for nervous-system disorders, particularly those related to chemotherapy-induced peripheral neuropathies or CNS diseases in which axonal degeneration is a common factor.

Introduction

Neurons can extend meter-long axons to carry out their physiological functions (Debanne et al., 2011). However, this architec-

ture also makes axons highly susceptible to damage. Axonal degeneration is a common and early feature of many neurological disorders, including amyotrophic lateral sclerosis (ALS), Parkinson's disease (PD), and Alzheimer's disease (AD). Axonal degeneration also commonly occurs after traumatic or chemical insults (Coleman, 2005; Benarroch, 2015; Gerds et al., 2016).

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The degenerative process of mechanically isolated axons, known as Wallerian degeneration (Waller, 1850), has been broadly used as a model to unveil the cellular and molecular mechanisms of axonal degeneration (Coleman and Freeman, 2010; Conforti et al., 2014). Likewise, chemotherapeutic agents have also been used to model chemotherapy-induced peripheral neuropathies, a serious and common side effect of cancer treatment (Taillibert et al., 2016). Mechanistic studies in transgenic models displaying delayed Wallerian degeneration (Calixto et al., 2012; Osterloh et al., 2012; Di Stefano et al., 2017) have shown evidence of a regulated axonal self-destruction program (Court and Coleman, 2012; J.T. Wang et al., 2012). The expression of the *Wld^s* protein is the most studied model of delayed axonal degeneration (Mack et al., 2001; Adalbert et al., 2005; Hoopfer et al., 2006). Loss-of-function studies have also identified mutants that exhibit *Wld^s*-like protection of severed axons, such as *dSarm/Sarm1* (Osterloh et al., 2012; Essuman et al., 2017) and *Axundead* (Neukomm et al., 2017), supporting the idea that Wallerian degeneration is indeed an active and regulated process.

Although axonal degeneration is apoptosis-independent (Finn et al., 2000; Whitmore et al., 2003), both axonal degeneration and apoptosis exhibit some related steps, including a latent phase, where cell death can be delayed, and a subsequent execution phase of cellular breakdown (Geden and Deshmukh, 2016). Mitochondrial dysfunction, by activation of the mitochondrial transition pore (mPTP), calcium imbalance, and reactive oxygen species (ROS) increase, are key events for axonal degeneration (Barrientos et al., 2011; Calixto et al., 2012; Villegas et al., 2014). Mitochondrial fragmentation is also present in diseases in which axon degeneration is a prominent feature (Knott et al., 2008; Court and Coleman, 2012). These events connecting mitochondrial dysfunction with axonal degeneration are also shared features of programmed necrotic cell death, currently known as necroptosis.

Necroptosis is characterized by cellular and organelle swelling, plasma membrane rupture, and the release of cellular contents, inducing a proinflammatory response (Vandenabeele et al., 2010; Orozco and Oberst, 2017). Both apoptosis or necroptosis can be triggered upon TNF α -receptor activation (Degterev et al., 2005). Normally, TNF α induces apoptosis through caspase-8 activation, which in turn inhibits necroptosis by RIPK1 and RIPK3 cleavage (Sun and Wang, 2014). Under caspase-8 inhibition, both kinases are phosphorylated, triggering phosphorylation and oligomerization of MLKL (Mixed Lineage Kinase Domain-Like) and membrane permeabilization (Cai et al., 2014; Tummers and Green, 2017). Other necroptotic activators have been recently described, including increased intracellular Ca²⁺ (Nomura et al., 2014; Wallach et al., 2016) and ROS (Zhang et al., 2017; Yang et al., 2018).

Necroptosis participates in nervous-system pathologies with prominent axonal degeneration, such as multiple sclerosis (Ofengeim et al., 2015), ALS (Re et al., 2014; Ito et al., 2016), PD (Iannielli et al., 2018), AD (Caccamo et al., 2017), and neuroinflammation (Bian et al., 2017; Daniels et al., 2017), among others (Huang et al., 2013; Vieira et al., 2014). This research has established that there are both neuronal-autonomous roles for necroptosis and noncell autonomous mechanisms participating in nervous-system dysfunction (Tonnus and Linkermann, 2017). However, no one has determined whether necroptosis is activated in a specific neuronal compartment, such as the axon.

Here, we demonstrate that pharmacological inhibition and genetic knock-down of central necroptotic players strongly delayed morphological and functional loss of axons in both pe-

ripheral nervous system (PNS) and CNS models of Wallerian degeneration. Furthermore, we observed that, after injury, necroptosis is activated early in axons, followed by mitochondrial fragmentation and progressive degeneration of axons. Inhibition of mitochondrial fission delayed axonal degeneration, supporting the idea that mitochondrial fragmentation is a crucial axon-degeneration effector after necroptosis activation. Together, our results identify necroptosis as the programmed mechanism for axonal degeneration, revealing novel molecular targets for neuroprotective therapies in which axonal degeneration is a common factor.

Materials and Methods

Animals. Wild-type (WT) C57BL/6J [International Mouse Strain Resource (IMSR) catalog #JAX:000664, RRID:IMSR_JAX:000664] adult mice (25 g) of either sex and Sprague Dawley [Mouse Genome Informatics (MGI) catalog #5651135, RRID:MGI:5651135] pregnant rats were obtained from the animal facilities of Pontificia Universidad Católica de Chile and Universidad Mayor. Thy1-YFP mice (IMSR catalog #JAX:003782, RRID:IMSR_JAX:003782), of either sex, were purchased from the Jackson Laboratory and maintained in the Universidad Mayor animal facility. Experiments with animals followed protocols approved by the institutional animal care and use committees and complied with National Institutes of Health guidelines.

Nerve explant cultures. To study axonal degeneration *ex vivo*, sciatic and optic nerve segments of 10 and 4 mm, respectively, were dissected from adult WT mice ($n = 3$) and cultured in 24-well dishes containing 400 μ l of Neurobasal medium (Invitrogen, 21103-049), 2% B27 (Invitrogen, 17504-044), 0.3% L-glutamine (Invitrogen, 25030-081), and 1% streptomycin/penicillin. Explants were cultured at 37°C and 5% CO₂ over different periods of time and thereafter fixed with 4% paraformaldehyde in 1 \times PBS for 1 h, followed by three 10 min washes in 1 \times PBS.

Sciatic nerve crush. Mice ($n = 3$) were intraperitoneally anesthetized with Avertin (0.37 mg/g; Sigma-Aldrich, T48402) and the sciatic nerve was exposed at the mid thigh and crushed three times for 5 s each using Dumont #5 forceps. The crushed site was marked with graphite powder applied in the forceps, and the wound was closed using surgical clips. The sciatic nerve was intraperitoneally injected with necrostatin-1 (Nec-1; 1.65 mg/kg; Sigma-Aldrich, N9037) or its vehicle (3% DMSO in PBS) before crushing as described by Catenaccio et al. (2017).

Teased nerve fibers. Sciatic nerve explants, obtained from Thy1-YFP mice ($n = 3$), were removed from culture medium and immersed in a 1.5 ml tube with 4% paraformaldehyde in 1 \times PBS for 1 h at room temperature (RT). Then, nerves were washed three times for 10 min in 1 \times PBS and transferred into a 35 mm culture dish with 1 \times PBS to remove the perineurium. The bundles were separated with acupuncture needles. Each bundle was placed in a drop of 1 \times PBS on a slide coated with 3-aminopropyltriethoxy silane. The fibers were teased and attached to the slides. After the drops dried, the slides were immersed in cold 100% acetone for 20 min at -20°C to proceed with the immunofluorescence protocol detailed below.

Dorsal root ganglion cultures. Briefly, embryonic day 16 rat embryos were decapitated, and the limbs and organs were removed. The spinal cord with dorsal root ganglia (DRGs) was dissected and placed in a Petri dish containing cold L-15 medium (Invitrogen, 11415-064). For DRG explants, complete DRGs were cultured in 24-well dishes containing 400 μ l of Neurobasal medium, 2% B27, 0.3% L-glutamine, 1% streptomycin/penicillin, 4 μ M aphidicolin (Sigma-Aldrich, A0781), 7.5 μ g/ml 5-fluoro-2-deoxyuridine (Sigma-Aldrich, F0503), and 50 ng/ml nerve growth factor (Invitrogen, 13257-019). The mixture of aphidicolin and 5-fluoro-2-deoxyuridine inhibits proliferation of Schwann cells by inhibition of DNA polymerase (Spadari et al., 1985; Wallace and Johnson, 1989), thus constituting a highly pure sensory neuron culture (Heermann et al., 2012). DRGs were cultured for 7–11 d at 37°C in 5% CO₂. Axotomy of DRG explants was made using a micropipette tip to separate all the axons from the somas. Chemical injury was achieved by adding 1 μ M vinblastine (Sigma-Aldrich, V1377) to the culture medium.

Western blot. Axons obtained by axotomy from rat DRG explants ($n = 3$) were homogenized in radioimmunoprecipitation assay buffer with 1% protease inhibitor mixture (Sigma-Aldrich, P8340) plus phenylmethylsulfonyl fluoride and phosphatase inhibitors NaVO_3 and NaF. Proteins were quantified by bicinchoninic acid assay (Thermo Fisher Scientific, 23225). Loading buffer and 40 μg of lysate were combined, subjected to 10% SDS-PAGE electrophoresis, and transferred to polyvinylidene fluoride membranes (Thermo Fisher Scientific, 88518). The following primary antibodies were used at 1:1000 dilution: anti-RIPK3 (Abcam, catalog #ab16090, RRID:AB_302252), anti p-MLKL (Abcam, catalog #ab196436, RRID:AB_2687465), and anti-MLKL (Abcam, catalog #ab172868, RRID:AB_2737025). Secondary antibodies used were goat anti-mouse horseradish peroxidase (HRP; Bio-Rad, catalog #170-6516, RRID:AB_11125547) and goat anti-rabbit HRP (Bio-Rad, catalog #170-6515, RRID:AB_11125142) at 1:3000. Western blots were revealed by enhanced chemiluminescence in a Chemi-doc Gel Imaging System (Bio-Rad, XRS+). Band analysis was performed using Image Lab (RRID:SCR_014210) and ImageJ (National Institutes of Health, RRID:SCR_003070) software.

Quantitative real-time PCR. The mRNA levels of *RIPK3* and *MLKL* were analyzed in DRG neurons infected with lentivirus containing short hairpin *RIPK3* (shRIPK3) or shMLKL before and after axotomy. Neurons were homogenized in Trizol reagent (Invitrogen) for extraction of total RNA based on the manufacturer's protocol. Total RNA was quantitated using a Multiskan Go ultraviolet/visible spectrophotometer (Thermo Fisher Scientific, 1510). Complementary DNA from 1.0 μg of total RNA was synthesized using iScriptTM reverse transcription Supermix for RT-qPCR (Bio-Rad). mRNA levels in each sample were determined by real-time PCR using EvaGreen qPCR dye. The reverse-transcription product was included in the 5 \times HOT FIREPOL EvaGreen qPCR Mix Plus (ROX; Solis BioDyne), along with rat *RIPK3* and *MLKL* primers, and the mixture was placed in the StepOnePlus Real-Time PCR System (Thermo Fisher Scientific, Lightcycler System). Thermal cycling parameters were 30 s at 95°C, followed by 40 cycles of 5 s at 95°C, 5 s at 60°C, and 30 s at 72°C. At the end of the program, melting curve analysis was performed at 72°C for 30 s, followed by a cooling step at 37°C for 30 s. Each sample was also run with primers for a housekeeping gene. Primer sequences were obtained from the literature (Liu et al., 2016): rat *RIPK3* forward primer 5'-CTGTGCGCTGCTAGAGGAAG-3', and reverse primer 5'-TCTGCTAAGTTGGCGTGGAG-3'; rat *MLKL* forward primer 5'-CCCGAGTTGTCAGGAGAT-3', and reverse primer 5'-TCTCCAAGATTCC ATCCGAG-3'; and *Gadph* forward primer 5'-TCCCTCAAGATTGTCAGCAA-3', and reverse primer 5'-AGATCCA CAACGGATACATT-3'. For relative comparison of each gene, we analyzed the Ct value with the $\Delta\Delta\text{Ct}$ normalizing method.

Immunofluorescence. For immunofluorescence analysis, the nerve explants were fixed by immersion in 4% paraformaldehyde in 1 \times PBS for 1 h, followed by three 10 min washes in 1 \times PBS, sucrose gradient (5, 10, and 20% in 1 \times PBS), and then embedded in Optimal Cutting Temperature compound (Sakura Tissue Tek, 4583). Cryostat sections from the middle of the explants were cut transversely at 10 μm thickness and mounted on Superfrost Plus slides (Thermo Fisher Scientific, J1800AMNZ). Sections were washed in 1 \times PBS for 10 min and then blocked/permeabilized in 0.1% Triton X-100, 5% fish skin gelatin (Sigma-Aldrich, G7765) in 1 \times PBS for 1 h at RT. Sections were incubated in primary antibodies in 0.1% Triton X-100, 1% fish skin gelatin solution overnight at 4°C, washed in 1 \times PBS 3 \times 10 min, and incubated in secondary antibodies for 2 h at RT. Sections were washed 3 \times 10 min in 1 \times PBS and mounted in Fluoromount G (Electron Microscopy Sciences, 17984-25). For immunofluorescence analysis, the embryonic DRG explants were fixed in 4% paraformaldehyde in 1 \times PBS for 20 min, followed by three 10 min washes in 1 \times PBS. The following antibodies were used for immunofluorescence analysis: rabbit anti-neurofilament heavy chain (NFH; Sigma-Aldrich, N4142) at 1:5000; chicken anti-neurofilament medium chain (Millipore, catalog #AB5735, RRID:AB_240806) at 1:2000; anti-acetylated tubulin (Sigma-Aldrich, catalog #T6793, RRID:AB_477585) at 1:1000, and COX IV (Abcam, catalog #ab16056, RRID:AB_443304) at 1:1000. We used secondary antibodies

conjugated to AlexaFluor dyes at 1:1000 (Jackson ImmunoResearch Laboratories).

Electron microscopy. Nerves were fixed overnight by immersion in 2.5% glutaraldehyde, 0.01% picric acid, and 0.1 M cacodylate buffer, pH 7.4. Nerves were rinsed in the same buffer, immersed in 1% OsO_4 for 1 h, followed by in block incubation with 2% uranyl acetate for 2 h. Nerves were dehydrated with a graded series of ethanol and propylene oxide, and infiltrated with Epon (Ted Pella). Ultrathin sections from the middle of the explants were contrasted with 1% uranyl acetate and lead citrate. Grids were examined with a Philips Tecnai 12 electron microscope operated at 80 kV. Negative films were developed and scanned.

Axonal degeneration quantification. The number of axons per area of nerve tissue was assessed in confocal images of neurofilament-immunostained explant sections (matched for laser power, photomultiplier tube gain/offset, and postprocessing) using the particle analysis macro of ImageJ. Relative neurite integrity was based on the ratio of the area of fragmented axons to total axonal area (Villegas et al., 2014). Degenerated axon fragments were detected using the particle analyzer algorithm of ImageJ and the total fragmented axon area versus total axonal area was used to estimate a degeneration index.

Electrophysiology. The viability of sciatic nerve explants was tested by the recording of the compound action potential (CAP). Sciatic nerve explants from mice were desheathed from the perineurium and transferred to a recording chamber filled with HBSS, supplemented with 5 mM HEPES buffer, pH 7.35 (Alcayaga et al., 1998). The proximal end of each explant was pinned onto a stimulating bipolar tungsten electrode at the bottom of the recording chamber, while the distal end was pinned onto the recording electrode at the top of the chamber and immersed in mineral oil. Stimulating pulses of 50 μs each were delivered at 1 s intervals. The maximum fast monophasic A-wave was obtained by increasing the intensity of the pulse. The maximal peak height and area of the CAP was analyzed.

Virion production and lentiviral transduction. Human embryonic kidney 293T cells (ATCC) were transfected into 15 cm^2 dishes using the calcium chloride (1.25 M) method. Media was replaced 4 h after the initial transfection, and 48 h later lentivirus-enriched media were collected and cleared by slow centrifugation at 1500 rpm for 5 min. Subsequently, the conditioned media was filtered through a 0.45- μm -pore conic tube filter. The lentiviruses were concentrated using the Lenti-X Concentrator reagent (Clontech), and then stored at -80°C . Four plasmids for lentiviral transduction of *RIPK3* shRNA were designed by Biosettia using the *Rattus norvegicus* *RIPK3* mRNA sequence [National Center for Biotechnology Information (NCBI) reference sequence NM_139342.1] and cloned in a pLVRNAi vector that coexpresses GFP as reporter gene. The *ripk3* shRNA oligo vector used the following sequences: GAAAGGCTTCTA AAGCAA; GAAGCATCATTGGGCAT; GAAGAAACAGCAATCCTT; CCTACAGTCTATTGTCTT. One plasmid for lentiviral transduction of *MLKL* shRNA was designed and packaged by Vigene Bioscience using the *Rattus norvegicus* *MLKL* mRNA sequence (NCBI reference sequence XM_008772571.2) and cloned in a pLV-RNAi vector that coexpresses GFP as a reporter gene. The *mlkl* shRNA oligo vector used the following sequences: GACCAAAGTGAAGACAAATAA; TCCCAACATCCTGCG TATATT. One plasmid for lentiviral transduction of scramble shRNA was designed by Vigene Bioscience and cloned in a pLV-RNAi vector that coexpresses GFP as reporter gene. Transduction of DRG explants was performed in 300 μl of free-serum media with 1×10^6 TU (transduction units) per explant in 24-well plates at 2 d *in vitro* (DIV) and maintained in culture until axotomy or vinblastine treatment at 7 DIV.

Statistical analysis. Statistical significance was established at $p < 0.05$ by one-way ANOVA with Tukey's post-test or two-way ANOVA for multiple comparisons. The analysis was performed using GraphPad Prism Software v6.0 and performed with data obtained from ≥ 3 independent experiments.

Results

RIPK1 inhibition delays axonal degeneration in both the PNS and CNS

To determine whether necroptosis is associated with axonal degeneration, we first used an *ex vivo* sciatic nerve model of injury-

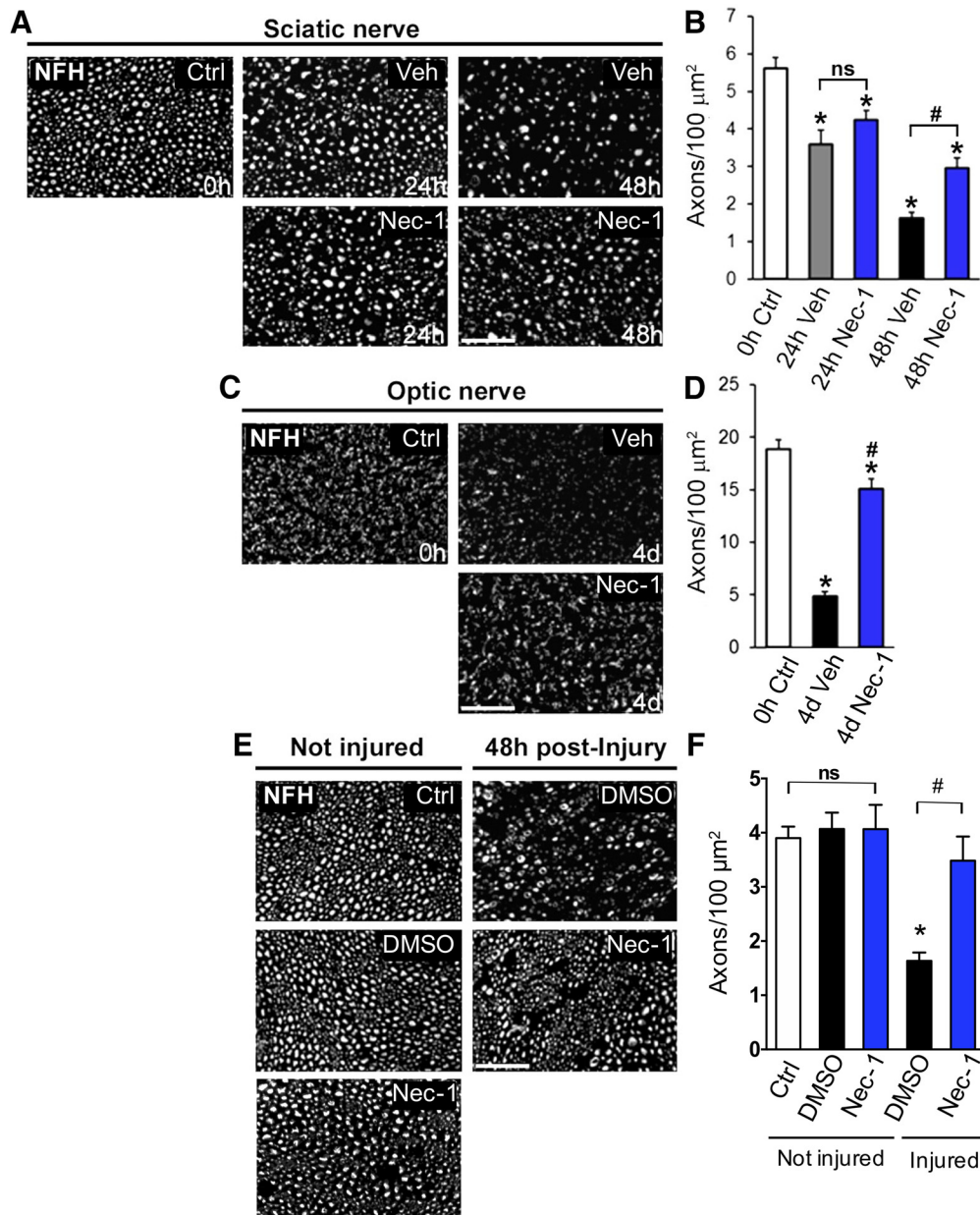


Figure 1. Pharmacological inhibition of RIPK1 delays degeneration of both peripheral and central axons. The degree of axonal degeneration was quantitatively assessed by immunofluorescence using NFH immunostaining in transverse nerve sections. **A–D**, Sciatic and optic nerve explants were treated with Nec-1 (100 μM) or vehicle (DMSO) for 48 h and 4 d, respectively. **E**, Axonal degeneration was evaluated *in vivo* in crushed sciatic nerves injected with 200 μl of 0.64 mM Nec-1 or vehicle (3% DMSO in PBS) 48 h after injury. Nec-1 treatments strongly delay axonal degeneration triggered by nerve injury. Scale bar, 20 μm . **B, D, F**, Quantification of NFH-positive axons in nerve cross sections was expressed as axonal density for both *ex vivo* explants (**B, D**) and *in vivo* crushed nerve (**F**). In injured nerves, statistically significant protection from axonal degeneration was seen after Nec-1 treatment compared with vehicle-treated nerves ($^{\#}p < 0.0001$) or noninjured controls ($^*p < 0.0001$). Mean values are shown, error bars indicate SEM, $N = 3$ per group; analysis was performed by one-way ANOVA test.

induced axonal degeneration. This model allowed us to study axonal degeneration in the absence of any effects from the neuronal cell body or the immune system (Barrientos et al., 2011). We tested the effect of Nec-1, a drug that inhibits the kinase activity of RIPK1 and prevents RIPK1/RIPK3 complex formation, thus strongly inhibiting cell death by necroptosis (Degterev et al., 2008). In vehicle-treated explants, >50% of axons degenerated by 48 h [0 h control (Ctrl): 5.612 ± 0.29 axons/100 μm^2 ; 48 h vehicle (veh): 1.625 ± 0.26 axons/100 μm^2 , $F_{(10,22)} = 94.65$, $p < 0.0001$, ANOVA], as observed by NFH immunostaining and quantitative analysis of axonal density (Fig. 1A,B). In explants treated with Nec-1 (100 μM), significant protection from axonal degeneration was observed at 48 h (48 h veh: 1.625 ± 0.26 axons/

100 μm^2 ; 48 h Nec-1: 2.948 ± 0.28 axons/100 μm^2 , $F_{(10,22)} = 94.65$, $p = 0.0094$, ANOVA; Fig. 1A, B). To determine the protective effect of Nec-1 in damaged CNS axons, we analyzed optic-nerve explants using the same approach described above. Vehicle-treated optic-nerve explants showed a dramatic reduction in axonal density after 4 d in culture (0 h Ctrl: 18.84 ± 0.80 axons/100 μm^2 ; 4 d veh: 4.86 ± 0.41 axons/100 μm^2 , $F_{(2,18)} = 96.97$, $p < 0.0001$, ANOVA), which was strongly inhibited by Nec-1 (100 μM) treatment (4 d veh: 4.86 ± 0.41 axons/100 μm^2 ; 4 d Nec-1: 15.07 ± 0.89 axons/100 μm^2 , $F_{(2,18)} = 96.97$, $p < 0.0001$, ANOVA; Fig. 1C,D). To test whether RIPK1 inhibition protects degenerating axons *in vivo*, we injected 2 μl of 0.64 mM of Nec-1 or its vehicle (3% DMSO in PBS) locally in mouse sciatic

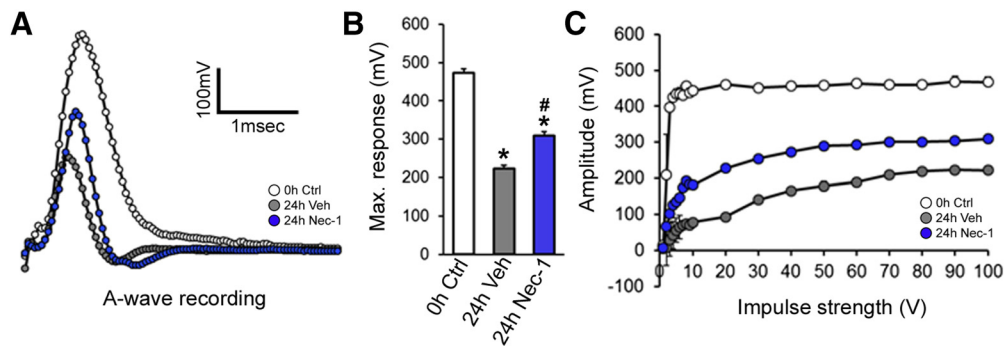


Figure 2. Nerve function is protected by RIPK1 inhibition. The size of the CAP was measured in sciatic nerve explants cultured for 24 h in the presence or absence of Nec-1 (100 μM). **A**, CAP A-wave length and amplitude were registered in vehicle-treated nerves. **B**, The maximum peak of the CAP was analyzed. **C**, The recruitment of excitable fibers within the nerve was measured at fixed stimulus intensity and the maximum fast monophasic A-wave was obtained by increasing the intensity of the pulse. Nerves cultured for 24 h with Nec-1 showed a significantly higher response than nerves cultured under vehicle conditions. One-way ANOVA with *post hoc* Tukey; # indicates statistically significant compared with Veh; * indicates statistically significant compared with Ctrl.

nerves. Injected nerves were then crushed proximal to the injection site, and axon density in the distal stump was evaluated 48 h after injury by NFH immunostaining. We detected a significant decrease in axonal density in injured nerves treated with vehicle compared with undamaged nerves (Ctrl: 4.04 ± 0.11 axons/100 μm^2 ; injured: 1.63 ± 0.15 axons/100 μm^2 , $F_{(3,20)} = 16.12$, $p < 0.0001$, ANOVA). Meanwhile, Nec-1 effectively protected axons from degeneration after injury (injured: 1.63 ± 0.15 axons/100 μm^2 ; injured+Nec-1: 3.48 ± 0.44 axons/100 μm^2 , $F_{(3,20)} = 16.12$, $p < 0.0001$, ANOVA; Fig. 1E,F). No significant effect was observed for Nec-1 in noninjured axons compared with control conditions. These results suggest that RIPK1 is required for the progression of axonal degeneration in *ex vivo* and *in vivo* models of Wallerian degeneration of both the CNS and PNS.

RIPK1 inhibition preserves nerve function during Wallerian degeneration

Isolated axons progressively lost their capacity to conduct action potentials (Donaldson et al., 2002). To evaluate whether necroptosis inhibition preserves axonal function during Wallerian degeneration, we analyzed the CAP of sciatic nerve explants cultured for 24 h in the presence or absence of Nec-1 (100 μM). At 24 h after injury, the CAP A-wave of vehicle-treated nerves was severely affected, shortening both length and amplitude to half the A-wave of intact nerves (0 h Ctrl: 473.6 ± 0.67 mV; 24 h veh: 223.3 ± 4.74 mV, $F_{(2,6)} = 557.4$, $p < 0.0001$, ANOVA; Fig. 2A). In Nec-1-treated nerves, a significant inhibition of the injury-induced A-wave amplitude decay was observed compared with damaged and vehicle-treated nerves (24 h veh: 223.3 ± 4.74 mV; 24 h Nec-1: 309.6 ± 5.69 mV, $F_{(2,6)} = 557.4$, $p < 0.0001$, ANOVA; Fig. 2A,B). We also evaluated the recruitment of excitable fibers within the nerve at a fixed stimulus intensity. Myelinated A-fibers of freshly dissected sciatic nerves reached their maximum response at low stimulus intensities. In contrast, nerve explants cultured for 24 h showed a smaller maximum response, which was only detected under high stimulus intensities, reflecting a lower number of functional axons due to the degenerative process. On the other hand, nerves cultured for 24 h with Nec-1 showed a significantly higher response than nerves cultured under vehicle conditions. Results were similar at lower stimulus intensities (Fig. 2C). Altogether, these results demonstrate that RIPK1 inhibition, besides protecting axons from morphological degeneration after transection, also preserves some of their ability to function after nerve damage.

Mitochondrial fragmentation and Drp1 activity are crucial for late phases of axonal degeneration

Axonal degeneration after mechanical damage is associated with mitochondrial dysfunction, including a decline in mitochondrial membrane potential and opening of the mPTP (Barrientos et al., 2011). Mitochondrial fragmentation has also been documented as a consequence of necroptosis activation (Young et al., 2010; Z. Wang et al., 2012; Yamada et al., 2016). Therefore, we evaluated changes in mitochondrial morphology during axonal degeneration and the effect of pharmacological RIPK1 inhibition. In uninjured nerves, axonal mitochondria exhibited long, tubular shapes, which become rounded by 48 h after injury as axons progressively degenerate (Fig. 3A,B). Using confocal microscopy, we measured mitochondrial length in axons from teased fibers of sciatic nerves at 0, 24, and 48 h after injury. In intact axons, mitochondria displayed a wide range of lengths from <2 μm up to >8 μm . At 24 and 48 h after injury, the distribution of mitochondrial lengths became much narrower and shifted to smaller values (Fig. 3C), with significant decreases in the mean mitochondrial length (0 h Ctrl: 4.116 ± 0.178 μm ; 24 h veh: 1.737 ± 0.123 μm ; 48 h veh: 0.8024 ± 0.064 μm , $F_{(2,6)} = 171.9$, $p < 0.0001$, ANOVA; Fig. 3D). Furthermore, a significant increase in mitochondrial density at 24 and 48 h after injury (0 h Ctrl: 22.05 ± 1.392 mitochondria/100 μm^2 ; 24 h veh: 34.64 ± 1.734 mitochondria/100 μm^2 ; 48 h veh: 36.21 ± 1.007 mitochondria/100 μm^2 , $F_{(2,51)} = 30.33$, $p < 0.0001$, ANOVA; Fig. 3E) suggests that the changes in mitochondrial length after axonal damage are associated with a fragmentation process. By electron microscopy of longitudinal sections of sciatic nerves, evidence for mitochondrial fragmentation and swelling was commonly observed at 24 and 48 h postinjury, respectively (Fig. 3F).

We next tested whether axonal mitochondrial fragmentation after nerve injury was associated with RIPK1 activation. To this end, the effect of Nec-1 on injury-induced mitochondrial fragmentation was evaluated. At 24 h after injury, mitochondrion shortening was inhibited in nerves treated with Nec-1 compared with nerve explants treated with vehicle (24 h veh: 1.351 ± 0.046 μm ; 24 h Nec-1: 2.698 ± 0.072 μm , $F_{(2,1269)} = 115.0$, $p < 0.0001$, ANOVA; Fig. 4A,B). In addition, the increase in mitochondrial density 24 h after damage was inhibited by Nec-1 when compared with vehicle-treated nerves (24 h veh: 33.35 ± 2.68 μm ; 24 h Nec-1: 27.87 ± 1.62 μm , $F_{(4,40)} = 4.354$, $p < 0.0001$, ANOVA; Fig. 4C).

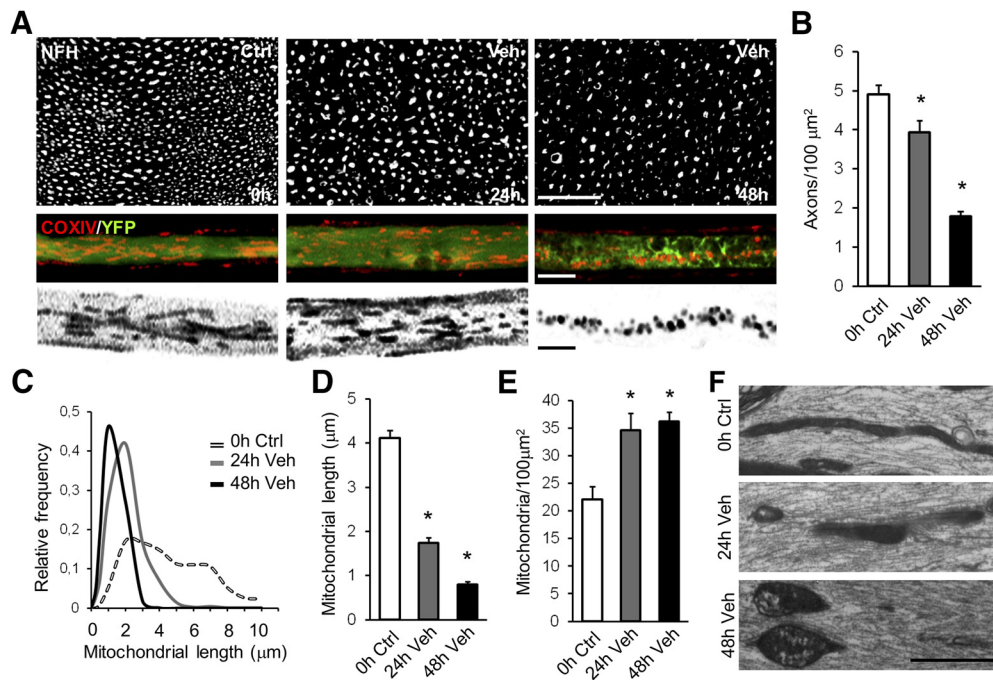


Figure 3. Mitochondrial fragmentation is associated with axonal degeneration after nerve injury. Mitochondrial morphology was evaluated along with axonal degeneration in teased fibers of sciatic nerve explants at 0, 24, and 48 h after injury. **A, B**, Axonal degeneration progression is shown in nerve transverse sections immunostained for neurofilament (**A**) and quantified as axonal density from NFH-positive axons per area (**B**). Scale bar, 20 μm . Mitochondria were immunodetected with a COX IV antibody (mitochondria, red) in teased sciatic nerves dissected from Thy1-YFP mice (axons, green). Morphological analysis of mitochondria was performed from COX IV+ labeling of binarized images (**A**, bottom). Scale bar, 5 μm . **C, D**, Mitochondrial length was plotted as relative frequency (**C**) and as mean values (**D**). **E**, Decreased mitochondrial length along axonal degeneration was associated with increased mitochondrial density, measured as number of mitochondria per area. **F**, Longitudinal sections of sciatic nerves were analyzed by electron microscopy, showing mitochondrial fragmentation and swelling at 24 and 48 h after injury, respectively. Scale bar, 1 μm . One-way ANOVA with *post hoc* Tukey, * indicates statistically significant compared with Ctrl.

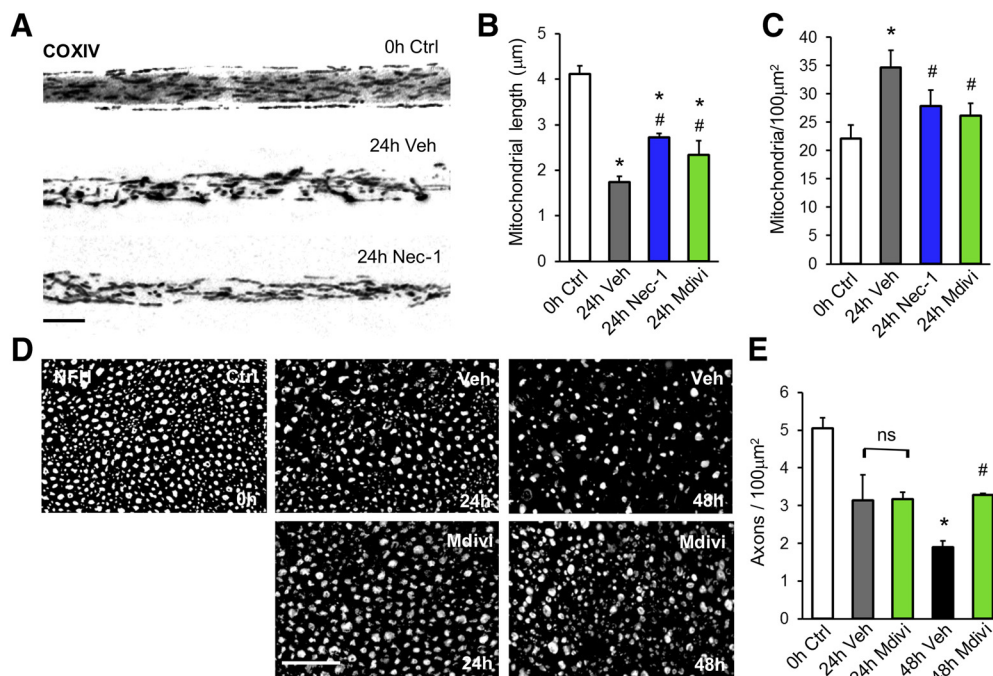


Figure 4. RIPK1 inhibition prevents axonal degeneration-associated mitochondrial fragmentation. **A**, Mitochondrial fragmentation was evaluated at 24 h after injury from binarized images of COX IV-immunostained teased nerves. **B**, Mitochondrion shortening was inhibited in nerves treated with Nec-1 (100 μM) or Mdivi (200 μM) compared with nerve explants treated with vehicle. **C**, Increased mitochondrial density was also inhibited by both treatments. **D, E**, Axonal integrity after Mdivi treatment was also evaluated by NFH staining of transversal nerve sections at 24 and 48 h after injury. One-way ANOVA with *post hoc* Tukey, * indicates statistically significant compared with Ctrl; # indicates statistically significant compared with Veh.

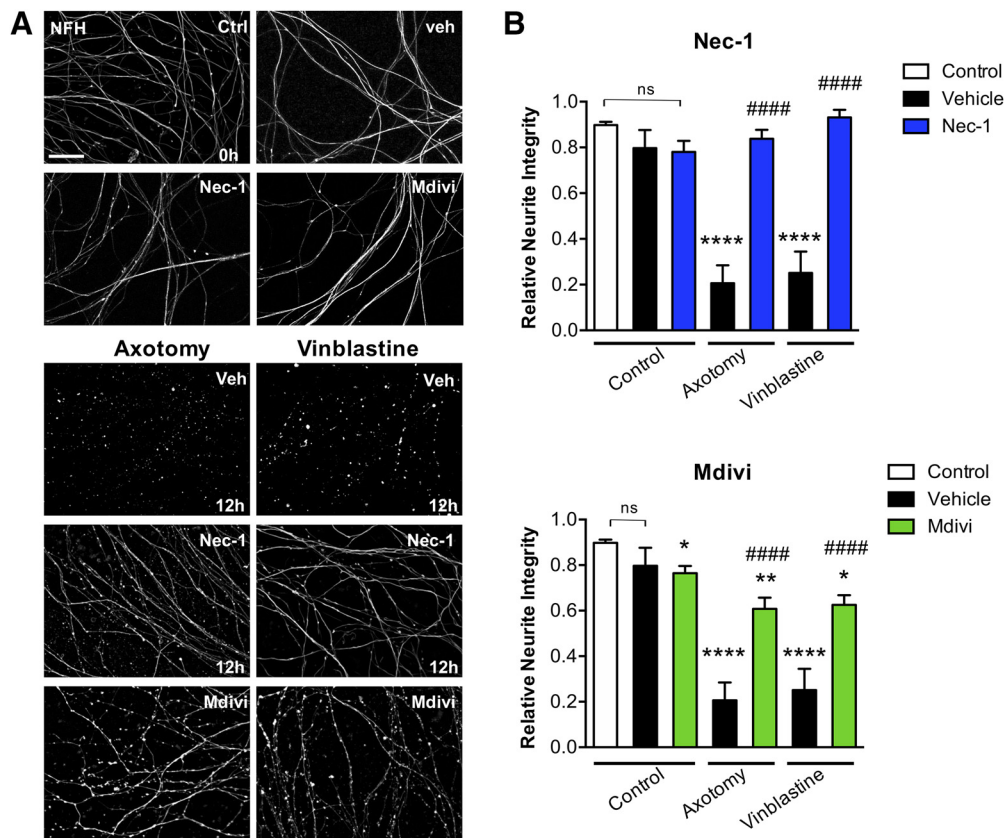


Figure 5. RIPK1 activation and mitochondrial fragmentation contributes to axonal degeneration by a cell-autonomous mechanism. DRG explants were damaged by axotomy or vinblastine ($1 \mu\text{M}$) treatment in the presence of Nec-1, Mdivi, or vehicle as a control, and analyzed after 12 h. *A, B*, Degeneration was evaluated in distal axons immunostained for NFH (*A*), quantitatively assessed, and represented as relative neurite integrity (*B*). Scale bar, $30 \mu\text{m}$. Two-way ANOVA with *post hoc* Tukey, * indicates statistically significant compared with Ctrl; # indicates statistically significant compared with Veh.

Drp1-dependent mitochondrial fragmentation has been associated with the execution stages of necroptosis in cultured cell lines after treatment with $\text{TNF-}\alpha$ (Wang Z et al., 2012). To test the contribution of mitochondrial fragmentation to axonal degeneration, sciatic nerve explants were cultured in the presence or absence of Mdivi, a specific pharmacological inhibitor of Drp1 (Cassidy-Stone et al., 2008). Treatment with Mdivi ($200 \mu\text{M}$) partially inhibited mitochondrial fragmentation and the increase in mitochondrial density at 24 h after injury compared with vehicle-treated nerve explants (Fig. 4*B, C*). The inhibitory effects of Mdivi in these mitochondrial parameters were comparable to those of nerve explants treated with Nec-1. When axonal integrity after Mdivi treatment was evaluated by NFH immunostaining, a significant inhibition of axonal degeneration was observed 48 h after injury compared with control nerves (48 h veh: 19.98 ± 1.31 axons/ $100 \mu\text{m}^2$; 48 h Mdivi: 34.24 ± 0.86 axons/ $100 \mu\text{m}^2$, $F_{(2,9)} = 50.4$, $p = 0.0029$, ANOVA; Fig. 4*D, E*). Cotreatment with Mdivi plus Nec-1 had no additive protective effects, suggesting that Drp1-dependent mitochondrial fragmentation and RIPK1 activation uses the same signaling pathway to regulate axonal degeneration.

RIPK1 activation and mitochondrial fragmentation contributes to axonal degeneration in a cell-autonomous manner

During Wallerian degeneration, Schwann and immune cells contribute to axonal degeneration (Catenaccio et al., 2017). To analyze whether necroptosis and mitochondrial fragmentation are

involved in axonal degeneration in a cell-autonomous manner, pharmacological treatments with Nec-1 and Mdivi were performed in cultures of sensory neurons from DRGs devoid of glial cells (Villegas et al., 2014). Even though embryonic DRG neurons *in vitro* show very different timing of degeneration compared with sciatic nerve explants (Barrientos et al., 2011), they have been widely used as a model for Wallerian degeneration, since both embryonic DRG neurons and adult sciatic nerves degenerate by a mechanism that depends on Wld^s and Sarm1 (M.S. Wang et al., 2001; Osterloh et al., 2012). Axonal degeneration was induced by either DRG axotomy or treatment with vinblastine ($1 \mu\text{M}$), a chemotherapeutic agent that induces Wallerian-like degeneration in peripheral nerves (Bradley et al., 1970; Taillibert et al., 2016) in the presence of Nec-1, Mdivi, or vehicle as a control, and analyzed after 12 h. Axons were immunostained for NFH and axonal integrity was quantitatively assessed. After either axotomy or vinblastine treatment, axons completely degenerate by 12 h (0 h Ctrl: 0.898 ± 0.014 integrity index; 12 h veh+axotomy: 0.206 ± 0.079 integrity index; 12 h veh+vinblastine: 0.2515 ± 0.093 integrity index, $F_{(6,44)} = 33.74$, $p < 0.0001$, two-way ANOVA; Fig. 5*A, B*). By contrast, Nec-1 treatment completely inhibits axonal degeneration 12 h after axotomy or vinblastine treatment compared with vehicle-treated axons (axotomy: 12 h veh: 0.206 ± 0.079 integrity index; 12 h Nec-1: 0.838 ± 0.039 integrity index; vinblastine: 12 h veh: 0.2515 ± 0.093 integrity index; 12 h Nec-1: 0.931 ± 0.034 integrity index; $F_{(6,44)} = 33.74$, $p < 0.0001$, two-way ANOVA; Fig. 5*A, B*). No significant effect on axonal degeneration was observed in control axons treated

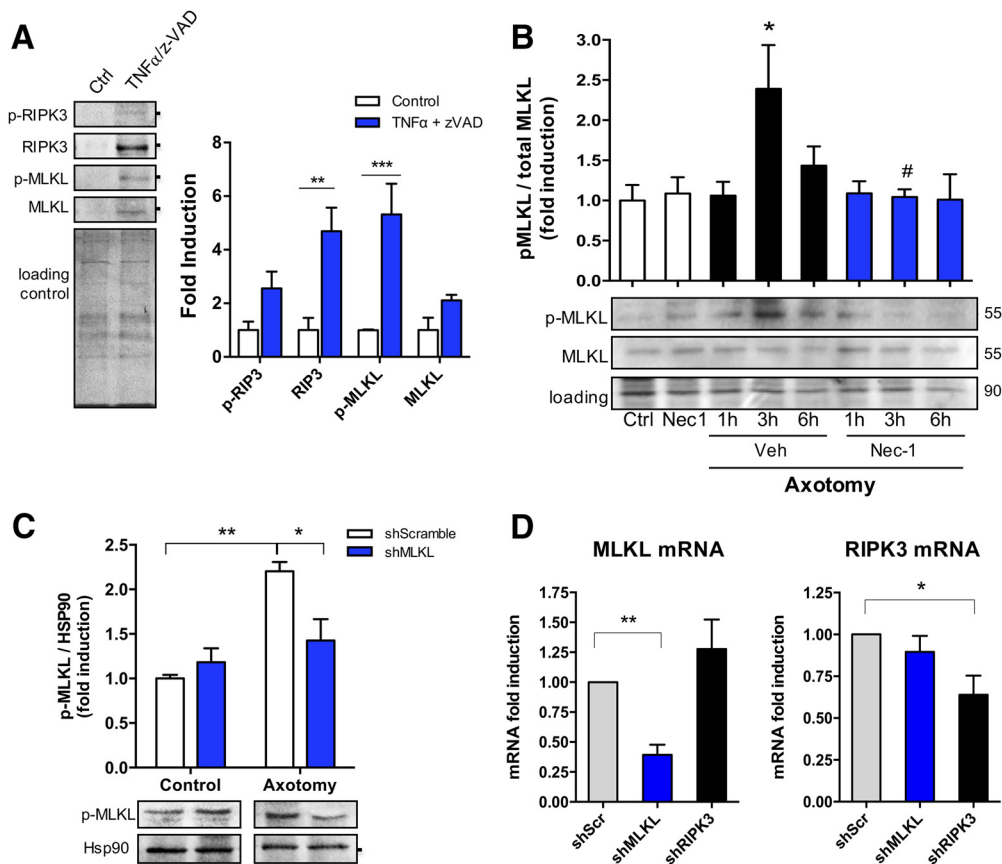


Figure 6. Necroptosis is activated after axonal injury. Necroptotic pathway activation was evaluated after TNF α /z-VAD treatment through the detection of RIPK3, MLKL, and their phosphorylated forms by Western blot. **A**, Total protein (Coomassie staining) was used as a loading control to normalize the expression of the different proteins, which were expressed as fold induction compared with control ($n = 3$). Downstream activation of the pathway was determined in mechanically injured axons in a time course (≤ 6 h) through the detection of MLKL and p-MLKL. Results are represented as pMLKL/MLKL ratio from $n = 5$ independent experiments. Total protein was used as a loading control. Two-way ANOVA with *post hoc* Tukey, * indicates statistically significant compared with Ctrl; # indicates statistically significant compared with 3 h + Veh. **B**, **C**, Specificity of p-MLKL antibody was tested in injured axons devoid of MLKL by using DRGs transduced with lentiviral vectors containing MLKL shRNA (**B**) and compared with a control nontargeting shRNA (scramble; $n = 3$; **C**). **D**, The magnitude of MLKL or RIPK3 knock-down was evaluated by real-time qPCR in dissociated DRGs transduced with lentivirus containing shMLKL or shRIPK3, respectively ($n = 3$). One-way ANOVA with *post hoc* Tukey, * indicates significant compared with Scramble shRNA.

with Nec-1. Treatment of mechanically or chemically injured axons with Mdivi partially inhibits their degeneration and confers significant protection when compared with vehicle-treated axons (axotomy: 12 h veh: 0.206 ± 0.079 integrity index; 12 h Mdivi: 0.608 ± 0.049 integrity index; vinblastine: 12 h veh: 0.2515 ± 0.093 integrity index; 12 h Mdivi: 0.625 ± 0.043 integrity index; $F_{(6,44)} = 33.74, p < 0.0001$, two-way ANOVA; Fig. 5A,B). Indeed, treatment of uninjured axons with Mdivi for ≤ 12 h leads to progressive axonal degeneration *in vitro* (Ctrl: 0.898 ± 0.014 integrity index; Mdivi: 0.764 ± 0.031 integrity index, $p = 0.0491$; Fig. 5B). This is consistent with the physiological role of mitochondrial fission in cellular homeostasis, which might explain the partial protection conferred by Mdivi treatment in injured axons. These results strongly suggest that both necroptosis and mitochondria fragmentation are involved in axonal degeneration by a cell-autonomous mechanism and triggered by diverse prodegenerative insults.

The downstream effector of necroptosis MLKL is transiently activated after axonal damage

To evaluate the capability of embryonic DRG cultures to activate necroptosis, DRG explants were exposed to TNF α (100 ng/ml) and the pan-caspase inhibitor z-Val-Ala-Asp(Ome)-fluoromethylketone (z-VAD; 20 μ M), which in combination are known to

induce necroptosis in diverse cell types (Holler et al., 2000; Degterev et al., 2005). After 72 h of incubation, only axons were removed and immediately lysed to evaluate the axonal protein levels of key participants of necroptosis by Western blot. All markers, including RIPK3, MLKL, and their phosphorylated forms, increased after treatment with TNF α /z-VAD. Interestingly, the phosphorylated form of the downstream effector MLKL was significantly increased after TNF α /z-VAD treatment compared with control axons (Ctrl: 1.0003 ± 0.022 p-MLKL fold induction; TNF α /z-VAD: 5.3180 ± 1.148 p-MLKL fold induction, $F_{(3,16)} = 3.242, p = 0.0006$, ANOVA; Fig. 6A). To determine whether necroptosis is activated during axonal degeneration, DRG explants were damaged by axotomy and the axonal levels of p-MLKL were measured. A significant and transient increase of p-MLKL was observed 3 h after axotomy (Ctrl: 1.00 ± 0.194 p-MLKL fold induction; 3 h veh: 2.390 ± 0.546 p-MLKL fold induction, $p = 0.0208$), an effect that was significantly inhibited by Nec-1 treatment (3 h veh: 2.390 ± 0.546 p-MLKL fold induction; 3 h Nec-1: 1.044 ± 0.094 p-MLKL fold induction, $F_{(7,28)} = 3.307, p = 0.0276$, ANOVA; Fig. 6B). Since necroptosis regulators are poorly expressed in the nervous system, we evaluated the specificity and sensitivity of the pMLKL antibody by using lentivirus expressing shRNA sequences designed against MLKL in axons 3 h after axotomy, a condition that we already showed to

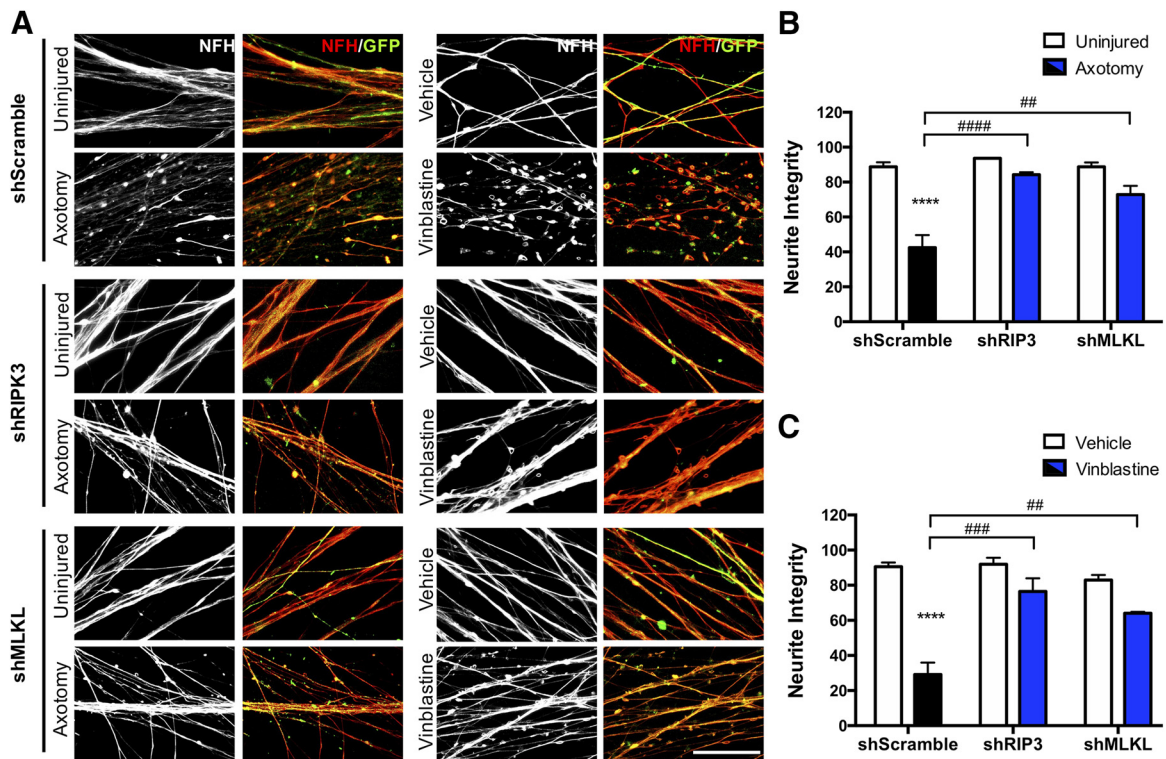


Figure 7. Knock down of *RIPK3* and *MLKL* in sensory neurons protects them from axonal degeneration *in vitro*. DRGs were transduced with lentiviral vectors containing *RIPK3* or *MLKL* shRNA versus a control nontargeting shRNA (scramble). All plasmids express GFP for identification of transduced neurons. **A**, Representative images before and after injury (axotomy or vinblastine) are shown (NFH in red, GFP signal in green). Scale bar, 5 μ m. **B**, **C**, Quantifications of axonal degeneration 8 h after axotomy (**B**) or 16 h after vinblastine (1 μ m) treatment (**C**). Protection with shRIPK3 or shMLKL is represented with blue bars under both conditions. Two-way ANOVA with *post hoc* Tukey, * indicates statistically significant compared with Veh; # indicates statistically significant protection compared with axotomized or vinblastine-treated axons.

increase p-MLKL levels (Fig. 6B). A significant decrease in pMLKL was found in shMLKL transduced and injured axons [shScramble (shScr) axotomy: 2.203 ± 0.105 p-MLKL fold induction; shMLKL axotomy: 1.425 ± 0.240 p-MLKL fold induction, $F_{(1,8)} = 9.743$, $p = 0.0297$, ANOVA; Fig. 6C]. These results indicate that necroptosis is induced in embryonic DRG axons after mechanical injury and that necroptotic regulator levels can be specifically detected by Western blot. Furthermore, to evaluate the magnitude of the knock down, *MLKL* and *RIPK3* mRNA levels were studied by real-time PCR in dissociated DRGs transduced with lentiviral vectors containing shMLKL or shRIPK3 sequences. shMLKL produced a 60% decrease in *MLKL* mRNA levels compared with shScramble (shScr: 1.000 ± 0.00 p-MLKL fold induction; shMLKL: 0.3935 ± 0.084 p-MLKL fold induction, $F_{(2,9)} = 9.056$, $p = 0.0070$, ANOVA), while *RIPK3* levels showed nonsignificant changes after *MLKL* shRNA. shRNA against *RIPK3* showed a 36% fall in *RIPK3* mRNA levels (shScr: 1.000 ± 0.00 p-MLKL fold induction; shRIPK3: 0.6393 ± 0.114 p-MLKL fold induction, $F_{(2,9)} = 4.647$, $p = 0.0411$, ANOVA), while *MLKL* levels were not significantly affected (Fig. 6D).

RIPK3 and MLKL knock down delays axonal degeneration

To evaluate the participation of downstream effectors of necroptosis in injury-induced axonal degeneration at a genetic level, DRG neurons were transduced with lentivirus expressing shRNA sequences designed against *RIPK3* and *MLKL*, or with a scramble sequence as a control. The lentiviral vector coexpresses *GFP* as a reporter gene to identify transduced neurons in the culture for the subsequent morphological analysis. After transduction, axons were injured by axotomy or chemically damaged with

vinblastine. Integrity was evaluated 8 or 16 h after injury, respectively, by NFH immunostaining in GFP-expressing axons. In neurons transduced with the control shRNA, both injury paradigms induce a massive degeneration of axons (shScr uninjured: 88.787 \pm 2.610% of integrity; shScr axotomy: 42.491 \pm 7.191% of integrity, $p < 0.0001$, $F_{(1,12)} = 55.47$, two-way ANOVA; shScr veh: 12 h veh: 90.568 \pm 2.342% of integrity; shScr vinblastine: 29.147 \pm 6.838% of integrity, $p < 0.0001$, $F_{(1,12)} = 69.67$, two-way ANOVA; Fig. 7). In contrast, shRNA for *RIPK3* and *MLKL* effectively protects from axonal degeneration triggered by axotomy or vinblastine treatment (axotomy: shScr: 42.491 \pm 7.191% of integrity; shRIPK3: 84.251 \pm 1.386% of integrity; or shMLKL: 72.808 \pm 5.057% of integrity, $p < 0.0001$ and $p = 0.0016$, respectively, $F_{(2,12)} = 18.15$, two-way ANOVA; vinblastine: shScr: 29.147 \pm 6.838% of integrity; shRIPK3: 76.436 \pm 7.520% of integrity; or shMLKL: 64.097 \pm 0.793% of integrity, $p = 0.0001$ and $p = 0.0021$, respectively, $F_{(2,12)} = 13.56$, two-way ANOVA; Fig. 7A, B). Together, these results demonstrate that key players of the necroptotic pathway, namely *RIPK1*, *RIPK3*, and *MLKL*, are required for the degeneration of axons triggered by diverse degenerative stimuli and in a cell-autonomous manner.

Discussion

Axonal degeneration is an active self-destruction program that shares morphological and mechanistic features with necroptosis (Barrientos et al., 2011; Calixto et al., 2012; Villegas et al., 2014; Wallach et al., 2016). Here, we demonstrated that *RIPK1* inhibition with Nec-1 protects PNS as well as CNS axons from injury-induced degeneration, suggesting that necroptosis is a common degenerative program. Nec-1 has the potential to inhibit other

proteins, such as indoleamine-pyrrole 2,3-dioxygenase, p21 activated kinase-1, and protein kinase A (Cho et al., 2011; Eswaran et al., 2012; Takahashi et al., 2012; Vandenabeele et al., 2013), or to cause inactivation of other cell-death pathways, such as ferroptosis (Conrad et al., 2016). However, we showed by genetic interference of necroptosis activation and execution, through RIPK3 and MLKL knock down, respectively, delayed axonal degeneration in a cell-autonomous fashion, supporting our pharmacological results with Nec-1.

The progression of degeneration was associated with an early activation of necroptosis in axons, followed by mitochondrial fragmentation, which was also prevented by RIPK1 inhibition, indicating that necroptosis activation is triggered upstream of mitochondrial defects after axonal damage. Actually, inhibition of mitochondrial fission delayed axonal degeneration with the same protective magnitude as necroptosis inhibition, suggesting that both events converge in the same cascade for axonal degeneration.

Necroptosis is associated to axonal loss and functional impairment

Neurons contain at least two molecularly distinct cell-death programs, one for caspase-dependent apoptosis and another for selective axon degeneration. The last one does not require caspases (Finn et al., 2000; Wang JT et al., 2012).

The molecular mediators of necroptosis, RIPK1, RIPK3, and MLKL, regulate neuronal cell death for several diseases, including ALS (Re et al., 2014; Ito et al., 2016), multiple sclerosis (Dhib-Jalbut and Kalvakolanu, 2015; Ofengeim et al., 2015), AD, and PD (Caccamo et al., 2017; Iannielli et al., 2018). However, to date there is no direct and functional evidence that points out necroptosis as the main program controlling axonal self-destruction.

Many neuropathies are preceded by axonal degeneration, beginning at the distal ends of diseased axons, spreading retrogradely to the soma to trigger neuronal death (Cavanagh, 1979; Coleman, 2005). This suggests that neuronal viability depends on axonal integrity, which makes axons important targets for attenuating disease progression. In fact, inhibition of axonal degeneration protects from cell death and disease progression in several disorders (Ferri et al., 2003; Dadon-Nachum et al., 2011; Yang et al., 2017). This observation is quite relevant considering our results identifying necroptosis as a triggering factor for axonal degeneration by an autonomous-cell mechanism that specifically occurs within the axon. Our results are based on models of Wallerian degeneration; therefore, it will be important to determine whether necroptosis is involved in other non-Wallerian processes of axonal degeneration.

Necroptosis, originally described as a proinflammatory signaling pathway (Silke et al., 2015), occurs in many diseases. CNS diseases are no exception. It has been recently established that RIPK1 and RIPK3 are critical mediators of axonal degeneration in ALS by promoting inflammation and necroptosis activation in microglia (Ito et al., 2016). Similarly, necroptosis is required for astrocyte-triggered degeneration and death of motor neurons (Re et al., 2014; Fan et al., 2016). This evidence supports the role of necroptosis under neurodegenerative conditions. Our results showed that after injury, necroptosis is intrinsically activated within axons to trigger axonal degeneration.

RIPK3 is a key component for necrosome formation to trigger necroptosis (Newton et al., 2014; Orozco et al., 2014; Wu et al., 2014). Ablation of RIPK3 in *Fadd* or caspase-8-deficient mice rescues embryonic lethality (Kaiser et al., 2011) particularly by suppressing necroptosis during embryogenesis (Zhao et al., 2017).

Our RIPK3 knock-down studies propose necroptosis as a common pathway required for developmental processes under physiological conditions, but also for axonal destruction after injury or toxic insults. MLKL, the RIPK3 downstream effector (Sun et al., 2012), is responsible for the necrotic plasma membrane disruption, but also can translocate to other compartments as mitochondria (H. Wang et al., 2014), where it could modify oxidative phosphorylation to increase ROS levels (Zhao et al., 2012), events that also occur during axonal degeneration (Calixto et al., 2012). These antecedents point out necroptosis activation in convergence with mitochondrial dysfunction as potential triggering mechanisms of axonal degeneration.

Necroptosis links mitochondrial fragmentation to axonal degeneration

Mitochondria are considered central sensors for degenerative stimuli (Court and Coleman, 2012). After axonal injury, mitochondrial defects, such as mitochondrial potential loss and swelling, precede axonal fragmentation, making axons more vulnerable (Sievers et al., 2003; Barrientos et al., 2011; Park et al., 2013). Altered mitochondrial dynamics have also been related to pathogenesis of several neurodegenerative conditions (Knott et al., 2008; Grohm et al., 2012; Itoh et al., 2013). Inhibition of the main fission regulator, Drp1 (Smirnova et al., 2001; Otera and Mihara, 2011), delayed axonal degeneration. However, protection is assured. Since Drp1 also modulates mitochondrial function (Grohm et al., 2012), mitochondrial shortening might reflect a progressive organelle dysfunction that could lead to energetic failures and loss of axonal integrity. Conversely, the physiological role of Drp1 in neuronal function should be considered. Drp1 is essential for synapse formation (Ishihara et al., 2009) and for neuronal differentiation and migration (L. Wang et al., 2014), processes in which Drp1 inhibition also affects mitochondrial membrane potential (Kim et al., 2015). These antecedents could explain why axons are not fully protected from degeneration under Drp1 inhibition, and indeed confirm that regulation of mitochondrial fragmentation is a key step in the degenerative process.

Mitochondria-associated membranes (MAMs) constitute the contact sites between the endoplasmic reticulum and mitochondria during fission (Friedman et al., 2011) and have been proposed to contribute to mitochondrial dysfunction during axonal degeneration (Villegas et al., 2014). Interestingly, the active necrosome can translocate to MAMs, where necroptosis could be initiated (Chen et al., 2013). Consistently, the phosphatase PGAM5, a key regulator of Drp1, has been identified as a novel necrosome component (Wang Z et al., 2012). Our results indicate that necroptosis could mediate axonal degeneration by inducing mitochondrial reshaping. In fact, Drp1 and RIPK1 inhibition reached similar magnitudes of axonal protection, suggesting that they belong on the same pathway on which necroptosis is first activated to then induce mitochondrial fragmentation and axonal degeneration. This observation is sustained by results showing delayed mitochondrial fragmentation under RIPK1 inhibition, strengthening the idea that mitochondrial fission is a convergent point between programmed necrosis and axonal death. According to our results, loss of RIPK3 can rescue Drp1-knockout Purkinje neurons from degeneration, modulating mitochondrial morphology from large spheres into elongated tubules (Yamada et al., 2016). Pharmacological studies with Nec-1 have indicated that RIPK1 inhibition improves mitochondrial function after spinal cord injury (Y. Wang et al., 2015). However, genetic approaches like those we took in our work should be attempted to directly determine the mechanism involved in

necroptosis-mediated degeneration and its connection with mitochondrial reshaping.

The role of mitochondrial fragmentation and mPTP formation in necroptosis has been a subject of debate, especially considering that MLKL can directly permeabilize the plasma membrane leading to cellular damage. More recent evidence demonstrates that the executory mechanisms of necroptosis can be diverse and are highly dependent on cellular type and context. In sepsis-mediated kidney injury, necrosome formation can lead to mitochondrial dysfunction and ROS production by inhibiting mitochondrial complex I and III (Sureshbabu et al., 2018). In other cells types, necroptosis activation can enhance aerobic respiration and ROS production (Yang et al., 2018). As ROS can trigger mitochondrial fragmentation by diverse mechanisms, including Drp-1 activation (Abuarab et al., 2017; Riba et al., 2017), necroptosis-dependent mitochondrial fragmentation could represent an executioner mechanism for axonal degeneration after injury.

The identification of the prodegenerative gene *Sarm1* provided evidence indicating Wallerian degeneration as a program driven by a conserved axonal death pathway (Osterloh et al., 2012; Gerdtts et al., 2013; Geisler et al., 2016). It has been shown that mitochondrial dysfunction also stimulates a *Sarm1*-dependent cell-destruction pathway by a caspase-independent mechanism (Summers et al., 2014), suggesting the participation of necroptosis. However, *Sarm1* deletion fails to prevent changes in mitochondrial dynamics occurring after axotomy, despite delaying degeneration (Loreto et al., 2015). Thus, researchers still do not know whether *Sarm1* participates in or interacts with the necrosome to promote axonal death. A recent study showed that *Sarm1*-Toll-interleukin receptor-1 domain itself has intrinsic nicotinamide-adenine dinucleotide (NAD)ase activity, which was required to promote axonal NAD⁺ depletion and axonal degeneration after injury (Essuman et al., 2017). Moreover, Yang et al., showed that the necrosome can translocate to the mitochondria to activate pyruvate dehydrogenase, increasing ROS generation, which in turn enhances necrosome formation and necroptosis activation (Yang et al., 2018). These antecedents support our evidence showing that mitochondria converge as a central effector in the necroptotic cascade to initiate axon degeneration.

The present results show that necroptosis participates in the execution program of axonal degeneration as a ubiquitous and cell-autonomous mechanism, regulating both PNS and CNS axonal integrity and function. Furthermore, we reveal the participation of mitochondrial fragmentation as a downstream effector of the necroptotic pathway. Further studies are required to determine what are the main downstream effectors of necroptosis and mitochondria, and how they are inter-regulated to trigger axonal degeneration.

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