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The nano-architecture of the axonal cytoskeleton

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Abstract | The corporeal beauty of the neuronal cytoskeleton has captured the imagination of generations of scientists. One of the easiest cellular structures to visualize by light microscopy, its existence has been known for well over 100 years, yet we have only recently begun to fully appreciate its intricacy and diversity. Recent studies combining new probes with super-resolution microscopy and live imaging have revealed surprising details about the axonal cytoskeleton and, in particular, have discovered previously unknown actin-based structures. Along with traditional electron microscopy, these newer techniques offer a nanoscale view of the axonal cytoskeleton, which is important for our understanding of neuronal form and function, and lay the foundation for future studies. In this Review, we summarize existing concepts in the field and highlight contemporary discoveries that have fundamentally altered our perception of the axonal cytoskeleton.

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The cytoskeleton is a cellular shapeshifter. Like these creatures of mythology and speculative fiction, the cytoskeleton can alter its physical form and shape to accommodate the immediate needs of the cell. Indeed, many a rapt student has watched the cytoskeleton of a dividing cell seemingly miraculously transform into an organized mitotic spindle and then dissolve into an indiscrete mass, right before their eyes. The extreme polarization of neurons (that is, their fundamental asymmetry, arising due to the presence of elongated processes), along with their lifelong plasticity, creates unique demands on the cytoskeleton. A remarkable example is the axon, which can grow to enormous lengths and must generate and maintain its form and function throughout life — a burden that rests largely on the cytoskeleton.

The axonal cytoskeleton has three major constituents: microtubules, neurofilaments and actin (BOX 1). Each is unique, associating with its own set of binding proteins and performing specialized roles within the axon. Most of these cytoskeletal proteins are synthesized in the neuronal soma and are transported along the axon. Such transport is a constitutive phenomenon that occurs throughout the life of the neuron. Thus, the axonal cytoskeleton is best understood by considering both its anatomical organization and its dynamics (including axonal transport).

Given the complex morphology and physiology of neurons, the field of neurobiology has traditionally been at the forefront of adopting new optical technologies as they arise. Advances in microscopy now allow us to observe cells with unprecedented spatial resolution and to follow dynamic processes with exquisite temporal detail[1]. For example, super-resolution strategies that circumvent the diffraction limit of optical microscopy appeared more than ten years ago (BOX 2) and have been used to reveal aspects of neuronal organization down to the scale of macromolecular complexes[2]. These techniques have provided key insights into the organization and function of the axonal cytoskeleton — and in particular the organization of actin and microtubules — revealing how it builds the axon and maintains its intricate architecture. Focusing on the cytoskeleton within the axon initial segment (AIS) and the more distal axon shaft, in this Review, we highlight these recent discoveries and place them in the context of earlier findings, giving the reader a tentative vision of the future.

Overview of the axonal cytoskeleton

The history of cytoskeletal research is essentially the pursuit of tools and techniques that allowed an ever-closer view of this elaborate structure, a quest that continues to this day. Early studies by 17th-century microscopy pioneers highlighted a network of 'neurofibrils', which we now know were most likely neurofilaments[3]. With the advent of electron microscopy, two types of fibrils were seen in axons: filaments measuring approximately 10 nm in diameter, corresponding to neurofilaments, and others measuring approximately 20–30 nm in diameter, corresponding to structures that eventually came to be known as microtubules[4,5].

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Box 1 | **The components of the cytoskeleton**

Microtubules are present in all eukaryotic cells and play essential roles in cell division, the movement of cellular organelles and structural support. They are composed of 13 protofilaments, which are organized into a cylinder of 25 nm diameter. Each protofilament is made of heterodimers composed of α-tubulin and β-tubulin, which are arranged in an alternating pattern. These heterodimers bind GTP, which is progressively hydrolysed to GDP after assembly. Microtubules are polar structures, with a fast-growing 'plus' end and a slower-growing 'minus' end.

Actin filaments (also known as F-actin) are found in all eukaryotes and play critical roles in cell motility, division, adhesion, signalling and many other fundamental processes. Actin filaments are approximately 8 nm in diameter, are composed of heterodimers of α-actin and β-actin (known as G-actin) and require ATP for polymerization. Actin filaments have a defined polarity, with a barbed (rapidly growing) end at which monomer exchange is greater than it is at the other (pointed) end.

Neurofilaments are the intermediate filaments of neurons, measuring approximately 10 nm in diameter. They provide mechanical support to the axon and regulate axon diameter. Neurofilaments are heteropolymers, composed of subunits of high, medium and low molecular weight as well as internexin or peripherin intermediate filaments. Each neurofilament subunit has a short amino-terminal globular head domain, a central conserved alpha-helical rod domain and a variable carboxy-terminal tail domain. The central rod domain co-assembles with other subunits to form dimers, protofilaments and filaments. Unlike actin and microtubules, neurofilaments are not polarized and do not require ATP or GTP for polymerization.

Electron microscopy
A group of methods that generate an image of a sample by using a beam of electrons. Electrons can be detected after passing through the sample (transmission electron microscopy) or after being reflected (scanning electron microscopy). Electron microscopy can routinely reach 1 nm resolution (the size of a single amino acid) but usually requires the sample to be placed in a vacuum and is therefore destructive and most applicable to fixed samples labelled with specific procedures.

Immuno-electron microscopy
An electron microscopy modality in which proteins labelled by using antibodies that are tagged with small gold beads are imaged, allowing their localization.

Microtubule plus-end-tracking proteins
A set of proteins that bind the growing plus ends of microtubules. The core components of this complex are end-binding proteins, dimeric proteins that interact with the specific tubulin conformation found at the plus end.

As electron microscopy techniques improved, they revealed a surprisingly complicated cytoskeletal network in axons, composed of intermingled neurofilaments and microtubules and numerous 'side arms' and 'cross bridges' (REFS 6,7). At the same time, biochemical studies indicated that these cytoskeletal polymers were composed of building blocks, or subunits, providing a fundamental insight into their biogenesis[8,9]. Although actin filaments were long known to be present in muscle, these linear polymers were eventually also identified in various non-muscle cells, including neurons[10].

The evolution of electron microscopy also led to the identification of a new anatomical and physiological compartment along the first 20–60 μm of the axon shaft: the AIS. The AIS is of exceptional interest because it is the region within which axon potentials are initiated (reviewed in REF. 11). The AIS is characterized by a unique cytoskeletal organization that has two striking features: small fascicles of microtubules and a dense sub-axolemmal undercoat[12–14]. Unlike the more distal axon, where microtubules are uniformly dispersed within the cross section of the axoplasm, microtubules in the AIS are arranged in fascicles in which 3–6 closely apposed microtubules are connected by electron-dense cross bridges[13] (FIG. 1). This unique clustering was suggested to have a role in directing the polarized trafficking of vesicles and organelles into the axon[15,16]. The undercoat[17] was shown to contain a dense complex of cytoskeletal proteins (including actin, spectrin and ankyrin-G, also known as ankyrin 3) that tethers an array of membrane proteins, including voltage-gated sodium and potassium channels and cell adhesion molecules (CAMs)[18–22]. The membrane proteins anchored at the AIS were later shown to create a diffusion barrier between the cell body and the axon[23,24].

Although the picture painted by electron microscopy suggested a static structure, other studies hinted that the axonal cytoskeleton was dynamic. One clue came from pulse-chase radiolabelling studies that examined axonal transport of the cytoskeletal components. In these experiments, newly synthesized proteins were labelled in the neuronal soma, and their egress into the axon, reflecting axonal transport, was inferred by examining the amount of label present in successive axon segments over time. These studies indicated that the three major cytoskeletal proteins are conveyed with overall velocities that are several orders of magnitude lower than those of vesicles and mitochondria. This sluggish rate class was called slow axonal transport[25–29]. Neurofilament proteins and a large fraction of tubulin proteins (microtubule subunits) are conveyed in the slowest form of this transport class, known as 'slow component a' (SCa), moving at rates of approximately 0.2–2 mm per day. In most peripheral nervous system (PNS) neurons, actin is conveyed a little faster, at rates of approximately 2–4 mm per day, known as 'slow component b' (SCb), though reported rates in CNS neurons are only approximately 0.4 mm per day[26,29]. Although these studies could detect overall axonal transport, the methods used could not visualize the transported cargo, and mechanistic details remained obscure. With the advent of fluorescent probes including green fluorescent protein (GFP), cytoskeletal structures in a variety of other cell types were visualized by live imaging, revealing a highly dynamic organization[30]. More recent studies, described below, have expanded this concept and ushered in a new era in which super-resolution microscopy and live imaging have played a crucial role.

Although electron microscopy gave an unprecedented view of the cytoskeleton and offered the first clues into its organization, there are two limitations. First, some of the fixatives used in electron microscopy, although invaluable for the visualization of the cytoskeleton, can be harsh, leading to disruption of delicate structures[31–34] (reviewed in REF. 35). Second, proteins associated with the cytoskeleton cannot be identified easily by this technique. Although immuno-electron microscopy provides a potential solution to the latter problem, the methodology is challenging and idiosyncratic. Fluorescence microscopy, when combined with cytoskeleton-binding probes, offers a number of ways to highlight the cytoskeleton (TABLE 1). However, the axonal cytoskeleton is tightly packed, and it is not easy to discern individual structures owing to the diffraction limit of optical microscopy (approximately 200 nm). This issue has now been somewhat resolved by the emergence of optical super-resolution imaging techniques (BOX 2), although these are also subject to fixation artefacts[2]. As described below, these studies have not only clarified the spatial relationships of the various cytoskeletal elements in axons — in some cases revealing new biology — but have also uncovered previously unknown assemblies that could not be appreciated by conventional light microscopy. In addition, our ability to visualize the cytoskeleton by live imaging has been transformative, leading to the demonstration of the transport of neurofilaments as assembled polymers[36,37] and the motility of microtubule plus-end-tracking proteins[38,39], for instance.

Moiré effect
The emergence of a third pattern due to the superposition of two patterns with distinct frequencies. In microscopy, this effect is exploited in structured illumination microscopy by illuminating the sample with periodic patterns of light and using the resulting Moiré pattern to infer sample details that are beyond the diffraction limit.
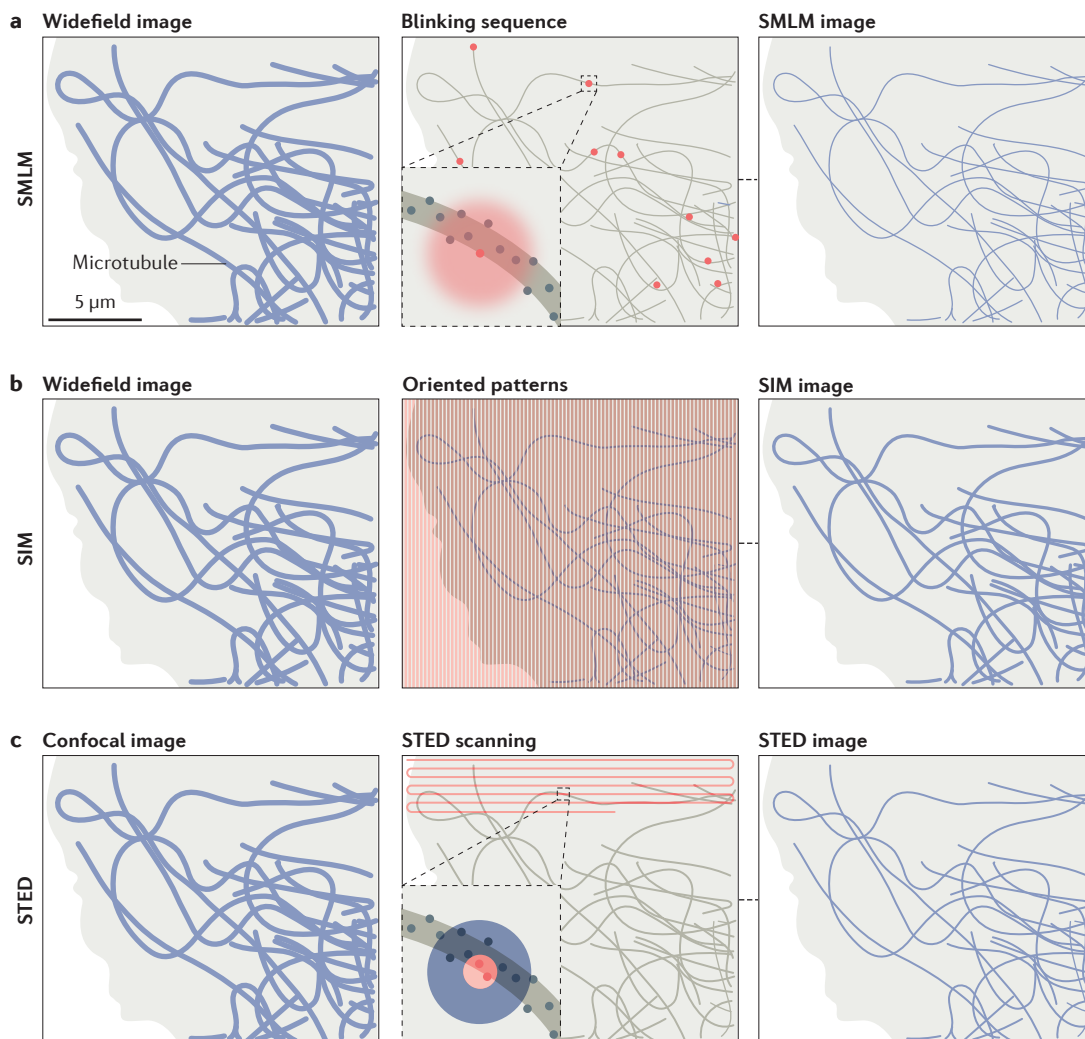
Box 2 | **Super-resolution optical microscopy techniques**

Several light microscopy techniques allow us to visualize the organization of the cytoskeleton at a resolution that is currently down to a few tens of nanometres. These techniques have also been extended to enable 3D imaging and continue to develop rapidly[177–179].

When a light microscope is used to acquire a traditional widefield image, all the fluorophores present in a cellular structure emit light at once. In single-molecule-localization microscopy (SMLM, see the figure, part **a**), only a few fluorophores emit light at any time. In the SMLM modality known as stochastic optical reconstruction microscopy (STORM)[180], a reducing buffer ensures that most of the organic fluorophores present in an immunolabelled cell enter a long-lasting dark state upon excitation. At any given time, only a few fluorescent molecules are stochastically switched to an active state. In photoactivated localization microscopy (PALM)[181], fluorophores are photoactivable or photoconvertible fluorescent proteins that are sparsely activated or converted by a low-power violet light, and then imaged and bleached. In both STORM and PALM, light emission from single molecules results in the generation of 'spots' of a size that is determined by diffraction (~200 nm, large red spot). The centre of each isolated spot can be fitted by a 2D Gaussian function, resulting in localization of the spot with ~20 nm precision (small red dot in inset). The single-molecule coordinates for all fluorophores in the imaged sample are accumulated over a blinking sequence (tens of thousands of frames) to generate an image (the resolution of which is typically 20–30 nm with current approaches).

Structured illumination microscopy (SIM; see the figure, part **b**) is based on the illumination of the sample with periodic light patterns (indicated in the schematic by an overlaid grating) that lead to a Moiré effect[182]. By acquiring the light emissions that result from 9–15 complementary patterns of excitation, an image can be reconstructed that provides a 2X gain in resolution (a resolution of ~100 nm). Nonlinear variants of SIM can go down to a resolution of ~50 nm (REF. 183).

Stimulated emission depletion microscopy (STED; see the figure, part **c**) is a scanning technique based on confocal imaging (left panel). In STED (middle panel), a focused laser scans the sample: this is composed of an excitation beam (inset, blue and pink) and a high-power, donut-shaped depletion beam (inset, blue) that restricts the emission from excited fluorophores to a small, tuneable area (inset, pink)[184]. A single-point detector collects light from this sub-diffractive area, giving a resolution of ~40 nm.
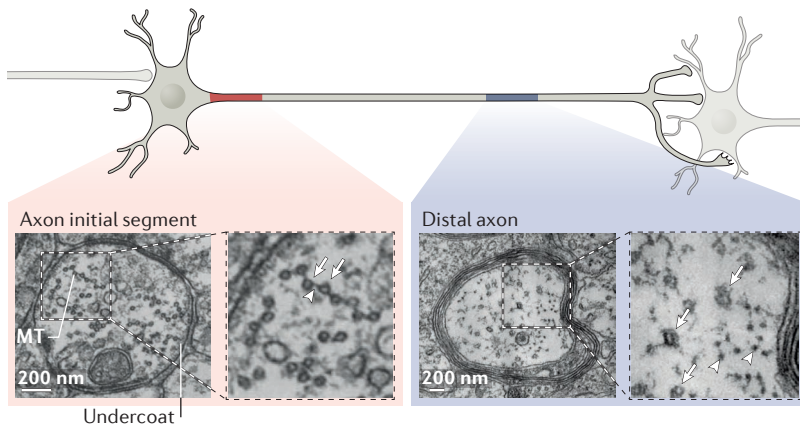
Figure 1 | **Ultrastructure of the axon shaft.** Classical electron microscopy studies were the first to allow us to visualize the organization of the cytoskeleton within axonal shafts. The schematic depicts the different compartments of a mature neuron, and the electron microscope images illustrate the cytoskeletal ultrastructure in different regions of the axon. The left panels show a thin-section electron micrograph of the axon initial segment from a pyramidal neuron of the rat cerebral cortex and a magnified view of part of this image. On this transverse section, the microtubule fascicles (MT) contain individual microtubules. These appear as beads (arrows in inset) that are connected by cross bridges (arrowhead in inset). The dense undercoat is visible along the plasma membrane. The right panels show a thin-section electron micrograph of a rat optic nerve, specifically the transverse section of a myelinated axon, and a magnified view of part of this image. The microtubules appear as hollow rings (arrows in inset) and neurofilaments as groups of smaller puncta (arrowheads in inset). Left panels adapted with permission from ©Peters *et al. J. Cell Biol.* **39**, 604–619 http://dx.doi.org/10.1083/jcb.39.3.604 (1968) (REF. 14). Right panels adapted with permission from ©Peters and Vaughn. *J. Cell Biol.* **32**, 113–119 http://dx.doi.org/10.1083/jcb.32.1.113 (1967) (REF. 47).

Microtubules in the axon shaft

Organization of axonal microtubules. As first revealed by electron microscopy, microtubules within the axon are uniformly oriented with their plus ends directed towards the axon tip, whereas dendrites contain a mixture of plus-end-out and minus-end-out microtubules[40–43]. More recently, axonal microtubule polarity was confirmed in cultured neurons and *in vivo* by tracking growing microtubules with fluorescent microtubule end-binding proteins (EBs) such as EB1 (also known as microtubule-associated protein RP/EB family member 1 (MAPRE1)) and EB3 (also known as MAPRE3)[38,39,44,45]. Notably, this revealed that microtubule polarity changes progressively over development from 80% plus-end-out in undifferentiated neurites (BOX 3) to 100% plus-end-out after axon specification[39].

In vertebrate axons, microtubules are densely packed, with an average of ten microtubules per square micrometre in cross sections of large-diameter axons and up to 100 per square micrometre in smaller ones[46,47]. Owing to their density and bundling, determining the lengths of axonal microtubules has been difficult. Using serial-section electron microscopy, the reported lengths of microtubules in vertebrates were 4 µm on average in cultured rat sympathetic neurons[48] but much longer when measured *in vivo* (including 100 µm in rat sensory axons[49], 400 µm in frog olfactory axons[50,51] and up to 750 µm along the mouse saphenous nerve[52]). The sparsity of axonal microtubules in *Caenorhabditis elegans* DA9 motor neurons recently allowed an optical method to be used to determine their microtubule organization and length. Using transgenic worms engineered to express tubulin–GFP, a recent study visualized 'steps' in fluorescence profiles along axons that correspond to the microtubule extremities[53]. This revealed that the average microtubule length varied between 3.5 µm in L2-stage larvae and 7 µm in adult worms[53], in line with previous estimates from electron microscopy (10–20 µm in touch receptor neurons[54]). These lengths are significantly shorter than those of mammalian neurons, where the higher density and longer length of microtubules preclude the use of this optical approach. Refinements in super-resolution techniques could help in this regard: for instance, single-particle tracking of retrogradely transported cargoes — by using quantum dots conjugated with nerve growth factor molecules — results in nanoscale trajectories that make it possible to discriminate between individual microtubules within axonal bundles[55]. New, smaller probes also allow the labelling and super-resolution imaging of individual microtubules, which are normally densely packed in neuronal processes[56].

As described above, microtubules have a unique organization at the AIS, forming tight bundles connected by electron-dense crosslinkers[13] (FIGS 1,2a). Although the identity of the crosslinkers is still unknown, a recently identified protein, tripartite motif-containing protein 46 (TRIM46), may be a candidate. Identified as the target of autoantibodies derived from patients with paraneoplastic neurological syndrome, TRIM46 is enriched along microtubules in the proximal AIS[57]. By use of a combination of super-resolution and live-cell imaging, TRIM46 was shown to drive the establishment of uniformly oriented microtubules at the axon entrance and also to generate fascicles of microtubules when overexpressed in HeLa cells[58]. The specific organization of AIS microtubules has long been thought to have a role in regulating cargo entry into the axon[59], and, interestingly, polarized trafficking is disrupted in TRIM46-deficient neurons[58]. Another microtubule-associated protein, microtubule crosslinking factor 1 (MTCL1), was recently shown to be enriched at the AIS of cerebellar Purkinje cells and to help in AIS maturation and maintenance[60]. Finally, microtubule-associated EBs provide a molecular link between the submembrane actin–spectrin complex (see below) and the deeper AIS microtubule fascicles[11]. Specifically, AIS microtubules contain a high density of non-hydrolysed GTP–tubulin clusters called GTP islands. These GTP islands can help recruit EBs to the AIS, as shown by electron and super-resolution microscopy[61,62]. EBs can bind to the AIS scaffold protein ankyrin-G[63,64], which is itself bound to βIV-spectrin (also known as SPTBN4), which interacts with actin[65] (FIG. 2a).

Dynamics of axonal microtubules. Once the axon is specified, it grows at a sustained speed (between 1 and 100 µm per hour), and there is a long-standing debate on how microtubules are supplied at a high rate to support this elongation. Issues include where the microtubules are nucleated — the process by which new microtubules start to polymerize — and how they are eventually

Table 1 | **Probes that label the axonal actin cytoskeleton and actin-binding proteins**

Probe	Imaging technique	Refs
Probes used to label the submembrane periodic scaffold		
Phalloidin	STORM	65,127,131,132,138
	STED	136,137
SiR–actin	STED*	130,131,133
	STED	140
	SIM	135
Anti-α-adducin antibody	STORM	127,139
Anti-βII-spectrin antibody	STORM	65,127,132,134, 138,139,172
	STED	136
βII-Spectrin–mMaple3	PALM*	139
βII-Spectrin–GFP	Epifluorescence*	139
Anti-βIV-spectrin antibody	STORM	65,127,131,132
Probes used to label actin waves		
Phalloidin	Epifluorescence	149
	SIM	153
GFP–β-actin; mRFP–β-actin	Epifluorescence*	155
paGFP–β-actin or dendra–β-actin	Photoactivation and/or conversion*	152
F-tractin–mCherry	Epifluorescence*	153
LYN–mCherry (membrane marker)	Epifluorescence*	153
THY1–YFP (neuronal marker)	Epifluorescence*	155
Anti-cortactin antibody	Epifluorescence	152
Anti-shootin antibody	Epifluorescence	152
Probes used to label actin trails and hot spots		
GFP–UTRN-CH	Epifluorescence*	148
LifeAct–mTurquoise	Epifluorescence*	148

CH, calponin homology domain; GFP, green fluorescent protein; LYN, tyrosine-protein kinase Lyn; mRFP, monomeric red fluorescent protein; PALM, photoactivated localization microscopy; SiR, silicon-rhodamine; SIM, structured illumination microscopy; STED, stimulated emission depletion microscopy; STORM, stochastic optical reconstruction microscopy; THY1, Thy-1 membrane glycoprotein; UTRN, utrophin; YFP, yellow fluorescent protein. *indicates techniques that were used for live imaging.

transported into the growing parts of the axon. One view is that microtubules are nucleated at the centrosome in the cell body and then severed by the enzymes katanin and spastin to generate smaller fragments. According to this view, the fragments are then transported into the axon, where they integrate into longer microtubules and elongate axons[66,67] (reviewed in REF. 68). However, laser ablation of the centrosome in neurons suggests that it is dispensable for axon growth[69]. Moreover, the free minus ends of most neuronal microtubules are not attached to the centrosome[48]. Genetic evidence in *Drosophila melanogaster* neurons also suggests that axonal growth is independent of centrosome nucleation activity[70]. Finally, the microtubule nucleation machinery at the centrosome is lost during neuronal maturation[69,71], and super-resolution microscopy reveals that microtubules lose their radial organization around the centrosome soon after maturation[72]. This raises the question of what

Centrosome
An organelle that nucleates and controls the organization of microtubules and regulates cell-cycle progression.

the alternative nucleation site for axonal microtubules might be. In non-neuronal cells, microtubules can be nucleated from isolated γ-tubulin clusters, the Golgi apparatus, plasma membrane or other microtubules[73]. Golgi-nucleated microtubules have been described in dendrites of *D. melanogaster* neurons[74,75], but it is unclear whether they also exist in axons. A nucleation mechanism that involves microtubule side-branching from γ-tubulin–augmin complexes has been recently proposed as a source of axonal microtubules[76] (FIG. 2a). As an alternative to nucleation, severing of existing microtubules by katanin and/or spastin along the axon, followed by growth of the resulting fragments, could increase and renew the axonal microtubule pool. An appeal of this mechanism is that it would intrinsically conserve the uniform plus-end-out orientation of the resulting network[77].

Even the highest estimates of microtubule length are far shorter than the total length of a typical axon, and axons must therefore contain a succession of overlapping microtubules throughout their length. The plus ends of these microtubules are usually in a state of dynamic instability, alternating between phases of growth (polymerization), pauses at a constant length and catastrophe (fast depolymerization)[78], whereas the non-centrosome-bound minus end is bound to stabilizing minus-end proteins[79,80]. Several minus-end-anchoring proteins in the axon have recently been identified. In *C. elegans* axons, PTRN-1 (the homologue of the *D. melanogaster* minus-end-binding protein patronin[81]) localizes as small puncta along the axons and stabilizes microtubules[82,83], as shown by live-cell imaging. In mammalian neurons, calmodulin-regulated spectrin-associated protein 2 (CAMSAP2) stabilizes microtubules by binding to their minus ends and is concentrated at the axon hillock and as puncta along the axon[72] (FIG. 2a). Other, yet unknown minus-end-anchoring proteins are likely to be present in neurons, particularly along axons[79].

Although microtubules are generally more stable in neurons than in non-neuronal cells, their stability depends on the maturation stage and cellular localization[84]. Axonal microtubules are more stable than their dendritic counterparts[85], and individual microtubules are known to have both stable and unstable domains along their length[86,87]. A current challenge is to visualize not only the growing ends of axonal microtubules (by using fluorescent EBs) but also the stable segments that represent the vast majority of axonal microtubules[77]. A recent study cut the microtubules at a precise point with laser microsurgery and then observed the growth of newly created microtubule ends labelled by fluorescent EBs, allowing the visualization of stable microtubules in axons by their subsequent growth[39]. However, this method does not reveal microtubules that do not grow after being cut. Here again, super-resolution microscopy with optimized probes[56] may provide answers in the near future by comprehensively mapping the organization of the axonal microtubule network.

Chemical modifications to tubulin are important for regulating microtubule structure and function. In addition to the hydrolysis of β-tubulin-bound GTP to GDP, tubulin undergoes several post-translational

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modifications (PTMs) — including detyrosination, polyglutamylation and acetylation — that accumulate on more stable microtubules[88]. A newly characterized PTM, polyamination, may be responsible for the maintenance of a stable, cold-resistant microtubule population in axons[84,89]. PTMs can alter the affinity of motors and microtubule-associated proteins[90,91] and are also likely to regulate the cytoskeletal network architecture by modifying the structural properties of microtubules[92]. AIS microtubules are rich in PTMs, although the role of these modifications in the selective recruitment of axonal cargoes is unclear[93–95]. The advent of new tools to visualize specific tubulin modifications in living cells[96,97], as well as techniques to locally perturb microtubules[98], will undoubtedly help us to dissect the extent and role of PTMs in the axonal architecture.

Axonal transport of tubulin. As noted above, pulse-chase radiolabelling studies showed that most tubulin is transported in the SCa, whereas a smaller fraction is conveyed in the slightly faster SCb[28]. Although the latter fraction of tubulin is largely soluble in a cold, low-salt detergent (Triton-X 100) — indicating that it may exist in a non-polymeric form — approximately half of the slower fraction in SCa is insoluble, suggesting compositional differences in the two transport components[99]; however, the importance of these differences has been unclear. Studies also suggest that only a small fraction of tubulin in axons is transported[100], which is relevant for the interpretation of studies that use fluorescent markers in cultured neurons. A number of studies in the 1980s and 1990s directly visualized the dynamics of fluorescent tubulin in axons. The consensus was that most microtubules in axons are stationary (reviewed in REF. 28). However, other studies showed that taxol-stabilized tubulin fragments injected into the squid axon can move with an anterograde bias and revealed the presence of short, motile microtubules at branch points of cultured neurons[101,102], suggesting that axonal microtubules can move.

In squid axons, injection of fluorescent tubulin resulted in the slow, anterograde advance of the bolus, at overall rates that were consistent with slow axonal transport[103,104]. Addition of the microtubule-depolymerizing drug nocodazole in this system did not impair this biased motion, indicating that tubulin may be able to move in an unpolymerized state[103]. In 2002, by use of a photobleaching-based paradigm, it was shown that linear structures containing fluorescent tubulin (presumably short microtubules) moved rapidly in cultured rat sympathetic neurons, with a significant overall anterograde bias[105]. However, the movement was very infrequent (approximately one event every 4 minutes) when compared with that of vesicle transport, and the moving structures spent about half of their time pausing (reviewed in REF. 106). Similar dynamics had also been seen for neurofilament transport, leading to the concept that the overall slow movement seen in the radiolabelling studies is a summation of rapid movement interrupted by long pauses (the 'stop and go' model, see below).
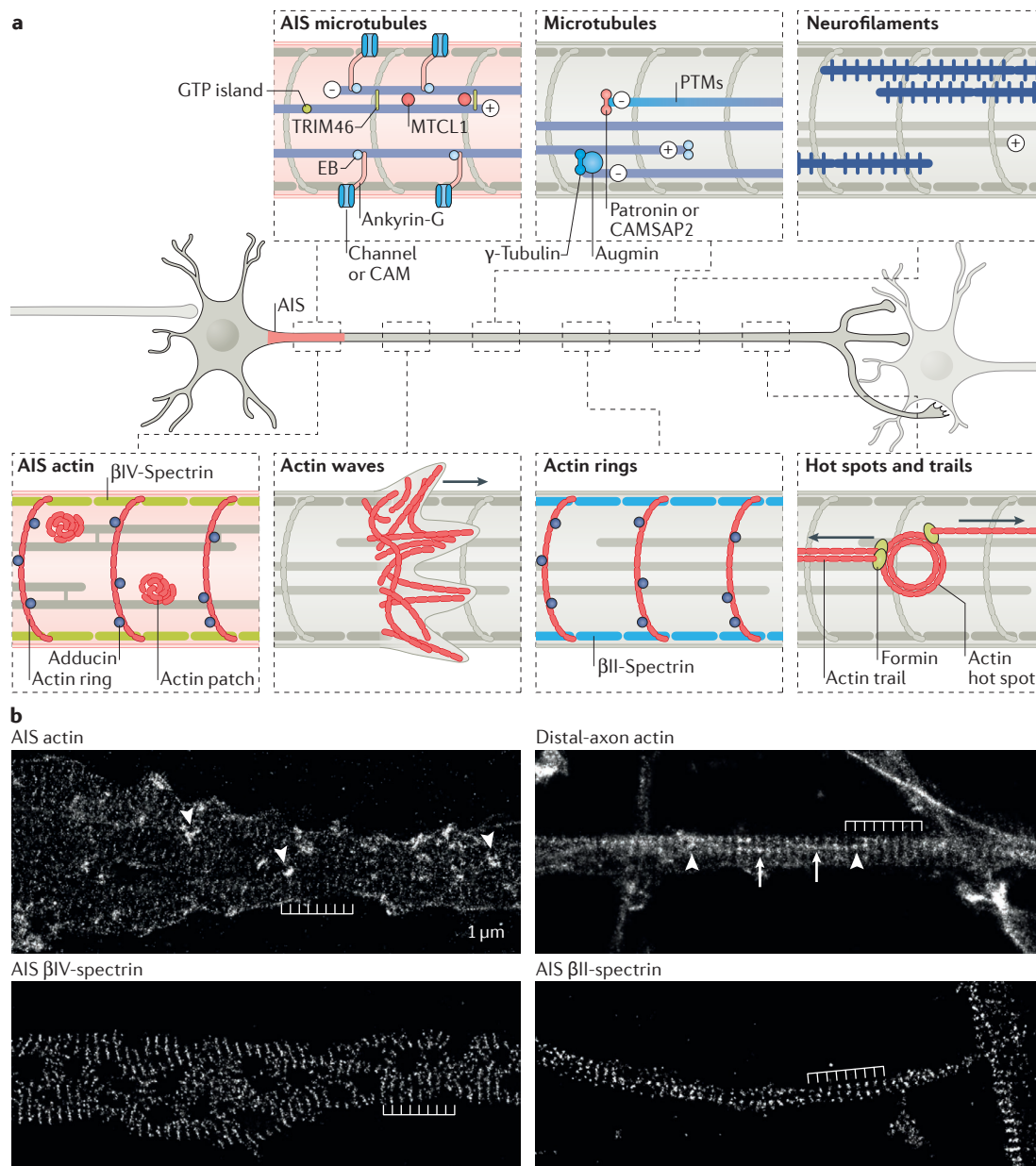
Figure 2 | **The cytoskeleton of the axon shaft. a** | The axon shaft is divided into the axon initial segment (AIS) and the distal axon shaft. In the AIS, microtubules (top left panel, blue) are oriented uniformly with the minus end (–) towards the cell body and the plus end (+) towards the distal axon. They form AIS-specific fascicles enriched in GTP islands and the associated proteins tripartite motif-containing protein 46 (TRIM46) and microtubule crosslinking factor 1 (MTCL1). End-binding proteins (EBs) link microtubules to ankyrin-G, which in turn binds to spectrin and anchors membrane proteins (channels and cell adhesion molecules (CAMs)). Microtubules in the distal axon (top centre panel) are also uniformly oriented and enriched in post-translational modifications (PTMs). They are anchored at the minus end by proteins such as patronin, calmodulin-regulated spectrin-associated protein 2 (CAMSAP2) and the γ-tubulin–augmin complex and are decorated with EBs at the growing plus end. Neurofilaments (top right panel, dark blue) are stable assemblies that fill the intra-axonal space. AIS actin (bottom left panel, red) is organized as submembrane rings enriched in adducin that are spaced every 190 nm by βIV-spectrin, as well as actin patches inside the axon. In younger neurons, actin waves (bottom centre panel) propagate along the axon shaft. In the distal axon, the actin rings are also spaced every 190 nm, but by βII-spectrin. Dynamic actin hot spots (bottom right panel) nucleate actin trails formed by fast, formin-mediated polymerization. **b** | Stochastic optical reconstruction microscopy (STORM) images of actin and spectrin in the AIS and distal axon. Actin in the AIS (upper left panel) organizes as 190 nm periodic rings (comb) and clusters (arrowheads). In the distal axon (upper right panel), actin rings are seen (comb) along with long actin polymers (arrows) and focal accumulations (arrowheads) that likely represent actin trails and hot spots[146]. Spectrins connect the actin rings and show the same 190 nm periodicity (indicated by the combs): βIV-spectrin in the AIS (lower left panel) and βII-spectrin along the distal axon (lower right panel). Left panels and lower right panel in part **b** are adapted with permission from REF. 65, upper right panel in part **b** is adapted with permission from ©Ganguly *et al. J. Cell Biol.* **210**, 401–417 http://dx.doi.org/10.1083/jcb.201506110 (2015) (REF. 148).

What are the mechanisms underlying this transport? Using live-cell imaging and fluorescent probes, more recent studies suggest that dynein motors are involved in both anterograde and retrograde transport of microtubules[107,108], whereas super-resolution microscopy has started to unravel the elusive structure of slow-transport cargoes[109]. Microtubule movement in other cell types is well established, providing a precedent[110,111]. However, further studies are needed to clarify the role of soluble tubulin pools in this transport, if any, and to characterize the short mobile microtubules.

Neurofilaments in the axon shaft

Organization of neurofilaments. Neurofilaments are heteropolymers composed of subunits of low, middle and high molecular weight, along with one of the two recently discovered forms of intermediate filament protein, α-internexin (in the CNS) or peripherin (in the PNS) and are organized as extensive parallel networks that fill the axoplasm (BOX 1; FIG. 1). In classical electron microscopy studies using the freeze-etching technique, neurofilaments appeared to be interconnected with inter-filament cross bridges that are approximately 3–5 nm in diameter[112]. Although the function of these cross bridges is not clear, they have been proposed to help to maintain the integrity and density of the neurofilament network and to keep the neurofilament network anchored to the plasma membrane and microtubules[112] (FIG. 2).

The assembly of neurofilaments starts with heterodimerization of two neurofilament proteins with their N-terminals aligned. These dimers then form a staggered tetramer with the N-terminals of the assembled dimer on each side, resulting in a nonpolar assembly. Approximately eight of these tetramers are assembled laterally to form unit-length filaments, which then anneal end to end to form neurofilaments[113] (BOX 1). Although neurofilaments are stable polymers with long half-lives, recent live imaging of fluorescently tagged neurofilaments revealed a dynamic process where neurofilament severing and annealing are common[114]. The mechanisms underlying this dynamic behaviour are largely unknown, and the precise relationship between neurofilaments and other cytoskeletal structures have not yet been explored by super-resolution microscopy (although a recent study used structured illumination microscopy (SIM) to examine intermediate filaments in glial cells[115]).

Axonal transport of neurofilaments. The slow axonal transport (SCa) of neurofilament proteins has been well documented by classic radiolabelling studies[25,116–118]; however, the form that is transported (monomer, oligomer or polymer) was debated for a long time[119,120]. Because neurofilaments are highly abundant in axons, the motility of individual fluorescently tagged neurofilaments could not be easily visualized. Taking advantage of the discontinuous neurofilament network in axons of cultured sympathetic neurons, live imaging studies have now unequivocally demonstrated the movement of neurofilament polymers[36,37,121]. Surprisingly, however, the movement of individual neurofilament polymers was rapid, resembling the velocities of fast transport cargoes. Unlike fast cargoes, neurofilament movement was very infrequent and the cargoes paused for prolonged periods during transit. This led to a 'stop and go' model for slow transport, which proposes that individual cargoes move rapidly but intermittently, leading to a slow overall movement of the population (reviewed in REF. 106). More recent studies suggest a model in which larger neurofilaments are largely stationary but are severed into smaller polymers that can be transported[114]. Conventional kinesins and dynein are involved in neurofilament transport[122], although the molecular mechanisms underlying the intermittent transport are still unclear.

Actin in the axon shaft

Organization of actin. Early electron microscopy studies showed a layer of sub-plasmalemmal actin along axon shafts[123], but intra-axonal actin was difficult to discern owing to the dense packing of cytoskeletal polymers and organelles within the lumen[6,7]. However, phalloidin staining of axoplasm extruded from squid giant axon showed extensive deep actin filaments[124]. 'Touch preparations' — in which a carbon-coated electron microscopy grid is touched against the surface of axoplasm to transfer a thin layer of the sample — suggested that actin filaments in axons were rather short, averaging only approximately 0.5–1 μm in length, and were in close apposition with microtubules[125]. However, such touch preparations are inherently disruptive and need to be interpreted with caution[125]. Experiments where skeletal myosin subfragment 1 was used to coat the actin filaments in a characteristic arrowhead pattern suggested that the polarity orientation of axonal actin filaments was mixed[125]. Phalloidin staining of axons in cultured neurons showed a patchy distribution, thought to reflect a combination of sub-plasmalemmal and deep actin[126]. Thus, until recently, the prevailing model of actin organization suggested that axon shafts have a coat of actin underneath the plasma membrane as well as deeper, short actin filaments with mixed polarity[35]. However, new information from super-resolution microscopy and live imaging has substantially revised this view.

In 2013, stochastic optical reconstruction microscopy (STORM, see BOX 2) was used to re-evaluate axonal actin organization, using the classical probe phalloidin[127]. Surprisingly, the patchy, apparently unremarkable actin seen only by routine light microscopy was shown to be composed of a periodic cytoskeletal assembly that can be seen by sub-diffraction microscopy techniques. Specifically, axons contain circumferential rings of actin that wrap underneath the plasma membrane and are spaced at approximately 190 nm (REF. 127) (FIG. 2b). Actin rings are connected by tetramers of spectrin, composed of βII-spectrin (also known as SPTBN1) and a still-unknown α-spectrin subunit, that are also approximately 190 nm long[128,129]. This periodic organization of actin was also seen in living neurons by using stimulated emission depletion microscopy (STED, see BOX 2)[130,131]. Moreover, the periodic actin–spectrin undercoat has been shown to be present along myelinated axons[131,132] and to be conserved in a variety of neuronal types and organisms[133–135].

Microtubule-associated proteins
(MAPs). The repertoire of proteins that bind to microtubules. They can associate with the microtubule lattice or with the minus end or plus end of the microtubules. Microtubule-associated molecular motor complexes are also MAPs.

It has been proposed that this actin–spectrin assembly could be an intrinsic property of cytoskeletal organization in thin processes, because similar periodicity in dendritic spine necks and the fine processes of glial have been reported[133,134,136,137]. Recently, a 2D polygonal lattice structure — resembling the erythrocyte membrane skeleton — was reported in larger compartments such as the cell body and large dendrites[138].

The actin–spectrin periodic complex appears to be immobile and stable, as shown by βII-spectrin fluorescence recovery after photobleaching and live-cell photoactivated localization microscopy (PALM, see BOX 2) experiments, and is also partially resistant to actin and microtubule depolymerization[139]. During development, actin rings first appear along the most proximal axon segment and have been proposed to 'propagate' along the shaft as the axon grows, although the rings themselves do not appear to move[139]. Mechanisms underlying the initial assembly of the rings and their progression along the shaft are unclear, but slow axonal transport of ring components may play a role (see below). Actin rings also contain adducin, a capping protein that binds to and stabilizes the ends of short actin filaments[127,140]. For this reason, it is thought that actin rings are composed of numerous short and stable actin filaments[141], although the actin rings have not yet been seen by electron microscopy. Interestingly, an absence of adducin increases the diameter of the actin rings, presumably because individual short filaments forming the ring can no longer be capped[140].

Actin rings are present along the entire axon, including the AIS; however, the β-spectrin subunits in the AIS and the more distal axon are different[65,127]. As the AIS is established, the βII-spectrin in this region is replaced by βIV-spectrin[139]. This switch contributes to the establishment of segregated submembrane scaffolds along the axon: βIV-spectrin–ankyrin-G scaffolds in the AIS and βII-spectrin–ankyrin-B scaffolds in the distal axon[142]; however, the relevance of this differential assembly is still unknown. In addition to actin rings, the AIS contains longitudinal actin filaments and actin clusters[131,143]. Some of these actin clusters are likely to correspond to postsynaptic sites, as the AIS is innervated by GABAergic neurons[131]. However, there are larger (approximately 1 μm in diameter) 'actin patches' in the AIS[143,144], and these may play a role in sorting vesicular transport by capturing myosin-bearing vesicles[145–147].

Although it was known that axons have both sub-plasmalemmal and deep actin[125], the characteristics of the latter (the filament length, dynamics and polarity, for example) were unclear. A recent study examined axonal actin by using filament-selective probes and low-light live imaging[148] (TABLE 2). Deep axonal actin was shown to be surprisingly dynamic: distributed along the length of the axon were focal actin 'hot spots', where actin continuously polymerized and depolymerized. Additionally, elongating actin polymers were seen along the long axis (these averaged approximately 9 μm in length, with the longest ones being up to 40 μm in length), extending in both anterograde and retrograde directions. These were termed 'actin trails' (FIG. 2b). The actin trails often seemed

Table 2 | Components of axonal actin assemblies and actin-binding proteins

Protein	Organization and/or function	Refs
Components of the submembrane periodic scaffold		
α-Adducin	Present at actin rings. Involved in capping of actin filaments and regulation of axon diameter	127,139,140
βII-Spectrin	Links actin rings in the distal axon. Involved in regulation of scaffold formation	65,127,134,139
βIV-Spectrin	Links actin rings at the AIS	65,127,139
Ankyrin-B	Bound to βII-spectrin in the distal axon	127
Ankyrin-G	Bound to βIV-spectrin at the AIS	65,127,139
Nav channels	Anchored to ankyrin-G at the AIS	65,127
Kv1.2	Associated with the actin rings at the AIS	132
Neurofascin	Anchored to ankyrin-G at the AIS	65,131
Components of actin waves		
Doublecortin	Controls the frequency of waves and promotes axon specification and branching	151
Shootin1	Couples actin filaments to the substrate	152
RAC1	Induces actin wave	153
ARP2/3, CDC42, RAP1B and cofilin, Lim kinase and slingshot	Function not explored in actin waves but likely to be involved in the movement of waves	153,155
Ezrin, GAP43 and cortactin	Function not explored in actin waves but likely to be involved in actin dynamics	149
Components of actin trails and hot spots		
Formins	Involved in the regulation of trail numbers and elongation	148

AIS, axon initial segment; ARP2/3, actin-related protein 2/3 complex; CDC42, cell division control protein 42 homologue; GAP43, growth associated protein 43; Kv1.2, voltage-gated potassium channel KCNA2; Nav channels, voltage-gated sodium channels; RAC1, Ras-related C3 botulinum toxin substrate 1; RAP1B, Ras-related protein 1B.

to arise from regions of hot spots, suggesting a scenario in which actin filament polymerization at hot spots created the nidus of assembly that initiated polymer elongation. The rate of elongation was rapid (approximately 1 μm s^{-1}) and depended on linear polymer growth mediated by formins and not on the actin-branching complex ARP2/3 (REF. 148). Collectively, the data indicate that axon shafts have two cytoskeletal networks: stable sub-plasmalemmal rings, composed of circumferential actin–spectrin assemblies and actin trails, a dynamic deep actin network consisting of focal actin hot spots and elongating filaments meandering along the shaft.

Besides rings and trails, developing axons also have other intriguing actin assemblies. Actin 'waves' are growth-cone-like structures that periodically emerge at the base of the axon, travelling along its length, up to the axon tip[149–154] (FIG. 2b). Best studied in cultured neurons, actin waves are present in all neurites before differentiation but are more frequent in the nascent axon[155]. These waves disappear as axons mature and make contacts with other neurons (reviewed in REF. 156). Similar waves of actin have also been described in other cell types such as fibroblasts, neutrophils, keratinocytes and *Dictyostelium discoideum*, where they are called travelling waves or t-waves[157]. Distal-axon 'actin patches' that may be similar to the ones found in the AIS have also been described in a variety of neurons (reviewed in REF. 158). Unlike actin waves, the patches are non-protrusive, focal accumulations within the shaft. Available data support the notion that actin patches in distal axons are involved in the emergence of axonal filopodia and branches[159,160].

Axonal transport of actin. Slow axonal transport of actin has been demonstrated in many different species by pulse-chase radiolabelling studies[28,29]. In rat motor neurons, for instance, the average rate of transport is approximately 3 mm per day, moving largely in SCb. However, actin rates in CNS neurons are much lower, and in most experiments, smaller 'peaks' of transport fractions were also seen (a smaller trailing peak of approximately 1.5 mm per day was seen in rat motor neurons, for example[117]). There is also a lagging trail of actin along axons as the radiolabelled 'wave' progresses, which may reflect actin deposition along axons[99,116]. Because actin is highly dynamic, any model of actin transport must also account for the dynamic exchange of monomeric and filamentous forms as well as explain the transport of actin in mature myelinated axons. Although the overall transport of actin in axons is well characterized, the underlying molecular mechanisms are essentially unknown. One study reported very-short-range saltatory motion of actin densities in cultured axons[161], but it is unclear how this bidirectional short-range movement can lead to an overall slow, anterogradely biased transport. The actin hot spots and trails described above[148] might play a role in the axonal transport of actin; however, mechanistic details are lacking. Finally, although actin waves seen in developing neurites have been proposed to play a role in actin transport, this appears unlikely because their infrequent nature, periodicity and transient nature seem inconsistent with the slow, constitutive transport observed in radiolabelling experiments.

Function of axonal actin assemblies. Studies in non-neuronal cells have shown that, although long-range cargo transport is dependent on microtubules, actin filaments may contribute to short-range transport[162]. Actin in axon shafts has also been thought to play a similar role in assisting vesicle transport[163]. The traditional view that axons contained only short actin filaments, together with the intermingling of actin filaments and microtubule bundles in axons, seemed compatible with this model[124,125]. However, the recent discoveries of new actin assemblies in axons suggest that this simplistic view is at best incomplete.

From a mechanical perspective, submembrane rigid actin rings connected to a supple elastic cylinder of spectrins could provide both high resistance and flexibility to axons that may be critical in maintaining the axon form and function[164,165]. In support of this idea, axons of *C. elegans* lacking β-spectrin readily break when these worms move[166], and the actin–spectrin undercoat is thought to maintain the mechanical properties of axons[167,168]. A combination of genetic mutants, pharmacologic treatments and SIM (see BOX 2) in *D. melanogaster* neurons allowed rapid screening of a large number of microtubule and actin regulators. This implicated the actin–spectrin undercoat in the maintenance of axonal microtubule organization[135]. This could be mediated by the classical partner of spectrin, ankyrin, because large ankyrin isoforms also contribute to microtubule spacing and organization in *D. melanogaster* axons[169]. However, the function or functions of actin rings have still not been established. Perhaps more focused ways of perturbing actin, such as the confinement of axons within microfluidic devices[170], the use of genetically encoded (and potentially subcellularly targeted) actin disassembly factors[171] or optogenetic perturbation, may help.

Because actin filaments are a necessary component of the AIS diffusion barrier that restricts membrane diffusion between the cell body and the axon[16,23], actin rings at the AIS might play a role in this diffusion barrier. Using correlative single-particle tracking and super-resolution microscopy, it was recently found that in the AIS (but not the distal axon), the lateral mobility of a glycophosphatidylinositol (GPI)-anchored protein was restricted to the plasma membrane between actin rings[172]. Specifically, the movement of the GPI probe was shown to be slowed when it was close to actin rings but not when it was midway between them, where membrane proteins (channels and CAMs) are anchored to ankyrin-G. The authors of this study thus favoured an 'actin fence' model — in which the actin rings act as barriers to lateral membrane diffusion — rather than a 'pickets' model in which tethered membrane proteins between the rings act as barriers[24,65,173]. As an alternative to corralling by actin itself, diffusion could also be impeded by the potassium channel Kv1.2 (also known as KCNA2), which (in contrast to other AIS membrane proteins) has been recently shown to localize precisely at actin rings[132]. It is also possible that other unidentified, AIS-specific 'pickets' might selectively impede the lateral diffusion of plasma membrane proteins at this locale, explaining the lack of restriction of diffusion seen in distal axons.

Microfluidic
Of a method to manipulate cells (for seeding, incubation or labelling) at the sub-millimetre scale by using small volumes of medium that are pumped into miniaturized culturing devices.

En-passant boutons
Presynaptic specializations along axons that make contact with downstream neurons, as opposed to synaptic terminals at the extremity of axons. In hippocampal and cortical neuronal cultures, most presynapses are *en-passant* boutons.

The functions of deep actin assemblies such as hot spots and trails are not well understood[174]. Available data suggest that these dynamics help deposit actin at en-passant boutons and sustain presynaptic function[148]. As noted above, it also seems likely that the hot spots and trails have a role in the slow axonal transport of actin (and perhaps actin-binding proteins as well); however, it is unclear how this could happen mechanistically. The elongating actin trails resemble 'actin comets' observed in *Listeria monocytogenes*[175], and one possibility is that they may generate a propulsive force that could move vesicles or other cargoes. However, inhibition of actin trails does not have much effect on overall vesicle transport[148] (although it remains possible that a subset of vesicles are transported in this manner). If all the hot spots and trails in an axon are considered to be part of an integrated actin 'network', the dynamic cytoskeletal organization could be seen as a source of assembly-competent actin monomers throughout the axon shaft. This may be an important homeostatic mechanism, allowing local actin-dependent physiological events (such as filopodial extension, axon branching, bouton formation and actin-dependent signalling) to occur rapidly, and future studies may provide clarity.

Conclusion

Easy to see but difficult to understand, the cytoskeleton of the axon shaft has long been the subject of speculative cartoons. Although much remains to be clarified, recent advances in super-resolution microscopy — along with the development of specific tools to probe or disrupt the cytoskeleton — now allow us to examine these assemblies with a spatial and temporal precision that was not possible before. The discovery of unique architectural assemblies of actin in axons forces us to question the roles traditionally assigned to actin, and one wonders if this might be true for the other cytoskeletal proteins as well. For instance, neurofilaments and mitochondria reportedly interact in axons[176], but the underlying structural and functional associations are unclear. Moving forward, we can predict four general lines of investigation that are likely to have an impact on our understanding of the axonal cytoskeleton: further definition of the nanoscale architecture and dynamics of the cytoskeletal proteins and the generation of quantitative models; clarification of the individual roles of diverse cytoskeletal assemblies; appreciation of the architectural and mechanistic links between the cytoskeletal elements and axonal organelles, cytosolic proteins and the plasma membrane; and, ultimately, understanding the guiding principles that unite the entire axonal cytoskeletal meshwork to orchestrate physiological processes.

The axonal cytoskeleton is clearly a complex and sophisticated machinery with a menagerie of players. Even with the seemingly detailed knowledge that we have today — and the sophisticated tools that are at our disposal — one feels that the axonal cytoskeleton holds many more secrets, which must await the next generation of cytoskeleton gawkers.

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