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Mechanistic Advances in Axon Pathfinding

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Abstract

The development of a functional nervous system entails establishing connectivity between appropriate synaptic partners. During axonal pathfinding the developing axon navigates through the extracellular environment, extending toward postsynaptic targets. In the early 1900s, Ramon y Cajal suggested that the growth cone, a specialized, dynamic, and cytoskeletal-rich structure at the tip of the extending axon, is guided by chemical cues in the extracellular environment. A century of work supports this hypothesis and introduced myriad guidance cues and receptors that promote a variety of growth cone behaviors including extension, pause, collapse, retraction, turning, and branching. Here we highlight research from the last two years regarding pathways implicated in axon pathfinding.

Introduction

Formation of appropriate neural networks requires that axons navigate to and connect with appropriate targets. Defining how the growth cone at the tip of the extending axon accomplishes its vast repertoire of responses associated with establishing neuronal connectivity remains an area of intense research. Growth cone motility and axonal pathfinding involve spatial and temporal remodeling of growth cone architecture resulting in mechanotransduction and motility (Fig. 1). Attractive and repulsive guidance cues interact with receptors on the axon surface and promote or constrain growth, respectively¹. Diffusion of soluble chemotactic cues influence axon dynamics over long distances, whereas interaction of the axon with adherent, haptotactic guidance cues influences dynamics locally.

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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These cues activate a host of diverse signaling pathways, which results in cytoskeletal and membrane remodeling within the growth cone. Here we review recent work on axonal pathfinding, with attention to the combinatorial synergy of pathways and parallels to morphogenesis of the dendritic spine.

Guidance cues direct the growth cone

Axon pathfinding begins with the guidance cues that direct growth. Over the last 30+ years, a host of guidance cues have been identified, classically divided into four families: semaphorins, ephrins, netrins, and slits. Identification of additional cues such as sonic hedgehog (Shh), Wnt, and growth factors provides additional complexity and control to axonal wiring². This list continues to grow: for example, recent work showed that the extracellular domains of the trans-synaptic adhesion molecule lasso (teneurin-2) are cleaved in rat brain, releasing soluble lasso³. As this suggested a novel, pre-synaptic role for lasso, subsequent work in cultured hippocampal neurons demonstrated soluble lasso binds growth cone localized latrophilin-1, initiating calcium signaling and attractive axon turning⁴. Identification of additional guidance cues and receptors will continue to illuminate how precise directional growth cone extension is tuned.

Beyond identification of cues and receptors, our understanding of axon pathfinding complexity continues to increase. Numerous cues originally proposed to promote a single axonal behavior are now known to promote multiple behaviors based on context, such as target cell receptors and extracellular factors (Fig. 1A). For example, netrin acts as an attractive or repulsive cue based on receptors, concentration, and the extracellular environment⁵⁻¹⁰. Furthermore, while netrin was historically described as a soluble cue that promotes growth over long distances^{11,12}, recent work suggests netrin exhibits local, haptotactic influences^{10,13-16}. These new studies may suggest that netrin has only an adherent haptotactic role, or alternatively that there is production of distinct pools of netrin in different brain regions; with netrin produced in the ventral zone promoting haptotaxis¹⁷ and floor plate netrin diffusing, promoting chemotaxis¹⁸. The diverse responses to a single guidance cue, the rich variety of guidance cues and receptors, and the variable downstream signaling cascades together highlight the combinatorial computation occurring within the neuron. The output of this computation is exquisite axon pathfinding tunability.

Synergy between axon guidance cues was first observed nearly two decades ago. Work in *Xenopus* spinal axons demonstrated netrin-1 attraction is mitigated by the binding of the repulsive cue Slit to its receptor Robo. Hindering response to netrin-1 is critical for midline crossing in the brain¹⁹. Since this discovery, a multitude of guidance cue combinations have been observed to coordinate axon pathfinding²⁰. In recent years, a synergistic effect between netrin-1 and the repulsive cue Ephrin-A5 was observed in both mouse and chick spinal lateral motor column (LMC) axons, yet the mechanism of this cooperation was unknown²¹. Subsequent work demonstrated explant treatment with Ephrin-A5 led to increased surface expression of the netrin receptor neogenin, therefore enhancing netrin-1 attractive axon turning²². Future work will question if this convergence between netrin and ephrin signaling impacts other stages of development, including synaptogenesis.

With the expanding repertoire of guidance cues, receptors, cell adhesion molecules and intersecting downstream signaling pathways, there are ample opportunities to investigate potential synergy and discord between diverse molecular pathways, as well as potential context dependence (i.e. cell type, receptor expression, or extracellular environment), that combine to direct the axon. For example, cell adhesion molecules can further sculpt pathfinding. A recent example in cultured murine midbrain dopamine (mDA) neurons demonstrated that the cell adhesion molecule CD166 (ALCAM) interacts *in trans* with the semaphorin receptor complex (L1cam, Chl1 and Nrp1). Interestingly, this interaction blocked Sema3A-induced axon growth and enhanced Sema3C-induced axon branching, likely by altering receptor availabilities²³ (Fig. 1A). Additional work is required to understand the convergence of numerous signaling events downstream of extracellular cues. For example, Shh-mediated guidance of commissural axons toward the floorplate requires activation of unconventional guanine exchange factors (GEFs) Dock3 and 4, their target Rac, and subsequent cytoskeletal remodeling, as well as Shh stimulated endocytosis of its receptor Boc and subsequently Ptch1^{24,25}. All these inputs eventually converge on production of mechanical forces that drive motility.

The growth cone transduces mechanical force

The classic “clutch hypothesis” of motility was first proposed three decades ago^{26,27}. In this model, neurons are linked to the extracellular matrix (ECM) by adhesion molecules. In turn, adhesion molecules are connected to the acto-myosin network by “clutch proteins”. Contraction of the actin cytoskeleton by myosin motors generates force, which is then transmitted to the ECM, facilitates motility (Fig. 1B). In non-neuronal cells, talin and FAK are understood as the clutch between integrins and the actin cytoskeleton. However, it is well appreciated that neuronal clutch molecules can differ between adhesion types, cell types, and extracellular substrates. For example, the cell adhesion molecule (CAM) L1-CAM is critical for adhesion during laminin-based haptotaxis in rat hippocampal neurons. Interestingly, L1-CAM mobility in the growth cone inversely correlates with traction force generated by the growth cone, indicating tunable adhesion and force generation. Human L1-CAM mutations associated with neurological syndromes and projection defects reduce L1-CAM adhesion to laminin and growth cone mechanotransduction²⁸, highlighting physiological importance of force generation. Shootin1a and cortactin serve as clutch molecules linking F-actin retrograde flow in the growth cone to L1-CAM^{29,30} to promote axon outgrowth. Notably, netrin-1 treatment of hippocampal neurons enhances L1-CAM and shootin1a interactions, facilitating growth cone motility and highlighting the irrevocable link between guidance cue signaling and mechanotransduction³⁰.

Transmembrane adhesion receptors are not alone in mechanotransduction of growth cones; stimulation of mechanosensitive ion channels, such as Piezo or TRP channels, feeds back into signaling pathways to promote outgrowth³¹⁻³³. In particular, membrane stretch triggers TRPV2 activation, increasing calcium signaling and enhancing axon outgrowth in cultured murine DRGs³⁴. To understand how TRPV2 is activated, subsequent studies in PC12 cells demonstrated TRPV2 localizes to the growth cone and clusters in response to mechanically applied force. TRPV2 also interacts with actin, facilitating cytoskeletal remodeling, increased filopodial length, and ultimately axonal outgrowth³⁵ (Fig. 1B). Critical questions

remain unanswered in neuronal mechanotransduction. Although new research has identified clutch molecules between various adhesion molecules and the cytoskeleton, the conservation of these interactions between cell types or with other receptors is unknown. Furthermore, while soluble netrin-1 treatment enhanced L1-CAM-shoottin interactions, it is unknown if haptotactic netrin-1 acts through this same pathway. As netrin-1 has numerous known receptors, this research also opens the door to discovery of new mechanotransduction pathways.

Cytoskeleton reorganization drives growth cone motility

Growth cone morphology, and in turn motility, are controlled by reorganization of the actin and microtubule cytoskeletons. Although neuronal studies have focused on Act β , six mammalian genes encode actin³⁶. Act α , Act β , and Act γ all localize to growth cone filopodia in cultured murine motoneurons. Knockdown of any isoform decreases filopodia initiation. Loss of Act α or Act γ decreases filopodial dynamics whereas Act β knockdown decreases growth cone size and motility³⁷. Actin-based myosin motors play numerous roles in growth cones, including traction force generation (myosin II) and vesicle/cargo transport (myosin V, VI and X). New work in cultured murine cortical neurons shows Myo1b localizes to the growth cone membrane and regulates actin dynamics, filopodia stability, and growth cone morphology, potentially via propagation of actin waves proposed to supply cytoplasmic material for growth cone motility³⁸. Additional work investigating the interplay of different actin genes and myosin motors in growth cone motility is warranted.

A critical relationship exists between microtubule stability—and consequently microtubule reorganization—and growth cone motility^{39,40}. This was highlighted in studies of both regulator of presynaptic morphology 1 (RPM-1), an E3 ubiquitin ligase, and the scaffolding protein WDR47. RPM-1 localizes to the growth cone of *C. elegans* PLM mechanosensory neurons and promotes collapse, potentially by opposing the function of microtubule stabilizers⁴¹. Similarly, WDR47 interacts with the microtubule destabilizing protein, superior cervical ganglion-10 (SCG10). *Wdr47*^{-/-} cortical and hippocampal neurons exhibit decreased growth cone area, which can be partially rescued by Epothilone D, a pharmacological microtubule stabilizer⁴². *Wdr47*^{-/-} mice exhibit microcephaly and corpus callosum agenesis, highlighting the critical function of microtubule stability in pathfinding.

Emerging motility research highlights links between microtubules and actin via *Xenopus* microtubule associated protein 215 (XMAP215) and Dishevelled-associated activator of morphogenesis (DAAM) (Fig. 1C). The microtubule polymerase XMAP215 regulates growth cone morphology and facilitates microtubule invasion into filopodia. This involves an interaction of XMAP215 with both actin and microtubules and likely contributes to growth cone remodeling during guidance⁴³. Likewise, the actin-nucleating formin DAAM also binds both microtubules and tip localized EB1. In *Drosophila* primary neurons, DAAM binds pioneer microtubules and nucleates new actin filaments to promote filopodial stability⁴⁴.

The rapid remodeling of the growth cone cytoskeleton in response to guidance cues highlights the demand for precise cytoskeletal regulation. Post-translational modification of

cytoskeletal elements poises the cytoskeleton for rapid regulation, and examples continue to emerge. For example, cortactin, an activator of the actin branch nucleating Arp2/3 complex, is a Src kinase substrate. Phosphorylation of cortactin promotes the formation of new growth cone filopodia and increased filopodial length⁴⁵. As Src family kinases act downstream of guidance cues, cortactin phosphorylation may initiate rapid filopodial changes in response to extracellular stimuli (Fig. 1D).

Ubiquitination also confers rapid signaling and changes in protein function. Classically, ubiquitination is considered a modification marking proteins for proteasomal degradation. Proper protein turnover is essential for neuronal function, and numerous neurodevelopmental and neurodegenerative diseases are associated with defective protein degradation⁴⁶. New work demonstrates the Siah1-mediated ubiquitination and degradation of protein kinase Akt3 (also known as PKB γ) is required for appropriate axon extension and branching. Furthermore, an Akt3 mutation that reduces Siah1 binding and ubiquitination is linked to focal malformations of cortical development⁴⁷. In addition to its degradative purposes, ubiquitination—particularly mono-ubiquitination—has emerged as a mechanism to regulate protein localization and activity. For example, reversible, non-degradative ubiquitination of the actin polymerase VASP regulates filopodial stability and axon guidance in response to netrin⁴⁸. New work also suggests competition between two class I TRIM E3 ubiquitin ligases, TRIM9 and TRIM67, that controls VASP ubiquitination⁴⁹ (Fig. 1D). Interestingly previous work demonstrating Madd-2, the single *C. elegans* class I TRIM ortholog, interacts with UNC-40 (DCC homolog) and is required for proper axon guidance^{50,51}. How duplication of TRIM genes adds additional complexity to signaling pathways integral to axon guidance will be an interesting avenue of future research. As TRIM9 and TRIM67 both localize to the growth cone, the identification of new ubiquitination targets—both regulatory and degradative—will further illuminate complex signaling pathways. Furthermore, with over 600 E3 ligases encoded in the mammalian genome, the role of ubiquitination remains a rich and understudied area of neuronal biology.

In neurons, the dynamic cytoskeleton also includes septins and neurofilaments. SEPT6 and SEPT7 function in filopodia formation and axon branching⁵², yet the role of septin family is unknown in the growth cone. Likewise, neurofilaments are present in the growth cone and filopodia⁵³, yet their precise role is understudied. For example, the intermediate filament protein nestin was previously thought only to be expressed in neuronal progenitor cells. Surprisingly, new work demonstrates maintenance of nestin expression in immature cortical neurons sensitizes neurons to Sema3A-induced collapse and retraction⁵⁴. The inclusion of both septins and neurofilaments in future growth cone studies is a necessary measure to fully understand axon guidance.

Calcium regulates growth cone signaling

Calcium signaling is essential for regulating the actin and microtubule cytoskeletons, and ultimately axon guidance^{55,56} (Fig. 1E). For example, the actin severing protein cofilin is activated downstream of the calcium dependent phosphatase calcineurin in response to serotonin⁵⁷. In serotonin-treated *Aplysia* bag neurons, cofilin activation is enhanced downstream of PKC and myosin II, which modulates actin density, traction stress in the

growth cone, and neurite outgrowth. This suggests myosin II regulates cofilin-mediated actin depolymerization downstream of calcium⁵⁸. The microtubule cytoskeleton is regulated via Stromal interacting molecule 1 (STIM1), which localizes to the ER and regulates calcium levels⁵⁹. STIM1 colocalizes with End binding protein 3 (EB3) and promotes microtubule invasion into filopodia and growth cone turning in cultured neurons, and proper axon guidance in zebrafish caudal primary (CaP) neurons⁶⁰.

Calpain is a calcium-dependent protease implicated in adhesion and motility. In immortalized cell lines, cleavage of numerous focal adhesion related proteins—including talin⁶¹ and FAK⁶²—by calpain is well appreciated to promote adhesion turnover during cell migration. New work shows talin and FAK are also targets of calpain in *Xenopus* spinal cord neurons, and their cleavage is required for repulsive growth cone turning *in vitro* and appropriate extension of Rohon–Beard peripheral axons *in vivo*⁶³. Furthermore, netrin-1 promotes calpain activity and cleavage of the netrin receptor DCC in rodent cortical neurons⁶⁴. This suggests DCC cleavage may present a negative feedback mechanism to regulate growth cone motility. Furthermore, if DCC is binding substrate adherent netrin as opposed to diffusible netrin, DCC proteolysis may highlight another role for calpain in adhesion. Whether cleavage of other guidance receptors is linked to calpain activation⁶⁵ is an open question.

Membrane trafficking delivers plasma membrane material during axon outgrowth

Unlike cytoskeleton remodeling or calcium signaling, membrane trafficking plays an underappreciated role in axon guidance. As highly polarized cells, neurons face the unique challenge of directionally targeting cargo to the growing axon. In particular, growth cone motility during neurite outgrowth is concomitant with increased plasma membrane surface area⁶⁶. Classic hypotheses propose insertion and remodeling of membrane material occurs via calcium regulated exocytosis and membrane flow^{67,68}. Recent empirical measurements of VAMP2 and VAMP7-mediated exocytosis in cultured neurons along with computational simulations of exocytosis, endocytosis, and membrane expansion suggest that VAMP2-mediated exocytosis supplies sufficient membrane material for neurite outgrowth early in murine cortical neuron development. Surprisingly, few exocytic events were observed in the growth cone, suggesting that membrane flow toward the tips of axons may play a role in axon outgrowth⁶⁹, yet this remains to be tested. However, this hypothesis contradicts earlier work that proposes asymmetric exocytosis as a mechanism for growth cone turning in chick dorsal root ganglion neurons⁷⁰. To understand these disparities, further work on sites of membrane addition and flow, as well as exocytic vesicle trafficking in the growth cone, are needed.

In addition to exocytic vesicles, other membrane bound compartments contribute to membrane addition and protein trafficking in the growth cone⁷¹. For example, lysosomes promote degradation of plasma membrane and extracellular substrates, exocytosis, and plasma membrane repair. Recent work indicates lysosomes are directionally transported to the axonal growth cone via kinesin-1 and a large ensemble of proteins including the

BLOC-1 related complex⁷². This biased motility maintains growth cone size and outgrowth in rat hippocampal neurons. Mitochondria also localize to the growth cone and facilitate axon outgrowth, presumably by generating ATP for actin polymerization⁷³. Mitochondria dynamics within the zebrafish growth cones *in vivo* suggested that a subset of mitochondria dock in the central zone to microtubules in a syntaphilin-dependent manner and a subset localize to growth cone filopodia, although the mechanism of transport is unclear⁷⁴. Future work is still required to fully connect membrane trafficking to guidance cue and mechanotransduction signaling to create a holistic understanding of growth cone motility.

Parallels between growth cones and dendritic spines

As cytoskeletal rearrangements dominate both growth cone motility and synaptogenesis, we asked whether pathways in the growth cone are conserved in dendritic spines (Fig 2). During synapse formation, filopodia grow and retract from dendrites, with the potential to form contacts with axons, and mature into actin-rich dendritic spines that receive synaptic transmission⁷⁵. Like in the growth cone, dendritic filopodia are dynamic structures that must respond to signals in the local environment. However, they also contain the proper machinery to develop into long-lived dendritic spines if they find the correct pre-synaptic partner. These distinct outcomes for dendritic filopodia may exploit parallel combinatorial usage of pathways for diverse outcomes as highlighted in distinct growth responses.

Numerous guidance cues and receptors discussed above are also linked to synaptogenesis and synaptic plasticity. For example, Netrin and DCC are enriched in the synapse and netrin increases cortical dendritic filopodia number, synapse number, and synapse strength⁷⁶. Netrin-1 facilitates synaptic plasticity by increasing surface localization of Glutamate receptor-1⁷⁷. DCC and downstream Src signaling are also required for long-term potentiation and spatial memory^{78,79}. Likewise, recent reviews highlight the role of semaphorins⁸⁰, wnt⁸¹, and ephrins⁸² in synaptogenesis. Furthermore, adhesion molecules implicated in axon guidance also localize to synapses and modulate cytoskeletal dynamics. For example, downstream of integrins, CaMKII is critical for actin reorganization during synaptic plasticity⁸³. Cadherins are also required for long-term potentiation in both the hippocampus⁸⁴ and ventral tagmental area⁸⁵. Notable recent work focuses on the binding of filopodial EphB (a receptor tyrosine kinase) to axonal ephrinB during synaptogenesis. Interestingly, the kinetics of EphB activation determined filopodial fate—whereas fast activation led to filopodial retraction, slower activation facilitated persistent binding⁸⁶. Finally, mechanotransduction, generated by linking these proteins to the cytoskeleton, is hypothesized to play a role in dendritic spine formation and plasticity, and is an avenue of future research⁸⁷.

The dendritic spine actin cytoskeleton has been equated to the branched actin network of lamellipodia^{88,89}, with formin and VASP-mediated protrusions contributing to motility⁹⁰. As a multitude of proteins can regulate actin polymerization—enhancing nucleation, elongation, bundling, severing, etc—it is notable that similar proteins play similar roles in both structures. This suggests mechanisms observed in growth cones may be conserved in spines, with the potential to inform future hypotheses regarding spine maturation and plasticity. For example, as our lab has found that TRIM9-mediated ubiquitination of VASP regulates

growth cone filopodia stability, we hypothesize that VASP ubiquitination may also impact dendritic spine morphology or function. Although key differences between these two structures are obvious—namely size and longevity—both structures must be poised to react rapidly remodel their cytoskeleton in response to extracellular cues, albeit the outcome leading to axon turning or plasticity.

Similar to the growth cone, additional research is required to understand the transient dendritic filopodia that preface spine development. However, interesting differences between growth cone and dendritic filopodia have been noted; unlike the growth cone, Arp2/3 and capping protein are enriched in dendritic filopodia that lack the actin-bundler fascin⁹¹. These disparities hint at the combinatorial power of actin regulators—the slight tweaking of the molecular makeup results in structures with distinct architecture or dynamics.

Conclusions

Although significant, unanswered questions remain regarding growth cone signaling and motility, a multitude of knowledge has accumulated over the past century. A theme of increasing complexity, specificity, and synergy emerges, likely lending exquisite tunability to growth cone navigation. As research continues, we expect additional convergence and specificity of pathways and cytoskeletal regulation to be revealed. Additional parallels and differences between the development of spines and growth cones will emerge, accounting for the dynamic tuning yet unique maturation and morphological changes of these neuronal structures.

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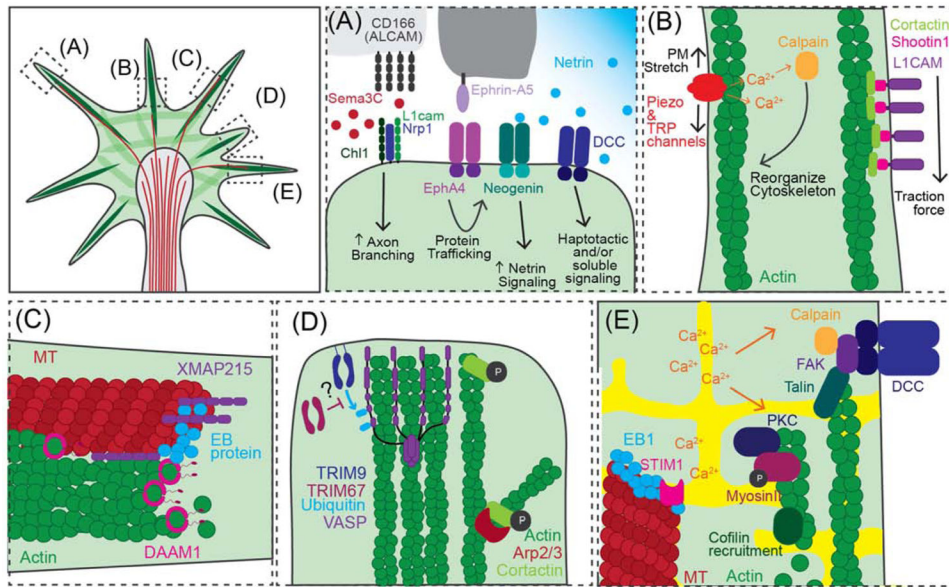


Figure 1. Recent research in growth cone motility.

Although guidance cues, mechanotransduction, cytoskeleton reorganization, and calcium signaling holistically contribute to growth cone motility, we've selected several new discoveries in each of these categories to highlight here.

(A) New work highlights contribution of adhesion molecules (CD166 enhances Sema3C-mediated axon branching), combinatorial effects (Ephrin-A5 signaling increases Neogenin surface expression and downstream Netrin-1 signaling) and both haptotactic and soluble signaling (Netrin-1) in axon guidance. (B) Mechanotransduction influences motility, both by the generation of traction force on the ECM (L1CAM) and the activation of Piezo and TRP channels by stretching of the plasma membrane. (C) Coupling of the microtubule and actin cytoskeletons in the growth, newly shown to be mediated by XMAP215 and DAAM1. (D) Post-translational modifications facilitate rapid changes in filopodial formation and stability, as indicated by phosphorylation of cortactin and VASP ubiquitination. (E) Calcium signaling is required for growth cone remodeling. Recent work includes: microtubule organization is regulated by STIM1, calpain cleavage of FAK, talin and DCC, and PKC-mediated Myosin-II activation promotes cofilin recruitment and actin severing.

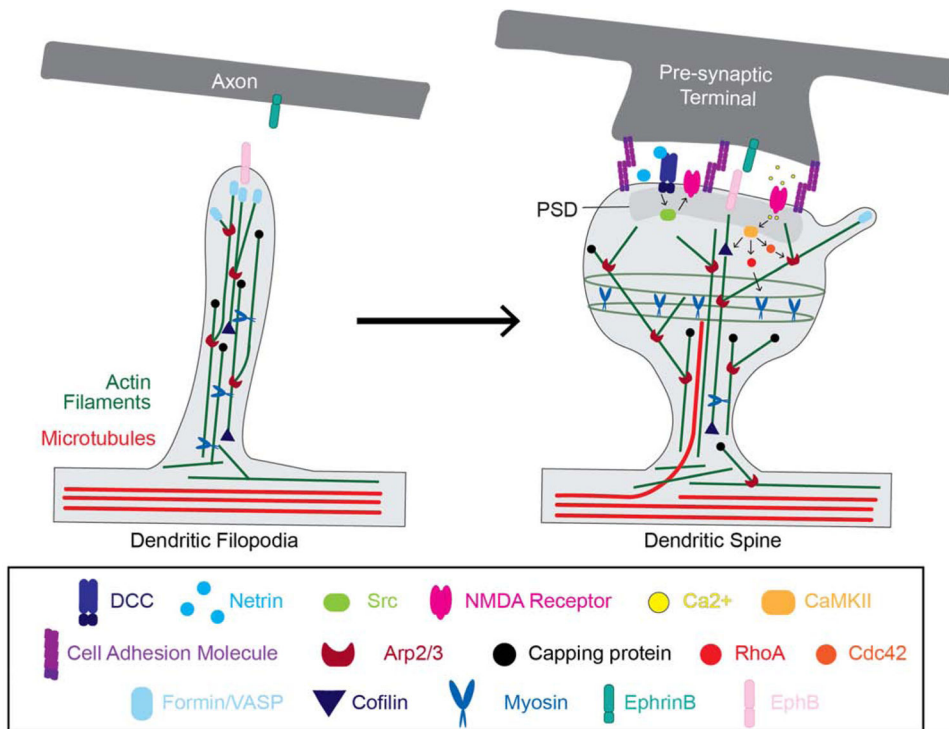


Figure 2. Dendritic filopodia and dendritic spine.

(A) Dendritic filopodia. Like in the growth cone, formin proteins have been observed at the tips of filopodia, elongating actin filaments. However, unlike in the growth cone, dendritic filopodia are known to contain the branching Arp2/3 complex and capping protein, while lacking the actin-bundling protein fascin. Sustained Eph-B signaling (following axonal EphrinB binding) facilitates synaptogenesis.

(B) Dendritic spine. Numerous and diverse cell adhesion molecules facilitate synaptic maturation and plasticity, by connecting the pre- and post-synapse (along with the ECM) and contributing to signaling pathway activation. Netrin signaling via the DCC receptor leads to downstream Src signaling required for long-term potentiation. NMDAR signaling results in the calcium-dependent activation of CaMKII, activating numerous cytoskeleton proteins including cofilin (leading to actin severing and depolymerization), RhoA (leading to myosin activation) and Cdc42 (ultimately leading to Arp2/3 mediated actin branching). The actin cytoskeleton of the dendritic spine is predominantly branched, akin to the lamellipodia, with formin and VASP-mediated protrusions.