Development/Plasticity/Repair

p75NTR and DR6 Regulate Distinct Phases of Axon Degeneration Demarcated by Spheroid Rupture

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The regressive events associated with trophic deprivation are critical for sculpting a functional nervous system. After nerve growth factor withdrawal, sympathetic axons derived from male and female neonatal mice maintain their structural integrity for \sim 18 h (latent phase) followed by a rapid and near unison disassembly of axons over the next 3 h (catastrophic phase). Here we examine the molecular basis by which axons transition from latent to catastrophic phases of degeneration following trophic withdrawal. Before catastrophic degeneration, we observed an increase in intra-axonal calcium. This calcium flux is accompanied by p75 neurotrophic factor receptor-Rho-actin-dependent expansion of calcium-rich axonal spheroids that eventually rupture, releasing their contents to the extracellular space. Conditioned media derived from degenerating axons are capable of hastening transition into the catastrophic phase of degeneration. We also found that death receptor 6, but not p75 neurotrophic factor receptor, is required for transition into the catastrophic phase in response to conditioned media but not for the intra-axonal calcium flux, spheroid formation, or rupture that occur toward the end of latency. Our results support the existence of an interaxonal degenerative signal that promotes catastrophic degeneration among trophically deprived axons.

Key words: degeneration; DR6; p75NTR; pruning; spheroids

Significance Statement

Developmental pruning shares several morphological similarities to both disease- and injury-induced degeneration, including spheroid formation. The function and underlying mechanisms governing axonal spheroid formation, however, remain unclear. In this study, we report that axons coordinate each other's degeneration during development via axonal spheroid rupture. Before irreversible breakdown of the axon in response to trophic withdrawal, p75 neurotrophic factor receptor-RhoA signaling governs the formation and growth of spheroids. These spheroids then rupture, allowing exchange of contents \leq 10 kDa between the intracellular and extracellular space to drive death receptor 6 and calpain-dependent catastrophic degeneration. This finding informs not only our understanding of regressive events during development but may also provide a rationale for designing new treatments toward myriad neurodegenerative disorders.

Introduction

Throughout nervous system development, axons, synapses, and even entire neurons are initially overproduced and then refined

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through a period of organized culling (Purves and Lichtman, 1980; Hamburger and Oppenheim, 1990; Kantor and Kolodkin, 2003). Sympathetic neurons in the PNS rely on NGF-TrkA neurotrophic signaling to stabilize these components (Hendry and Campbell, 1976; Campenot, 1977; Purves et al., 1988; Yan et al., 2010; Gamage et al., 2017). The absence of sufficient neurotrophic

 $DR6^{-/-}$ mouse line; and Bruce Carter (Vanderbilt University) for providing the p75NTR functional blocking 9650 antibody.

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signaling permits prodestructive cues originating from TNF receptor superfamily members, such as p75 neurotrophin receptor (p75NTR), death receptor 6 (DR6), and TNF receptor 1a, to promote regressive events, such as cell death, synapse restriction, or axon degeneration in the PNS (Singh et al., 2008; Nikolaev et al., 2009; Olsen et al., 2014; Wheeler et al., 2014; Gamage et al., 2017). Axon degeneration is critical for proper nervous system wiring during development, but it is also a hallmark of several neural pathologies, such as Alzheimer's disease, amyotrophic lateral sclerosis, and injury (Saxena and Caroni, 2007). Both developmental and pathological degeneration occurs in two phases: latent and catastrophic (Wang et al., 2012; Kristiansen and Ham, 2014).

During latent phase, axons are morphologically indistinguishable from those receiving trophic support (Deckwerth and Johnson, 1993). Readdition of trophic factor or mild depolarization during the latent phase is capable of rescuing degeneration (Edwards et al., 1991). After latency, the catastrophic/execution stage of degeneration occurs in a rapid and near synchronous manner. During this period, axonal transport ceases, axons develop swellings, neurofilaments become fragmented, the cytoskeleton disintegrates, and debris is removed by recruited phagocytes (Luo and O'Leary, 2005; Saxena and Caroni, 2007). Importantly, this stage of degeneration represents the point of no return where reintroduction of neurotrophic factors does not rescue degeneration (Edwards et al., 1991). Although we and others have shown that, in the absence of NGF-TrkA signaling, p75NTR and DR6 mediate axon degeneration (Bamji et al., 1998; Singh et al., 2008; Nikolaev et al., 2009; Olsen et al., 2014; Gamage et al., 2017), the relative contributions of p75NTR and DR6 specifically to latent and catastrophic phases of degeneration have largely been overlooked.

How do axons transition from latent to catastrophic phases? It was shown that calcium signaling in degenerating axons promotes activation of calpain, the calcium-sensitive protease required for the disassembly of cytoskeletal elements (George et al., 1995; Avery et al., 2012; Vargas et al., 2015). During development, it is thought that compartmentalized calcium transients act as temporal and spatial cues to trigger dendrite pruning (Kanamori et al., 2013). Moreover, it was recently shown that axoplasmic calcium increases before the emergence of gross morphological changes in NGF-deprived DRG cultures (Johnstone et al., 2019), suggesting that intra-axonal calcium signaling could play a role in all phases of degeneration. However, whether death receptors act permissively to allow calcium-dependent irreversible fragmentation in response to trophic withdrawal remains an open question.

In this study, we demonstrate that, after trophic deprivation, intra-axonal calcium increases before catastrophic degeneration. This is accompanied by the formation of calcium-rich spheroids that grow and then rupture, releasing their contents ($\leq 10 \text{ kDa}$) to the extracellular space while allowing an influx of extracellular molecules (e.g., calcium) into the intra-axonal space. The prodegenerative molecules released into the extracellular space are capable of hastening entry of trophically deprived axons into the catastrophic phase of degeneration. We show that p75NTR promotes spheroid formation, intra-axonal calcium rise, and membrane rupture in a Rho-dependent manner. In contrast, DR6 is not required for spheroid formation and rupture. Moreover, DR6 is required to hasten entry into the catastrophic phase of degeneration in response to conditioned media from degenerating axons, whereas p75NTR is dispensable. Together, these data place p75NTR and DR6 upstream and downstream of spheroidal rupture, respectively. This is consistent with separable roles for these receptors in early and late phases of degeneration induced by trophic withdrawal.

Materials and Methods

Mice. All experiments were performed in compliance with the Association for Assessment of Laboratory Animal Care policies and approved by the University of Virginia Animal Care and Use Committee. All mice are on a C57BL/6J.129S mixed background. $p75NTR^{-/-}$ animals were purchased from The Jackson Laboratory. $DR6^{-/-}$ animals were a generous gift from Genentech. Males and females are mixed in all of our experiments.

Primary sympathetic neuronal cultures. Sympathetic neuron cultures were established as described previously (Deppmann et al., 2008). Briefly, neurons were obtained by dissociation of P0–P2 mouse superior cervical ganglia. These neurons (from each litter of pups) were plated in compartmentalized microfluidic devices in DMEM supplemented with 10% FBS, penicillin/streptomycin (1 U/ml), and 45 ng/ml of NGF purified from mouse salivary glands. Glia are removed from cultures using 5 $\mu\rm M$ cytosine arabinofuranoside (Sigma-Aldrich) for 48–72 h. For NGF deprivation, cultures were washed three times with media lacking NGF and then maintained in NGF-deficient media containing a neutralizing antibody (1 $\mu\rm g/ml$ anti-NGF antibody, Millipore) through designated time points at 37°C.

Fabrication and use of microfluidic devices. Microfluidic devices were generated as described previously (Park et al., 2006). These chambers were affixed to coverglass coated with poly-D-lysine (50 μ g/ml) and laminin (1 μ g/ml).

Immunocytochemistry. Immunocytochemistry was performed as previously described (Singh et al., 2008). Briefly, at the indicated times, axons were fixed in 4% PFA (w/v)/PBS at room temperature for 20 min, washed 3×5 min with 1× PBS, and blocked/permeabilized (5% goat serum, 0.05% Triton X-100 in PBS) for 1 h at room temperature. Axons were then incubated overnight at 4°C with primary antibody diluted in blocking buffer. Cells were then washed 3×5 min with 1× PBS and incubated with fluorescent secondary antibody for 1 h at room temperature. Cells were again washed with 1 × PBS three times and imaged using a fluorescent inverted microscope. The antibodies used in this study are mouse anti-Tuj1 (1:1000, Covance) and goat anti-mouse Alexa-488 (1: 800, Invitrogen). For active Rho staining, axons were incubated with 50 μg/ml of GST-Rhotekin-Rho binding domain (RBD) fusion protein in blocking solution for 1 h at room temperature, washed 3×5 min with $1\times$ PBS. Cells were then incubated with anti-GST (1:800, Sigma-Aldrich, G1160) diluted in blocking buffer for 1 h, washed 3×5 min with $1\times$ PBS, and incubated with fluorescent secondary antibody for 1 h at room temperature. All in vitro experiments were performed in triplicate with at least two microfluidic devices used for each condition.

Live imaging. Sympathetic neuron cultures were washed 3 times with DMEM/F-12, Phenol Red free, and incubated for 30 min at 37°C and 10% CO₂ with live imaging dyes diluted in DMEM/F-12, Phenol Red free. Cells were then imaged under Leica SP5 X confocal microscope in W.M. Keck Center at the University of Virginia. Axons in grooves of microfluidic chamber were imaged after NGF deprivation. For assessing flipping of phosphatidylserine (PS) on axonal spheroids, Annexin V red reagent (IncuCyte, 4641) was diluted in DMEM/F-12, Phenol Red free (1:200) after NGF deprivation. For membrane rupture, dextran dyes diluted in DMEM/F-12, Phenol Red free were added to the microfluidic chamber after 17 h of NGF deprivation. The dyes used in this study are Fluo-4 AM (1 µM, F14201) and Dextran Texas Red, neutral, 3 kDa (50 μ M, D3329), 10 kDa (50 μ M, D1828), and 70 kDa (50 μ M, D1830). All dyes were purchased from Thermo Fisher Scientific. Drugs used for manipulating spheroid formation are Z-VAD-FMK (50 μM, Enzo, ALX-260-020), calpain inhibitor III (20 μM, Millipore, 208722), actinomycin D (1 μg/ml, Sigma-Aldrich, A9415), CT04 (1 μg/ml, Cytoskeleton), cytochalasin D (10 μg/ml, Sigma-Aldrich, C2618), CN03 (1 μg/ml, Cytoskeleton Inc), LM11A-31 (2 ng/ml, Cayman Chemicals, 21982), and anti-BDNF (20 µg/ml, Millipore, AB1779SP); 9650 immune serum was a gift from Bruce Carter in Vanderbilt University.

Image processing and analysis. Axon degeneration in culture was quantified from β 3-tubulin-stained fluorescence images by counting the number of individual axons at the leading edge that had at least three

beads/blebs as described previously (Zhai et al., 2003). A blinded investigator counted 10 representative pictures of the axons, in two microfluidic chambers per condition/time point. On each image, 10 50 µm boxes were randomly assigned to single axons. The investigator took care not to box bundles of axons, which may confound analysis. Then the number of boxes, which had ≥3 beads/blebs were counted and categorized as degenerating axons; ≥80% degeneration was considered maximum degeneration, and equal to or <10% degeneration of axons was considered as minimum degeneration. The percentage of the total number of degenerating axons was calculated using Microsoft Excel. At least 300 total axons were counted for each condition. The SEM was considered as error. In live imaging, Ca^{2+} intensity ($\Delta F/F0$), the size (S/S0), and number of axonal spheroids were quantified in selected ROI (single axon or axonal spheroid) by Fiji software (Schindelin et al., 2012). Each experiment was repeated at least 3 times with separate litters of mouse pups of the same genotype.

Conditioned media experiments. Established compartmentalized recipient neuron cultures were deprived of NGF by washing with NGF-free DMEM three times and incubating at 37°C with a neutralizing anti-NGF antibody for 12 h. To collect NGF deprivation conditioned media (NDCM), donor neuron cultures were washed with NGF-free DMEM three times and treated with a neutralizing anti-NGF antibody at 37°C for 24 h. For each device, a volume of $\sim\!100~\mu l$ of NDCM was collected, including the media in the wells of the device on the axonal side and the media in the axonal channel. NDCM or control conditioned media were applied on recipient axons for 5 h.

Calcium measurement. Conditioned media were diluted to MilliQ water (1:20) and mixed thoroughly; 100 μ l of reaction mixture was made with 1 μ l HEPES, 2 μ l 1 mm Fluo-4 (20 mm working concentration, Invitrogen), 10 μ l diluted conditioned media, and 87 μ l water. Black 96-well plate was used in spectrophotometric assay. Eight CaCl₂ standards (2.54, 4.87, 9.75, 19.5, 39, 78, 156, and 313 μ m) were used to calculate standard curve for analyzing Ca ²⁺ concentration in conditioned media.

Exogenous calcium assay. Sympathetic neuron cultures were washed three times with DMEM supplemented with 10% FBS, penicillin/streptomycin (1 U/ml), and then incubated in NGF-deficient media containing a neutralizing antibody (1 μ g/ml anti-NGF antibody, Millipore) for 12 h. For exogenous calcium concentration <1.8 mm (DMEM), cells were washed 3 times with DMEM Ca $^{2+}$ -free media. CaCl $_2$ was added to DMEM Ca $^{2+}$ -free media to indicated final concentrations, which was then applied to cultures 37°C and 10% CO $_2$ for 5 h.

Experimental design and statistical analysis. Statistical analysis was performed in Prism 8.0 software (GraphPad). All measurements are shown as mean \pm SEM. For samples defined by one factor, data were compared by unpaired two-tailed t tests for two samples or one-way ANOVA with Tukey's post hoc multiple-comparisons test for ≥ 3 samples. For samples defined by two factors, data were compared by two-way ANOVA with Sidak's post hoc multiple-comparisons test. Sample size (n) was defined as the number of axons or axonal spheroids counted in live imaging experiment, or the number of independent cultures that were quantified in each experiment. The null hypothesis was rejected at the 0.05 level. p values < 0.05 are considered significant. The statistical test, sample size (n), and the p values are reported in the figure legends.

Results

Intra-axonal calcium increases in trophic factor-deprived axons and accumulates in spheroids before catastrophic degeneration

To study the kinetics of degeneration induced by trophic withdrawal, we cultured sympathetic neurons from P0-P2 mice in microfluidic devices, which separate soma and axons (Fig. 1A). After establishment in 45 ng/ml of NGF for 5–7 DIV, WT sympathetic neurons were globally or locally deprived of NGF with media containing anti-NGF function blocking antibody for indicated times. After global NGF deprivation, the majority of axons remain intact for $\sim\!18$ h as measured by microtubule integrity ($\beta\!3$ -tubulin staining). This period of sustained axonal integrity is

referred to as the latent phase of degeneration (Coleman, 2005). At the conclusion of the latent phase, the majority of axons rapidly degenerate from $4.4 \pm 1.5\%$ to $90.4 \pm 2.2\%$ of degeneration within 3 h (Fig. 1*B*, *C*). This is referred to as the catastrophic or execution phase of degeneration (Wang et al., 2012).

What are the molecular events that signal the transition from latent to catastrophic phase? Axonal calcium waves have been reported to be critical in promoting axon degeneration after axotomy (Vargas et al., 2015). Given the importance of the calcium-dependent protease, calpain, in developmental axon degeneration, we speculated that calcium waves may occur at the transition between latent and catastrophic phases (Yang et al., 2013). We examined axonal calcium dynamics using Fluo4-AM imaging of axons that were withdrawn from NGF. Multiple small calcium peaks were detected during the first 12 h, while the largest axoplasmic calcium flux that we observed occurred between 17 and 19 h after global trophic withdrawal (Fig. 1D), a time reflecting the transition window between latent and catastrophic phases (Fig. 1C). Notably, axonal calcium increases up to threefold over baseline by 17 h 50 min after global NGF deprivation (Fig. 1F). After this initial increase, axonal calcium levels decrease \sim 1.5-fold. We hypothesize that axonal calcium increase may be a hallmark of the transition between latent and catastrophic degeneration phases.

In addition to axonal apoptosis induced by global NGF deprivation, pruning is critical for refinement of the nervous system during development. To examine whether axon pruning has similar calcium dynamics in relation to degeneration kinetics, we selectively deprived NGF in the axon compartment of sympathetic neurons grown in microfluidic devices (Fig. 1A). The majority of axons remain intact until 46 h after local NGF deprivation, indicating the delayed latent phase compared with global NGF deprivation, which is consistent with previous reports (Chen et al., 2012) (Fig. 1 B, C). Calcium levels in locally NGF-deprived axons spiked before entry into the catastrophic phase of degeneration (Fig. 1D). This relationship between calcium spike and catastrophic degeneration is similar to what we observed in neurons globally deprived of NGF, albeit with an overall delay in degeneration kinetics.

Upon closer inspection of calcium dynamics within the entire axon, we observed that much of this calcium is concentrated in nascent spheroids on NGF-deprived axons (Fig. 1E). The formation of axonal spheroids/beads has been observed in not only trophic withdrawal models of developmental degeneration but also in pathological scenarios (Takahashi et al., 1997; Griffiths et al., 1998; Probst et al., 2000; Beirowski et al., 2010; Mejia Maza et al., 2019). However, whether these beads/spheroids have a function as degeneration progresses remains an open question. Interestingly, after observing the initial formation of calcium-rich spheroids, these structures increase in size from 4.1 \pm 0.5 μ m² to $19.6 \pm 2.7 \,\mu\text{m}^2$ (~500%) within 1 h (Fig. 1*G*). In addition to the increase in average spheroid size, the number of spheroids increased from 1 to 13 per 100 µm between 17 and 18 h after trophic withdrawal (Fig. 1H). Spheroidal calcium levels had a ~1.7-fold increase by 17 h 50 min after NGF deprivation (Fig. 1*G*). However, the spheroidal calcium levels varied depending on spheroid size. This value likely underestimates the increase in calcium since the spheroidal area measured increases with time. As such, we also normalized the areas used for the ROI to quantify the spheroidal calcium density and found the same trend (Fig. 11).

To determine whether intra-axonal calcium elevation is required for the formation of spheroids, we depleted axoplasmic calcium by BAPTA-AM and examined the morphological

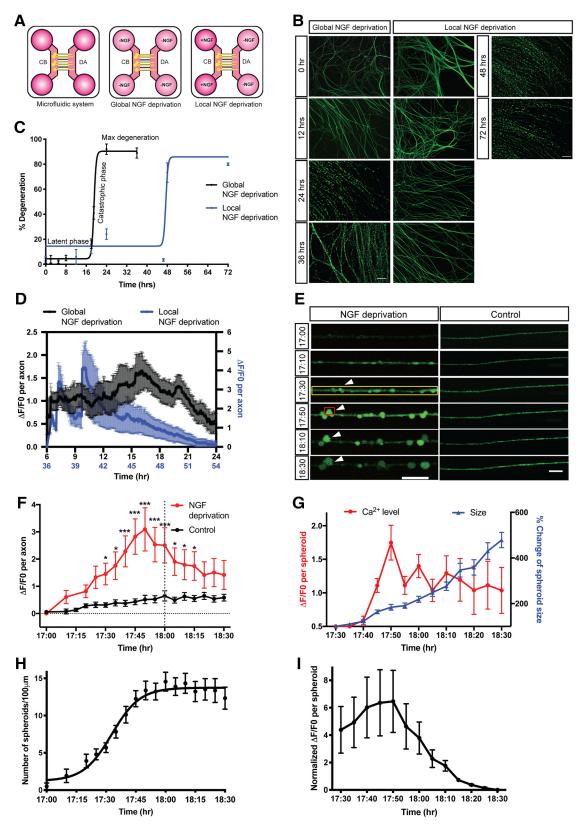


Figure 1. Axoplasmic calcium dynamics and formation of spheroids before catastrophic degeneration in response to NGF deprivation. **A**, Schematic representation of NGF deprivation paradigm in microfluidic devices. Cell bodies (CB) and distal axons (DA) are separated. For global NGF deprivation, sympathetic neurons were maintained in NGF-deficient media containing 1 μ g/ml anti-NGF antibody. For local NGF deprivation, sympathetic distal axons were maintained in 80 μ l per well of NGF-deficient media containing 1 μ g/ml anti-NGF antibody. Cell bodies were maintained in 150 μ l per well of media containing 45 ng/ml NGF. **B**, Representative images of β 3-tubulin-immunostained distal sympathetic axons before treatment (0 h) and 12, 24, 36, 48, and 72 h after global and local NGF deprivation. Scale bar, 50 μ m. **C**, Degeneration time course after NGF deprivation. Latent and catastrophic phases of degeneration are noted. Nonlinear regression curve was drawn according to the Hill equation. n = 3 for each time point. **D**, Normalized calcium fluorescence change of global (black, left y axis) and local (blue, right y axis) NGF-deprived axons over time. Total number of n = 6 (global NGF deprivation) and n = 11 (local NGF deprivation) axons from three independent litters were quantified. **E**, Fluo4-AM calcium (*Figure legend continues*.)

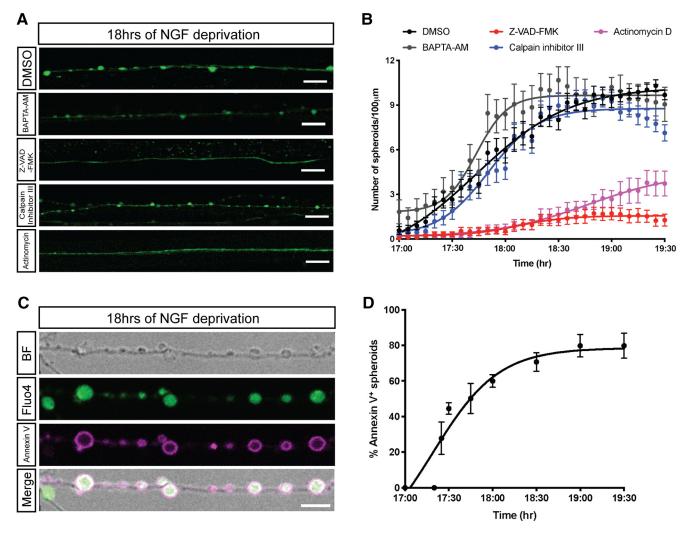


Figure 2. Transcription and caspase activation are required for formation of spheroids. *A, B,* Fluo4-AM calcium imaging (*A*) and quantification of axonal spheroid number per 100 μ m of sympathetic axons (*B*) at the indicated times after NGF deprivation in the presence of DMSO, 10 μ m BAPTA-AM, 50 μ m V-ZAD-FMK, 20 μ m calpain inhibitor III, and 1 μ g/ml actinomycin, respectively. Scale bar, 10 μ m. Nonlinear regression curves were drawn according to the Hill equation. Total number of n=26 (DMSO), n=26 (BAPTA-AM), n=25 (Z-VAD-FMK), n=20 (calpain inhibitor III), and n=17 (actinomycin) axons from three independent litters were counted. *C*, Representative images of axonal spheroids after 18 h of NGF deprivation. Fluo4-AM (green) indicates intra-axonal calcium, and annexin V (magenta) indicates exposure of PS on the outer leaflet on the spheroid membrane. Scale bar, 5 μ m. *D*, Quantification of the percentage of fluorescent annexin V-positive spheroids after 17–19.5 h of NGF deprivation. Total number of n=15 axons from three independent litters were counted. Data are mean \pm SEM.

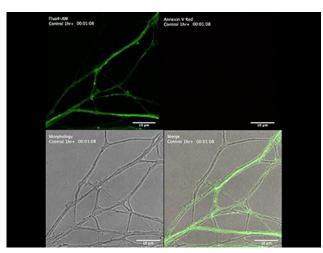
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(Figure legend continued.) imaging of sympathetic axons at the indicated times after global NGF deprivation. For the "NGF deprivation" condition, neurons were globally deprived of NGF for 17 h, and then incubated with Fluo4-AM for calcium imaging. For the "Control" condition, no NGF deprivation was performed. Yellow box represents individual axon as ROI. Red box represents axonal spheroid as ROI. White arrowheads indicate the formation and growth of spheroid. Scale bar, 10 μ m. **F**, Calcium fluorescence change of control or NGF-deprived axons over time. Gray vertical dotted line indicates the onset of catastrophic phase (18 h after NGF deprivation). Black horizontal dotted line indicates the baseline without any calcium change. Total number of n=8 (NGF deprivation) and n=18 (control) of axons from three independent litters were quantified. G, Calcium fluorescence and size change of axonal spheroid after 17 h of global NGF deprivation. Total number of n = 18 axonal spheroids from three independent litters were quantified. p = 0.0452 (17:30), p = 0.0059 (17:35), p < 0.0001 (17:40, 17:45, 17:50, 17:55, 18:00), p = 0.0037 (18:05), p = 0.0380 (18:10), p = 0.0293 (18:15); two-way ANOVA with Sidak's multiple-comparisons test. **H**, Quantification of axonal spheroid number per 100 μ m of axon at the indicated times after NGF deprivation. Nonlinear regression curve was drawn according to the Hill equation. Total number of n = 11 axons from three independent litters were counted. I, Quantification of normalized calcium fluorescence of axonal spheroids from 17.5 to 18.5 h of global NGF deprivation. Individual axonal spheroids were quantified: n = 18 spheroids from three independent replicates. Data are mean \pm SEM. *p < 0.05, ***p < 0.0001.

changes of NGF-deprived axons by live imaging. We find that axons retain their capacity to form spheroids in the absence of intra-axonal calcium elevation (Fig. 2A, B). These data indicate that, although intra-axonal calcium flux may correlate with entry into the catastrophic phase of degeneration, it is not required for spheroid formation.

Transcriptional induction of caspase activation is upstream of spheroid formation

Global NGF deprivation initiates apoptotic pathways to drive degeneration of axons and soma (Deckwerth and Johnson, 1993; Geden et al., 2019). Because caspases and calpain have been proposed to be involved in axon degeneration (Simon et al., 2012, 2016; Cusack et al., 2013; Ma et al., 2013; Yang et al., 2013), we next examined the requirement of these proteases in the formation of spheroids after NGF deprivation. Application of the pancaspase inhibitor Z-VAD-FMK, but not calpain inhibitor III, blocked spheroid formation (Fig. 2A). Less than two spheroids per 100 μ m of axons were observed in NGF-deprived cultures treated with Z-VAD-FMK, while the numbers of spheroids



Movie 1. Axoplasmic calcium dynamics without NGF deprivation. Fluo4-AM (green) and annexin V (red) live imaging of WT sympathetic axons with NGF. Scale bar, $10~\mu m$.



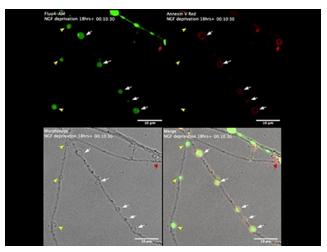
reached 10.2 \pm 0.4 and 8.8 \pm 0.2 per 100 μ m after 19 h of NGF deprivation in the DMSO and calpain inhibitor III treatments, respectively (Fig. 2*B*). Together, these data indicate that caspase activation is upstream of spheroid formation occurring at the transition between latent and catastrophic phases.

It has been suggested that transcriptional upregulation of proapoptotic proteins (e.g., PUMA) are required for axon degeneration after trophic withdrawal (Maor-Nof et al., 2016; Simon et al., 2016). To determine whether inhibition of early prodegeneration transcriptional events could delay spheroid formation, we treated cell bodies with a transcription inhibitor, actinomycin D, at the same time cells were deprived of NGF and examined spheroid formation between 17 and 19.5 h after trophic withdrawal. Interestingly, actinomycin D significantly diminished spheroid formation compared with the DMSO control (Fig. 2*A*,*B*). Based on previous reports, we speculate that a transcriptional program induced by NGF withdrawal is permissive for the activation of caspase pathways, which in turn promotes spheroid formation (Maor-Nof et al., 2016; Simon et al., 2016).

Once cells become apoptotic, plasma membrane PS asymmetry is lost and membranes undergo blebbing. Importantly, PS enrichment on the outer leaflet of the plasma membrane is read as an "eat me" signal for phagocytes to aid apoptotic cell recognition and clearance (Poon et al., 2014; Zhang et al., 2018). After NGF deprivation, the formation of axonal spheroids was detected by live imaging. We next assessed PS exposure on the extracellular surface of axonal spheroids by annexin V staining, which fluoresces upon binding to PS (Fig. 2C; Movies 1, 2). After 18 h of NGF deprivation, ~60% of spheroids, but not the rest of the axon, are annexin V-positive (Fig. 2D), suggesting PS flipping to the outer leaflet on the spheroid membrane.

Axonal spheroids develop membrane ruptures after NGF deprivation

Whereas annexin V staining is consistent with PS flipping to the outer leaflet of the plasma membrane, it is also formally possible that the membrane integrity of growing spheroids is disrupted, allowing access of the 45 kDa annexin V protein to the inner leaflet of the plasma membrane. To test this, we bathed control or



Movie 2. PS flipping of spheroidal membrane with NGF deprivation. Fluo4-AM (green) and annexin V (red) live imaging of WT sympathetic axons after 18 h of NGF deprivation. White arrows indicate calciumenriched spheroids with PS flipping. Red arrow indicates ruptured spheroids. Yellow arrowheads indicate calcium-enriched spheroids without membrane disruption. Scale bar, 10 μ m.



NGF-deprived axons in neutral fluorescent dextrans of different sizes (3, 10, 70 kDa). If there were any ruptures on the membrane after NGF deprivation, the dextran would immediately diffuse to the axoplasm (Fig. 3A). The exclusion of dextran was maintained until 17 h 40 min after NGF deprivation; however, by 17 h 50 min, the axoplasm began to fill with 3 kDa fluorescent dextran (Fig. 3B; Movies 3, 5). The fraction of labeled spheroids decreases with increasing size of fluorescent dextran, and no significant spheroidal uptake was observed with 70 kDa dextran (Fig. 3D), indicating that ruptures occur on a nanoscale and are permeable to molecules ≤10 kDa. Moreover, we observed nonpunctate dextran positivity of spheroids simultaneously as Fluo4-AM dye was lost (Fig. 3C), consistent with membrane rupture. We do observe some spheroids that have punctate dextran labeling, consistent with micropinocytosis; however, the events that we consider rupture are quite distinct and involve rapid and complete filling of the spheroid (Movie 4). If the labeling of annexin V on spheroids was due to entry of the dye through membrane rupture, the number of spheroids filled with dextran of similar or smaller sizes should be no less than the number of annexin V-positive spheroids. However, only $6.5 \pm 3.1\%$ of 10 kDa dextran-positive spheroids were observed after 18 h of NGF deprivation, significantly <45 kDa annexin V labeled spheroids (Figs. 2D, 3C), suggesting that the flipping of PS occurs before the entry of dextran or membrane rupture.

Rupture of axonal spheroids disrupts the electrochemical gradient and allows expulsion of axoplasmic material

Just as membrane rupture allows free diffusion of molecules \leq 10 kDa to enter spheroids, we predicted that these ruptures would allow small molecules to freely diffuse out of spheroids. Indeed, once spheroids reach \sim 400% of their original size, we observe a diminution of calcium signal, which we speculate is a function of diffusion of Fluo4-AM and calcium out of the spheroid. We sought to use calcium release to the extracellular space as a surrogate to test the hypothesis that small molecules may be released after spheroid rupture. We bathed axons in microfluidic devices in 100 μ l of calcium-free media and measured extracellular calcium before and after spheroid formation using a Fluo4 spectro-

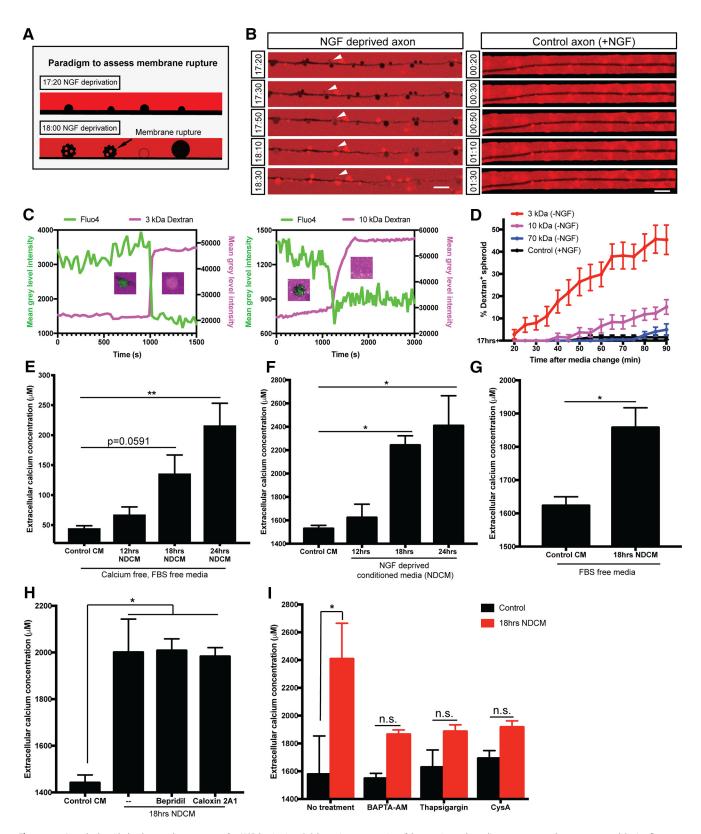


Figure 3. Axonal spheroids develop membrane rupture after NGF deprivation. *A*, Schematic representation of the experimental paradigm to assess membrane rupture model using fluorescent dextran. At 17 h 20 min of NGF deprivation, fluorescent dextran (red) is not taken up by the axon (black, negative space). However, by 18 h of NGF deprivation, as the plasma membrane loses integrity and ruptures, fluorescent dextran (red) can diffuse into spheroids, turning them red. Spheroids with intact membrane remain black. *B*, Representative images of dextran 3 kDa (red) entry to axonal spheroids (black) from 17 h 20 min to 18 h 30 min of NGF deprivation (left column), and dextran exclusion in untreated axons (right column). White arrowheads indicate that dextran 3 kDa enters axonal spheroids after 18 h of NGF deprivation. *C*, Examples of individual spheroidal tracing on NGF-deprived axons labeled with Fluo4-AM and bathed in 3 kDa (left) and 10 kDa (right) dextrans, respectively. Right, *y* axis indicates the mean gray level intensity of dextran fluorescence (magenta) after background subtraction. Left, *y* axis indicates the mean gray level intensity of Fluo4-AM (green) after background subtraction. Spheroids showing the red/green transition are captured. *D*, Quantification of the percentages of fluorescent 3 kDa (red), 10 kDa (magenta), and 70 kDa (blue) dextran-positive spheroids 20 –90 min after 17 h NGF deprivation. Black line (control) indicates the percentages of fluorescent 3 kDa dextran-positive (*Figure legend continues*.)



Movie 3. Dextran 3 kDa entry to axonal spheroids after NGF deprivation. Live imaging of dextran 3 kDa (red) entry to axonal spheroids (black) after 18 h of NGF deprivation. Yellow empty arrows indicate dextran 3 kDa entry to axonal spheroids. Scale bar, 10 μ m.



photometric assay. Sympathetic neurons were cultured in media containing FBS and 1.8 mM calcium, then switched to calcium-free, FBS-free media. We found that 24 h NDCM contained \sim 4-fold higher calcium (215.50 \pm 37.68 μ M) compared with Control CM (43.88 \pm 4.97 μ M) (Fig. 3*E*). Additionally, we were also able to observe an increase in extracellular calcium after spheroid rupture when experiments were performed using regular culture media containing FBS and 1.8 mM calcium. We found that, at 18 h, NDCM displayed a \sim 50% increase in calcium (2.24 \pm 0.08 mM) over Control CM (1.53 \pm 0.03 mM) (Fig. 3*F*). To rule out the

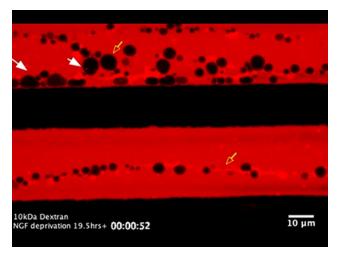
(Figure legend continued.) spheroids in the presence of NGF. Total number of n = 15 (3 kDa),

n = 15 (10 kDa), n = 12 (70 kDa), and n = 9 (control) axons from three independent litters were counted. **E**, Measurement of extracellular calcium expelled from axons into calcium-free, FBS-free media. All axons were grown in regular DMEM and then switched to calcium-free, FBS-free media before NGF deprivation. In the Control CM group, media were collected from axons grown in the presence of NGF. In NDCM groups, media were collected at 12, 18, and 24 h after NGF deprivation. Values were analyzed from n = 7 (Control CM), n = 5 (12 h NDCM), n = 76 (18 h NDCM), and n = 7 (24 h NDCM) independent replicates. Compared with Control CM, p =0.0591 for 18 h NDCM, p = 0.0003 for 24 h NDCM (one-way ANOVA with Dunnett's multiplecomparisons test). F, Measurement of extracellular calcium concentration of untreated and NGF-deprived conditioned media. Values were analyzed from n = 7 (Control CM), n = 4 (12 h NDCM), n = 6 (18 h NDCM), and n = 8 (24 h NDCM) independent replicates. Compared with Control CM, p=0.0227 for 18 h NDCM, p=0.0026 for 24 h NDCM (one-way ANOVA with Dunnett's multiple-comparisons test). **G**, Measurement of extracellular calcium extruded from axons into FBS-free, IS21 supplemented DMEM with or without 18 h of NGF deprivation. All axons were grown in FBS-free, IS21 supplemented media before NGF deprivation. n=8 (Control CM) and n=3 (18 h NDCM) independent replicates (p=0.0022, unpaired t test). H, Measurement of extracellular calcium concentrations in Control CM and 18 h NDCM in the absence and presence of NCX blocker 10 μ m Bepridril or PMCA blocker 0.5 mm Caloxin 2A1. n=4 for Control CM (p = 0.0038); n = 4 for 18 h NDCM); -(p = 0.0058); n = 3 for 18 h NDCM, Bepridril (p = 0.0079); n = 3 for 18 h NDCM, Caloxin 2A1 (one-way ANOVA with Tukey's multiple-comparisons test). I, Measurement of extracellular calcium concentration of untreated and NGF-deprived conditioned media in the presence and absence of intracellular calcium chelator BAPTA-AM (10 μ M, 1 h incubation before NGF deprivation), SERCA inhibitor thapsigargin (100 nm, overnight treatment before NGF deprivation), and mitochondrial mPTP blocker cyclosporin A (20 μ M, 1 h treatment before NGF deprivation). n=3 for each group. p=0.0375 for

No treatment; p = 0.7952 for BAPTA-AM; p = 0.8897 for thapsigargin; p = 0.9863 for CysA

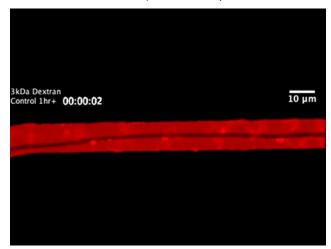
(two-way ANOVA with Sidak's multiple-comparisons test). Data are mean \pm SEM. *p < 0.05,

**p < 0.001. Scale bar, 10 μ m.



Movie 4. Dextran 10 kDa entry to axonal spheroids after NGF deprivation. Live imaging of dextran 10 kDa (red) entry to axonal spheroids (black) after 19.5 h of NGF deprivation. Yellow empty arrows indicate that dextran 10 kDa entry to axonal spheroids. White filled arrows indicate spheroids detaching from the axon. Some spheroids display punctate labeling consistent with macropinocytosis before filling, which is consistent with membrane rupture. Scale bar, 10 μ m.





Movie 5. Dextran 3 kDa exclusion of axons without NGF deprivation. Live imaging of dextran 3 kDa (red) exclusion of sympathetic axons in the presence of NGF. Scale bar, $10~\mu m$.



effect of FBS in regular culture media before calcium measurements, neurons were cultured in FBS-free, IS21-supplemented DMEM media. Calcium level in 18 h NDCM (1858.08 \pm 59.23 μ M) was significantly higher than calcium concentration in Control CM (1623.26 \pm 26.66 μ M) (Fig. 3G), suggesting that this calcium release occurred regardless of the presence or absence of FBS in the media.

Beyond membrane rupture, it is formally possible that calcium release could be in part due to enhanced permeability of plasma membrane Ca²⁺ ATPase (PMCA) and/or reversal Na⁺/Ca²⁺ exchanger (NCX) (Bano et al., 2007; Brini and Carafoli, 2011). To test this possibility, we treated NGF-deprived axons with PMCA inhibitor caloxin 2A1 or NCX inhibitor bepridil to test their ability to block elevation of extracellular calcium concentration. Axons deprived of NGF for 18 h and treated with

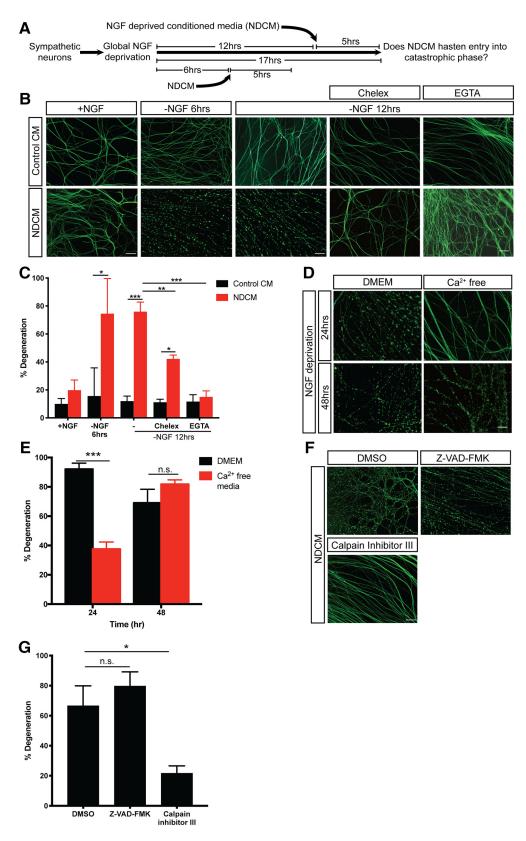


Figure 4. Calcium and calpain activation are required for NDCM-induced catastrophic axon degeneration. **A**, Conditioned media degeneration paradigm. WT sympathetic neurons were globally deprived of NGF for 6 or 12 h followed by addition of conditioned media to the axons for 5 h. NDCM was collected from degenerated axons after 24 h of NGF deprivation. **B**, **C**, Representative images (**B**) and quantification (**C**) of β3-tubulin-immunostained distal sympathetic axons after treatment with Control CM and NDCM for 5 h in the presence and absence of NGF, Chelex beads, and EGTA (6 mm). n = 4 or more for each group. For each repeat, at least 100 axons were scored for degeneration. Compared with Control CM, p = 0.0212 for - NGF 6 h, NDCM; p < 0.0001 for - NGF 12 h, Chelex, NDCM. Compared with - NGF 12 h, NDCM, p = 0.0002 for - NGF 12 h, Chelex, NDCM (two-way ANOVA with Sidak's multiple-comparisons test). **D**, **E**, Representative images (**D**) and quantification (**E**) of β3-tubulin-immunostained distal sympathetic axons after 24 and 48 h of NGF deprivation in media with calcium (DMEM) and calcium-free media. Axons were cultured in regular media (DMEM) and then switched to calcium-free media at the time of NGF deprivation. (*Figure legend continues*.)

caloxin 2A1 or be ridil retained their ability to release calcium to the extracellular space (Fig. 3*H*). Thus, spheroidal calcium is released to extracellular space independent of PMCA or NCX.

The observation that calcium in the media is elevated after membrane rupture is curious because the calcium concentration outside of the cell (1.8 mm) is known to be much higher than in the axoplasm (100 nm) (LaFerla, 2002; Clapham, 2007). How the axoplasm would have sufficient quantities of free calcium to appreciably elevate calcium levels in the media is unclear. To gain insight into this, we sought to determine the source of expelled intra-axonal calcium. We began by chelating all intracellular calcium with 10 μM BAPTA-AM added 1 h before NGF deprivation, which prevented calcium release to the extracellular space 18 h after trophic deprivation (Fig. 31). Intracellular calcium is buffered and stored by a contiguous network of ER and mitochondria, and calcium flux is thought to contribute to axon degeneration (Mattson, 2007; Villegas et al., 2014). Depletion of ER or mitochondria calcium, by treating neurons with 100 nm thapsigargin or 20 μ M cyclosporine A for 12 h before NGF deprivation, also blocked calcium increase in the extracellular space (Fig. 31). These results suggest that the elevated extracellular calcium comes from intracellular calcium stores.

We expect that, in physiological scenarios, membrane rupture would also allow disruption of the calcium electrochemical gradient between the inside and outside of the cell. In this way, elevated calcium level in the extracellular space may represent a surrogate for the release of other small molecules as well as a sustained elevation of intracellular calcium, which may lead to activation of calpain. Importantly, we are unable to observe sustained elevation of axoplasmic calcium after rupture (Fig. 1D) because the calcium indicator, Fluo4-AM, used in the above experiments would also diffuse out of the axon.

Expelled axoplasmic prodegenerative molecules hasten entry of axons into catastrophic phase

We speculated that contents released from ruptured spheroids may induce entry into the catastrophic phase of degeneration. To examine this, we collected media surrounding distal axons after 24 h of NGF deprivation (24 h NDCM) and incubated intact NGF-deprived axons for 5 h in 24 h NDCM (Fig. 4A). Importantly, in this paradigm, the total time that recipient axons are incubated without NGF are 11 and 17 h for "-NGF 6 h" and "-NGF 12 h" groups, respectively, before stereotypical morphological changes associated with degeneration. Neurons treated with Control CM (from untreated axons at time 0 of NGF deprivation) displayed minimal degeneration, which is consistent with the kinetics of the trophic deprivation time course presented in Figure 1C. However, we found that intact axons treated with NDCM displayed 74.5 \pm 25.1% and 76.0 \pm 6.8% degeneration at 11 and 17 h time points, respectively (Fig. 4B, C). Additionally, in the presence of NGF, NDCM failed to promote degeneration in 5 h

(Figure legend continued.) n=3 for each group. For each repeat, at least 100 axons were scored for degeneration. p<0.0001 for 24 h, p=0.4766 for 48 h (two-way ANOVA with Sidak's multiple-comparisons test). **F**, **G**, Representative images (**F**) and quantification (**G**) of β 3-tubulin-immunostained distal sympathetic axons after treatment with NDCM for 5 h in the presence of DMSO, 50 μ m Z-VAD-FMK, and 20 μ m calpain inhibitor III, respectively. All cultures were globally deprived of NGF for 12 h before NDCM incubation. Compared with DMSO (n=7), p=0.5328, n=9 for Z-VAD-FMK; p=0.0080, n=8 for calpain inhibitor III (one-way ANOVA with Dunnett's multiple-comparisons test). Data are mean \pm SEM. *p<0.005, ***p<0.001. Scale bar, 50 μ m.

of incubation (Fig. 4*B*, *C*), indicating that loss of NGF prosurvival signal is necessary to drive catastrophic axonal degeneration.

Extracellular calcium and calpain activation are required for entry into the catastrophic phase of degeneration

Small prodegenerative molecules may be released to extracellular space and reenter axoplasma via membrane rupture to promote degeneration. If sustained elevation of intracellular calcium is required for axons to trigger catastrophic degeneration, we would expect that blocking the prodegenerative-positive feedback step by depleting extracellular calcium would delay catastrophic degeneration. To this end, we performed trophic deprivation using calcium-free DMEM. NGF-deprived WT axons maintained in DMEM with 1.85 mm calcium showed classic degenerative hallmarks, such as beading, blebbing, and fragmentation by 24 h (Fig. 4D). Remarkably, when neurons were grown in calcium-free media, their rate of degeneration slowed. After 24 h of NGF deprivation, only 38% of axons were degenerated in calcium-free media; however, maximal degeneration was observed 48 h after NGF deprivation (Fig. 4D,E). Moreover, both chelex resin (which sequesters all divalent cations) and EGTA (which depletes calcium from NDCM) suppressed the ability of NDCM to promote degeneration in recipient axons (Fig. 4B,C). Thus, extracellular calcium is required to promote catastrophic degeneration in trophically deprived sympathetic axons. This is consistent with observations in trophically deprived sensory neurons and enucleated sympathetic neurons (George et al., 1995; Johnstone et al., 2019).

To explore the downstream effector of calcium influx during developmental axon degeneration, we applied NDCM to recipient axons in the presence of calpain inhibitor III or Z-VAD-FMK. Interestingly, after 5 h of incubation, only 21.6 \pm 5.0% of axons were degenerated in cultures treated with calpain inhibitor III, significantly less than the DMSO group (66.6 \pm 13.3%), whereas axons bathed in Z-VAD-FMK displayed 79.7 \pm 9.6% of degeneration (Fig. 4 F, G). These results indicate that calpain, instead of caspase, acts downstream of spheroid rupture and sustained calcium influx to promote catastrophic degeneration.

p75NTR, but not DR6, regulates the timing of spheroid formation and axoplasmic content expulsion to the extracellular space

Based on the above results, we have identified three events before catastrophic axon degeneration: (1) an intra-axonal calcium increase; (2) calcium-independent spheroid formation and membrane rupture; and (3) triggering of catastrophic degeneration by NDCM. What are the pathways upstream and downstream of these events? To address this, we examined the role of two TNFR family members, DR6 and p75NTR, which we and others have implicated in axon degeneration following trophic withdrawal (Park et al., 2010; Twohig et al., 2011; Olsen et al., 2014; Gamage et al., 2017).

We first examined whether p75NTR or DR6 acts upstream of spheroid formation and membrane rupture. To this end, we performed intra-axonal calcium imaging in NGF-deprived $DR6^{-/-}$ and $p75NTR^{-/-}$ sympathetic axons. Both WT and $DR6^{-/-}$ axons displayed up to a fivefold increase of spheroidal calcium levels; however, $p75NTR^{-/-}$ sympathetic axons did not display a significant increase in spheroidal calcium, even after 20 h 30 min of NGF deprivation (Fig. 5A,B). We next examined the role of p75NTR and DR6 in the accumulation of spheroids and the change in their size as a function of time after NGF deprivation. At 18 h after NGF deprivation, WT and $DR6^{-/-}$ neurons dis-

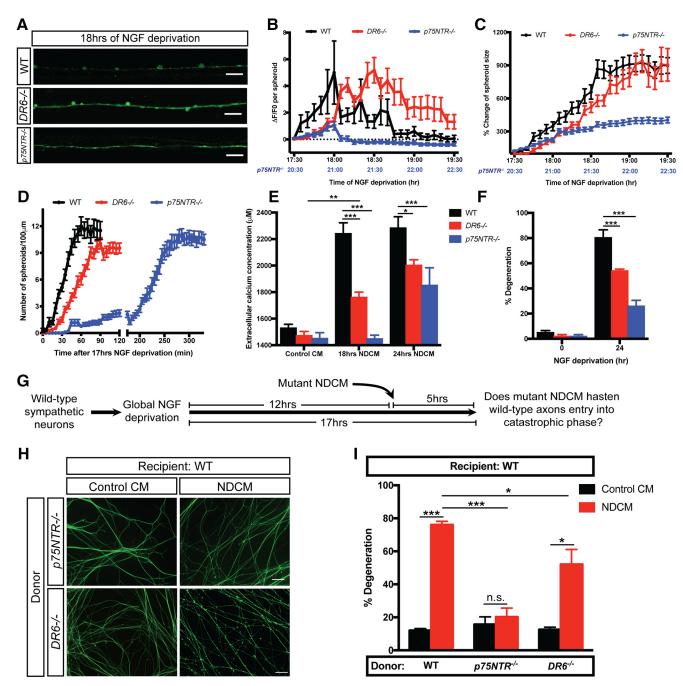


Figure 5. Depletion of p75NTR delays formation of spheroids and prodegenerative molecule exclusion after NGF deprivation. A, Fluo4-AM calcium imaging of WT, DR6-/-, and p75NTR-/sympathetic axons 18 h of NGF deprivation. Scale bar, 10 μ m. **B**, **C**, Spheroidal calcium fluorescence (**B**) and size change (**C**) of WT, $DR6^{-/-}$, and $p75NTR^{-/-}$ sympathetic axons after indicated time of NGF deprivation. B, Horizontal dotted line indicates the baseline without any calcium change. Lower x axis labeled with blue correlates with time of NGF deprivation in p75NTR^{-/-} sympathetic axons. Individual axonal spheroids were quantified: n = 32 (WT), n = 30 ($DR6^{-/-}$), and n = 22 ($p75NTR^{-/-}$) spheroids from cultured neurons harvested from three independent litters. **D**, Quantification of axonal spheroid number per 100 μ m of WT, $DR6^{-/-}$, and $D75NTR^{-/-}$ sympathetic axon at the indicated times after 17 h of NGF deprivation. Individual axonal spheroids were counted: n = 20 (WT), n = 29 (($DR6^{-/-}$), and n = 22 ($p75NTR^{-/-}$) axons from cultured neurons harvested from three independent litters. **E**, Measurement of extracellular calcium concentration of + NGF (Control CM), 18 and 24 h NDCM collected from WT, $DR6^{-/-}$, and $p75NTR^{-/-}$ sympathetic axons. n = 4 or more for each group. Compared with WT, 18 h NDCM, p < 0.0001 for $DR6^{-/-}$ $18 \text{ h NDCM}; p < 0.0001 \text{ for } p75NTR^{-/-}, 18 \text{ h NDCM}. \text{ Compared with WT, } 24 \text{ h NDCM}, p = 0.0026 \text{ for } DR6^{-/-}, 24 \text{ h NDCM}; p < 0.0001 \text{ for } p75NTR^{-/-}, 24 \text{ h NDCM}. \text{ Compared with } DR6^{-/-}, \text{Control of the property of the property$ CM, p = 0.0045 for $DR6^{-/-}$, 18 h NDCM (two-way ANOVA with Dunnett's multiple-comparisons test). F, Quantification of degeneration of WT, $DR6^{-/-}$, and $P75NTR^{-/-}$ sympathetic axons with or without 24 h of NGF deprivation. n = 3 or more for each group. For each repeat, at least 100 axons were scored for degeneration. Compared with WT, 24 h, p < 0.0001 for $DR6^{-/-}$, 24 h; p < 0.0001 for $DR6^{-/-}$. 0.0001 for p75NTR^{-/-}, 24 h (two-way ANOVA with Dunnett's multiple-comparisons test). **G**, Catastrophic axon degeneration paradigm to test the prodegenerative effect of mutant NDCM. WT sympathetic neurons were globally deprived of NGF for 12 h followed by addition of conditioned media derived from mutant axons for 5 h. Mutant NDCM was collected from degenerating p75NTR^{-/-} or DR6^{-/-} axons 24 h after NGF deprivation. H, I, Representative images (H) and quantification (I) of WT distal sympathetic axons immunostained for \(\beta\)3-tubulin after treatment with Control CM and NDCM collected from p75NTR^{-/-} and DR6^{-/-} axons for 5 h. Scale bar, 50 μ m. Left two columns represent percentages of degeneration of WT sympathetic axons treated with WT NDCM and Control CM, respectively. Compared with WT, NDCM (n = 10), p < 0.0001, n = 7 for $p75NTR^{-/-}$, NDCM; p = 0.0010, n = 7 for $DR6^{-/-}$, NDCM (two-way ANOVA with Dunnett's multiple-comparisons test). Compared with Control CM, p < 0.0001 for WT, NDCM; p = 0.8980 for $p75NTR^{-/-}$, NDCM, p = 0.0014 $DR6^{-/-}$, NDCM (two-way ANOVA with Sidak's multiplecomparisons test). Data are mean \pm SEM. *p < 0.05, **p < 0.001, ***p < 0.0001.

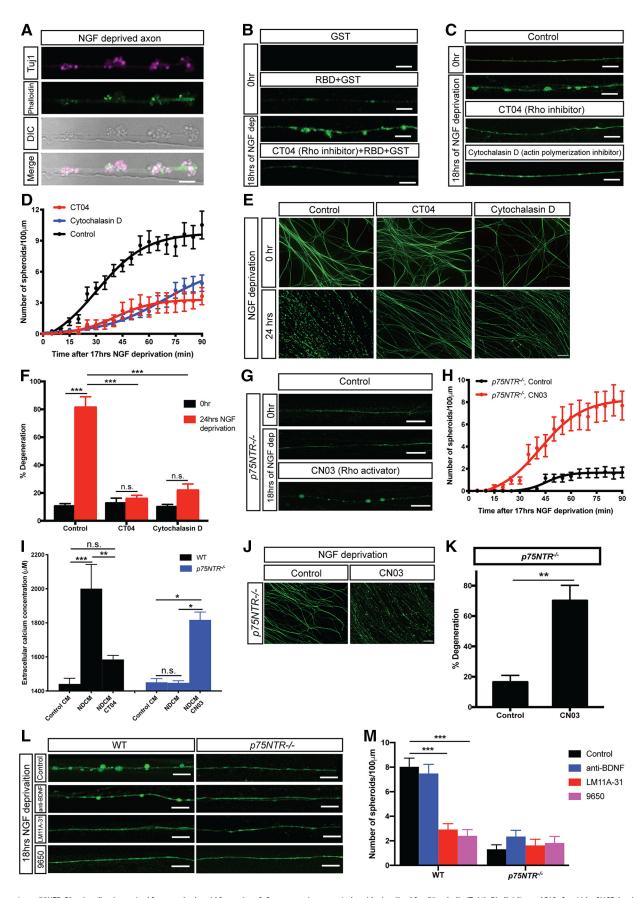


Figure 6. p75NTR-Rho signaling is required for axonal spheroid formation. **A**, Representative axons/spheroids visualized for β3-tubulin (Tuj1), Phalloidin, and DIC after 18 h of NGF deprivation. Scale bar, 5 μm. **B**, Representative images of WT sympathetic axons immunostained for GST tag with or without NGF deprivation. All images, except the first one, show axons incubated with Rhotekin-RBD GST-fusion protein that binds active Rho proteins after fixation. Bottom, Axon was NGF-deprived and treated with 1 μg/ml Rho inhibitor CTO4 for 3 h. (*Figure legend continues*.)

played 12.0 \pm 1.0 and 6.1 \pm 0.7 spheroids per 100 μ m of axons, respectively (Fig. 5 A, D). However, $p75NTR^{-/-}$ neurons displayed an \sim 2 h delay in spheroid formation (Fig. 5D), which is consistent with delayed onset of catastrophic degeneration (Gamage et al., 2017). We also observed that loss of DR6 displayed an \sim 800% increase in spheroid size after trophic deprivation similar to WT controls, whereas p75NTR^{-/-} neurons displayed a delayed accumulation of spheroids with a total size increase of <300% (Fig. 5C). If spheroid formation is a prerequisite for membrane rupture, NGF-deprived neurons isolated from p75NTR^{-/-} mice would show diminished extracellular calcium expulsion, whereas DR6^{-/-} would display normal calcium release, a surrogate for membrane rupture. Indeed, p75NTR^{-/-} axons showed no difference in extracellular calcium levels before or after 18 h of NGF deprivation, whereas DR6^{-/-} neurons displayed a modest but significant increase (Fig. 5E). We next examined whether NGFdeprived conditioned media from p75NTR^{-/-} or DR6^{-/-} neurons were capable of inducing degeneration of WT axons (Fig. 5G). Consistent with calcium release phenotypes, we found that NDCM derived from *p75NTR*^{-/-} axons was incapable of inducing degeneration in WT axons (Fig. 5H,I). In contrast, NDCM from $DR6^{-/-}$ neurons contains prodegenerative activity (Fig.

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(Figure legend continued.) Scale bar, 5 μ m. C, Fluo4-AM calcium imaging of WT sympathetic axons with or without drug treatment. For the CTO4 group, WT axons were incubated in SCG media containing 1 μ g/ml Rho inhibitor CT04, for 2 h before 17 h of NGF deprivation. For the cytochalasin D group, WT axons were incubated in SCG media containing 10 μ g/ml actin polymerization inhibitor for 2 h before 17 h of NGF deprivation. Scale bar, 10 μ m. **D**, Quantification of axonal spheroid number per 100 μ m of WT sympathetic axons at indicated time points after 17 h of NGF deprivation in the absence and presence of CT04 or cytochalasin D. Total number of n=9 (Control), n=8 (CT04), and n=22 (cytochalasin D) axons from cultured neurons harvested from three independent litters were quantified. Nonlinear regression curves were drawn according to the Hill equation. **E**, **F**, Representative images (**E**) and quantification (**F**) of WT distal sympathetic axons immunostained for β 3-tubulin in the absence and presence of CTO4 or cytochalasin D. Scale bar, 50 μ m. Compared with 0 h, p < 0.0001, n = 3 for Control, 24 h; p = 0.9256, n = 7 for CT04, 24 h; p = 0.1058, n = 6 for cytochalasin D, 24 h (two-way ANOVA with Sidak's multiple-comparisons test). Compared with Control, 24 h, p < 0.0001 for CT04, 24 h; p < 0.0001 for cytochalasin D, 24 h (two-way ANOVA with Dunnett's multiplecomparisons test). $\textbf{\textit{G}}$, Fluo4-AM calcium imaging of p75NTR $^{-/-}$ sympathetic axons grown in the presence or absence of NGF with or without CN03 treatment. For the CN03 group, $p75NTR^{-/-}$ axons were incubated in SCG media containing 1 μ g/ml Rho activator CN03 for 2 h before 17 h of NGF deprivation. Scale bar, 10 μ m. H, Quantification of number of spheroids per 100 μ m of p75NTR^{-/-} sympathetic axons at indicated time points after 17 h of NGF deprivation in the absence and presence of CN03. Individual axons were counted: n = 16($p75NTR^{-/-}$, Control) and n = 12 ($p75NTR^{-/-}$, CNO3) axons from three independent replicates. Nonlinear regression curves were drawn according to the Hill equation. I, Measurement of extracellular calcium concentration of untreated and NGF-deprived conditioned media in the presence and absence of CT04 or CN03 collected from WT (black) and $p75NTR^{-/-}$ (blue) sympathetic axons. Compared with WT, Control CM (n = 4), p < 0.0001, n = 4 for WT, NDCM; p = 40.4701, n = 6 for WT, NDCM CT04. Compared with WT NDCM, p = 0.0004 for WT, NDCM CT04. Compared with $p75NTR^{-/-}$, Control CM (n=4), p>0.9999, n=5 for $p75NTR^{-/-}$, NDCM; p=0.0088, n=3 for $p75NTR^{-/-}$, NDCM CN03. Compared with $p75NTR^{-/-}$, NDCM, p=0.00880.0055 for p75NTR^{-/-}, NDCM CN03 (one-way ANOVA with Tukey's multiple-comparisons test). **J**, **K**, Representative images (**J**) and quantification (**K**) of $p75NTR^{-/-}$ distal sympathetic axons immunostained for β 3-tubulin after treatment with or without CN03 for 5 h. All cell cultures were pretreated with 12 h of NGF deprivation. Scale bar, 50 μ m. p = 0.0006, n = 3 for CN03 and n = 7 for control (unpaired t test). L, M, Fluo4-AM calcium imaging (L) and spheroid number per 100 μ m of WT and p75NTR $^{-/-}$ sympathetic axons (M) after 18 h of NGF deprivation in the absence and presence of 20 µg/ml anti-BDNF, 2 ng/ml p75NTR ligand/functional blocker LM11A-31, or 9650 immune serum (1:100). Scale bar, 10 μ m. Individual axons were counted: n = 15 (WT, Control), n = 16 (WT, anti-BDNF), n = 37 (WT, LM11A-31), n = 22 (WT, 9650), n = 16 (p75NTR^{-/-}, Control), n = 19 (p75NTR^{-/-}, anti-BDNF), n = 18 $(p75NTR^{-/-}, LM11A-31)$, and n = 16 $(p75NTR^{-/-}, 9650)$ axons from three independent replicates. Significance is determined by two-way ANOVA with Dunnett's multiplecomparisons test. Data are mean \pm SEM. *p < 0.05, **p < 0.001, ***p < 0.0001.

5*H*,*I*). Together, these data suggest that p75NTR, but not DR6, is upstream of spheroid formation and membrane rupture.

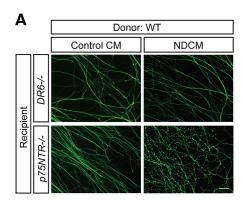
p75NTR-Rho signaling is necessary and sufficient for spheroid formation and entry into catastrophic phase

How does p75NTR regulate spheroid formation before the catastrophic phase of degeneration? Previous work has shown that neurotrophins regulate growth cone dynamics by suppressing RhoA activation that occurs through p75NTR signaling (Gehler et al., 2004). p75NTR promotes RhoA activation to mediate sympathetic axonal degeneration in response to degenerative triggers, such as myelin (Yamashita and Tohyama, 2003; Park et al., 2010). The major downstream target of Rho activation is actin remodeling. We observed actin and β 3-tubulin enrichment in axonal spheroids (Fig. 6A), similar to previous observations (Beirowski et al., 2010). Active Rho in axonal spheroids was detected by incubation with GST-rhotekin RBD fusion protein after 18 h of NGF deprivation but not in cultures treated with Rho inhibitor C3 transferase (CT04) (Fig. 6B). Moreover, inhibiting Rho family members using CT04 (1 μ g/ml, 2 h before 17 h NGF deprivation) prevented spheroid formation in WT axons (Fig. 6C,D). We next tested whether actin remodeling is required for spheroid formation. Indeed, the actin polymerization inhibitor cytochalasin D (10 μ g/ml) inhibited spheroid formation (Fig. 6*C*,*D*). Consistent with this, axons treated with CT04 or cytochalasin D delayed degeneration for up to 24 h after NGF deprivation (Fig. 6E,F). Together, these data suggest that activation of Rho and actin remodeling is necessary for spheroid formation and entry into the catastrophic phase of degeneration. Another downstream effector of Rho is the Rho-associated protein kinase 1. During apoptosis, Rho-associated protein kinase 1 has been shown to be cleaved by caspase-3, resulting in constitutive kinase activity to promote phosphorylation of myosin light chain and membrane blebbing (Coleman et al., 2001; Sebbagh et al., 2001). This may explain our finding of suppressed spheroid formation after NGF deprivation in the presence of caspase inhibitor (Fig. 2*A*, *B*).

If Rho is downstream of p75NTR, ectopic activation of Rho would promote spheroid formation and axon degeneration in the absence of p75NTR. To test this, we incubated NGF-deprived $p75NTR^{-/-}$ axons with Rho activator, CN03 (1 μ g/ml, 2 h before 17 h NGF deprivation). CN03 rescued the diminished spheroid formation and extracellular calcium release through membrane rupture in $p75NTR^{-/-}$ axons 18 h after NGF deprivation (Fig. 6G–I). We also found that CN03 could rescue the timing of degeneration in the absence of p75NTR (NGF deprived for 12 h with CN03 for the final 5 h) (Fig. 6J,K). Together, these data suggest that p75NTR-Rho signaling is necessary and sufficient to promote spheroid formation and entry into the catastrophic phase of degeneration.

p75NTR-dependent spheroid formation is ligand-dependent

We next sought to determine whether p75NTR requires ligand to induce spheroid formation. Application of LM11A-31, a small nonpeptide p75NTR modulator (Massa et al., 2006; Simmons et al., 2014), suppressed spheroid formation on WT sympathetic axons after NGF deprivation (Fig. 6L,M). Since there have been conflicting reports about whether LM11A-31 specifically blocks ligand binding (Massa et al., 2006; Xie et al., 2019), we also used the ligand-blocking antibody 9650 specific for p75NTR extracellular domain (Kraemer et al., 2014). Compared with control serum, 9650 significantly inhibited spheroid formation on WT sympathetic axons after NGF deprivation and did not show additive effect on $p75NTR^{-/-}$ axons (Fig. 6L,M). However, block-



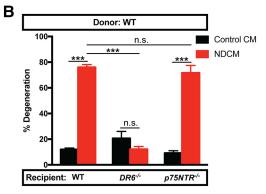


Figure 7. DR6 is required for catastrophic degeneration induced by NDCM. **A, B,** Representative images (**A**) and quantification (**B**) of distal sympathetic axons from $DR6^{-/-}$ and $p75NTR^{-/-}$ animals immunostained for β3-tubulin after treatment with NDCM and Control CM collected from WT neurons for 5 h, respectively. All cultures were NGF-deprived for 12 h before the addition of CM. Left two columns represent percentages of degeneration of WT sympathetic axons treated with WT NDCM and Control CM, respectively. Compared with NDCM, p < 0.0001, n = 10 for WT, Control CM; p = 0.5437, n = 3 for $DR6^{-/-}$, Control CM; p < 0.0001, n = 7 for $p75NTR^{-/-}$, Control CM, two-way ANOVA with Dunnett's multiple-comparisons test. Compared with WT, NDCM (n = 10), p < 0.0001, n = 9 for $DR6^{-/-}$, NDCM; p = 0.5422, n = 11 for $p75NTR^{-/-}$, NDCM, two-way ANOVA with Dunnett's multiple-comparisons test. Data are mean \pm SEM. ****p < 0.0001. Scale bar, 50 μ m.

ing BDNF, which has been shown to activate p75NTR apoptotic signaling (Kohn et al., 1999), failed to inhibit development of spheroids on NGF-deprived axons (Fig. 6L,M). These results suggest that p75NTR-mediated spheroid formation can be activated by binding prodegenerative ligands other than BDNF.

DR6, but not p75NTR, gates entry into the catastrophic phase of degeneration in response to NDCM

We next asked whether DR6 or p75NTR is downstream of the prodegenerative activity of NDCM using the paradigm described in Figure 4A. We applied NDCM collected from WT axons to recipient neurons derived from $DR6^{-/-}$ and $p75NTR^{-/-}$ mice. After a 5 h treatment of NDCM, $p75NTR^{-/-}$ axons showed 71.6 \pm 5.8% degeneration, whereas $DR6^{-/-}$ axons displayed minimal degeneration (Fig. 7*A*, *B*). These data suggest that DR6, but not p75NTR, is downstream of the prodegenerative effects of NDCM.

Discussion

By investigating calcium dynamics during the transition from latent to catastrophic phases of degeneration after trophic withdrawal, we identified several novel events: (1) Intraaxonal calcium rise and spheroid formation between latent and catastrophic phases of degeneration. This spheroid formation is triggered by p75NTR-Rho signaling, caspase activation, and actin remodeling. The timing of p75NTR-Rho-actin signaling and upregulation of prodegenerative transcriptional programs may define the duration of the latency phase. (2) The membranes of growing spheroids rupture, leading to expulsion of axoplasmic material to the extracellular environment. This may represent a novel mechanism by which prodegenerative molecules are released to influence neighboring cells. (3) The material expelled from ruptured spheroids triggers catastrophic degeneration. DR6 is required for responding to this prodegenerative signal, thereby gating entry into the catastrophic phase of degeneration (Fig. 8).

Formation of axonal spheroids has been reported in many neurological disorders, including Alzheimer's disease, amyotrophic lateral sclerosis, and neurocysticercosis (Griffin and Watson, 1988; Beirowski et al., 2010; Mejia Maza et al., 2019). After injury or oxidative stress, spheroids arise along the length of axons with accumulation of cytoskeleton degradation products and calcium overload (Beirowski et al., 2010; Barsukova et al., 2012). While spheroids have been extensively described in the literature,

whether they have a function in promoting degeneration has remained unknown. We propose that the function of these spheroids is to trigger catastrophic degeneration by releasing prodegenerative molecules into the extracellular environment. Here, we report localized annexin V positivity (i.e., PS flipping) (Fig. 2C), which suggests that they are marked for phagocytic engulfment reminiscent of apoptotic bodies (Ravichandran, 2010; Segawa et al., 2014). Indeed, we observed several instances of spheroids detaching from the axon, which is a unit that presumably would be phagocytosed (Movie 4). Importantly, the exposure of PS on the outer leaflet of the spheroid membrane, but not the axon shaft, indicates that spheroids might be "hot-spots" on degenerating axons to recruit nearby glia or macrophages, clearing damaged components. In correlation with the growth of spheroids, we also observe that 3 and 10 kDa fluorescent dextran entering the cytosol consistent with disrupted or ruptured plasma membrane (Fig. 3B-D). It is unclear where membrane integrity is disrupted; it may be at the belly of the spheroid and/or at axon/spheroid junctions. Whether or not spheroid rupture also occurs in aforementioned neurodegenerative disorders remains an open question.

What signaling pathways are upstream of spheroidal rupture? We found that disrupting caspase activation, p75NTR, Rho, or actin polymerization delays spheroid formation after NGF deprivation (Figs. 2A, B, 6C,D). The intracellular domain of p75NTR has been shown to modulate RhoA activation by interacting with Rho GDP-dissociation inhibitor to inhibit axonal outgrowth and promote axon pruning (Yamashita et al., 1999, 2002; Park et al., 2010). In the presence of NGF, the interaction of p75NTR with Rho GDP-dissociation inhibitor-RhoA complex is disrupted, leading to RhoA inactivation and neurite outgrowth (Mathew et al., 2009). Consistent with these findings, we show that activation of RhoA in the absence of p75NTR promotes entry into catastrophic phase by triggering spheroid formation (Fig. 6G–K). Proneurotrophin activation of p75NTR in the absence of Trk signaling leads to caspase-mediated neuronal death (Troy et al., 2002; Gentry et al., 2004). Sublethal executioner caspase activity plays an important role in nervous system development (Unsain and Barker, 2015). Interestingly, inhibition of caspases suppressed spheroid formation but failed to protect axons from NDCM-induced catastrophic degeneration (Fig. 4F, G). Thus, in the absence of NGF, spheroid formation is induced by p75NTR,

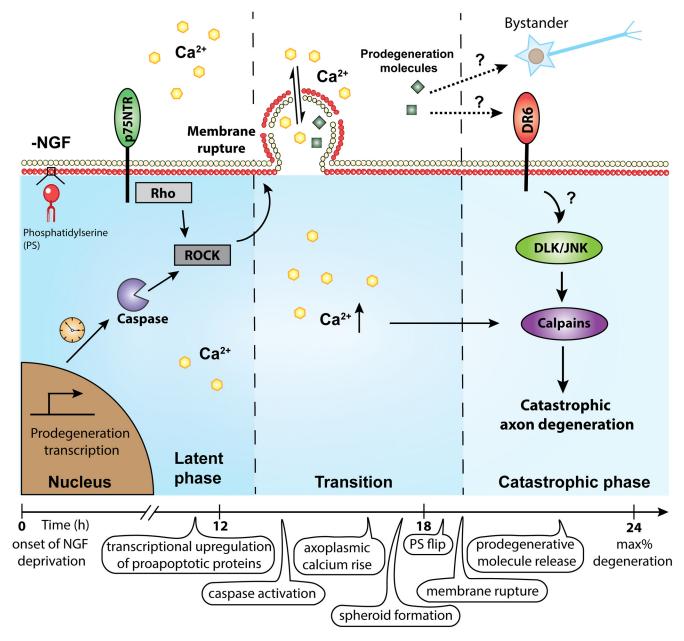


Figure 8. Time course of events associated with NGF deprivation-induced axon degeneration of sympathetic neurons. After NGF deprivation, prodegenerative transcription is upregulated. Axoplasmic calcium is increased and enriched in spheroids before catastrophic phase. Spheroid formation is regulated by p75NTR, Rho activity, and caspase activation. The calcium electrochemical gradient across the membrane is disrupted by spheroidal rupture, which may also lead to the release of intra-axonal prodegenerative molecules to extracellular space, acting as extrinsic factors to promote degeneration in a paracrine or autocrine manner. We speculate that p75NTR plays an important role in calcium dynamics during the latent phase, whereas DR6 can be activated by prodegenerative NDCM to mediate downstream catastrophic degeneration pathways (e.g., DLK/JNK, calpastatin, calpain).

caspase, RhoA, and actin remodeling, which are all requisite for transitioning from latent to catastrophic degeneration.

Our findings are consistent with the notion that p75NTR controls the window of latency before catastrophic degeneration. What is the molecular basis for the putative timer controlling the window of latency after trophic withdrawal? Classically, biological timers have been associated with transcriptional programs that form feedback loops (Mitrophanov and Groisman, 2008; Santos and Ferrell, 2008). Inhibition of transcription by applying actinomycin D to the soma during NGF deprivation blocked spheroid formation and degeneration (Fig. 2*A*, *B*), indicating the involvement of a degenerative transcriptional program. This pathway has been shown to upregulate TNF α -converting enzyme, which is necessary for p75NTR cleavage (Kenchappa et al.,

2010). Liberation of the p75NTR intracellular domain retrograde degenerative signal is accompanied by secondary activation of JNK and upregulation of proapoptotic factors, such as p53-upregulated modulator of apoptosis (Kenchappa et al., 2010; Simon et al., 2016; Pathak et al., 2018). The time it takes to engage this transcriptional program and cleave p75NTR by TNF α -converting enzyme and γ -secretase may represent a molecular timer governing the duration of the latency window (Pathak and Carter, 2017). This molecular timer may account for the delay of spheroid formation in NGF-deprived $p75NTR^{-/-}$ axons (Fig. 5D).

What signaling pathways are downstream of spheroidal rupture? Similar to p75NTR, DR6 is required for axon degeneration after trophic withdrawal (Olsen et al., 2014; Gamage et al., 2017). Does DR6 regulate the same signaling events as p75NTR after

trophic factor deprivation? We conclude that this is unlikely given that these TNFR family members govern distinct phases of degeneration. Unlike $p75NTR^{-/-}$, $DR6^{-/-}$ axons are capable of expelling prodegenerative axoplasmic materials to the surrounding environment via membrane rupture after NGF deprivation (Fig. 5H,I). $DR6^{-/-}$ axons are protected from entry into catastrophic phase in response to a 5 h incubation of NDCM (Fig. 7 A, B). These findings suggest that DR6 is downstream of spheroidal rupture and the destructive factor(s) found in NDCM; however, whether there is a direct or indirect interaction between these factors and DR6 remains to be determined. DR6 was previously shown to interact with β -APP to regulate developmental axonal pruning and synapse restriction (Nikolaev et al., 2009; Kallop et al., 2014). Given the fact that depletion of APP could partially protect sensory axons from degeneration after NGF withdrawal (Olsen et al., 2014), it is possible that DR6 promotes developmental catastrophic degeneration via an APP-dependent pathway. Previous findings show that DR6 potently activates NF-κB and JNK to induce apoptosis (Benschop et al., 2009; Hu et al., 2014). Consistent with this, loss of DR6 suppressed phosphorylation of JNK after injury (Gamage et al., 2017). Thus, we speculate that direct or indirect activation of DR6 by the prodegenerative factor emanating from ruptured spheroids may promote the activation of JNK and calpain, which are known to trigger catastrophic degeneration (Fig. 8). It is also possible that activation of DR6 independently activates catastrophic degeneration by engagement of receptor-mediated caspase pathways.

Although the calcium-dependent protease, calpain, is known to be required for catastrophic degeneration (Fig. 4F, G), it is unclear why the initial intracellular wave of calcium before catastrophic degeneration is insufficient to trigger this event. One clue may come from Tessier Lavigne and colleagues who suggest that depletion of the calpain inhibitor, calpastatin, is required for catastrophic degeneration (Yang et al., 2013). The membrane ruptures that we observe at the onset of catastrophic degeneration is likely to cause equilibration of external and internal calcium concentrations. After rupture, intraxonal calcium levels are likely to be persistently high. However, depletion of calpastatin is likely prerequisite for calcium-dependent catastrophic degeneration. Recent studies show that chelation of extracellular calcium by EGTA in late, but not early, phases of degeneration rescues axons from trophic deprivation-induced degeneration (Johnstone et al., 2019). Additionally, the transient receptor potential vanilloid family member 1 cation channel is required for calcium influx to promote developmental sensory axon degeneration, whereas plasma membrane nanoruptures allow entry of extracellular calcium to drive axon degeneration in multiple sclerosis (Johnstone et al., 2019; Witte et al., 2019). Thus, extracellular calcium may enter the axoplasma via calcium channels, such as transient receptor potential vanilloid family member 1 or spheroidal ruptures. Both routes of sustained intracellular calcium elevation converge on calpain to trigger catastrophic degeneration.

In conclusion, our data reveal novel signaling emanating from two different death receptors to govern latent and catastrophic phases of degeneration. After NGF deprivation, prodegenerative transcription, caspase activation, and p75NTR-Rho-dependent actin remodeling promote formation of axonal spheroids. Axoplasmic materials containing prodegenerative molecules, such as calcium, are then released to the extracellular space via membrane rupture, which may act as a positive feedback loop to hasten the entry of axons into catastrophic phase degeneration. Entry into the catastrophic phase of degeneration appears to be gated by DR6 (Fig. 8). Whether or not this represents a mech-

anism whereby axons might coordinate their degeneration requires further study. Nevertheless, the notion of destructive spheroids and subsequent interaxonal communication during degeneration facilitates our understanding of neural refinement during development and presents intriguing possibilities with respect to therapeutic intervention to alleviate bystander degeneration in disease.

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