Title: Unrestrained growth of correctly oriented microtubules instructs axonal

microtubule orientation

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Abstract:

In many eukaryotic cells, directed molecular transport occurs along microtubules. Within

neuronal axons, transport over vast distances particularly relies on uniformly oriented

microtubules, whose plus-ends point towards the distal axon tip (+end out). However, axonal

microtubules initially have mixed orientations, and what breaks this orientation symmetry is

poorly understood. Using live imaging, physical modeling and simulations, we found that +end

out microtubules surpass a growth transition and undergo persistent long-term growth. In

contrast, oppositely oriented microtubules remain short. We confirmed experimentally and *in*

silico that the enhanced growth of +end out microtubules is critical for achieving uniform

microtubule orientation. Our data suggest that the accelerated growth kinetics of correctly

oriented microtubules instructs overall axonal microtubule orientation, thus enabling efficient

long-range transport essential for neuronal functioning.

One Sentence Summary: Enhanced growth of microtubules towards the axon tip is crucial for

establishing microtubule order needed for molecular transport.

Main Text:

Symmetry breaking is critical for many biological systems. An organism starts of as a single round

cell that divides and differentiates into many cells, tissues and organ systems. The neuron, with

its branched dendrites and sometimes exceedingly long axon, is one of the least symmetric cells

found in animals. Axons connect neurons with distant targets and thus enable long-distance

signal transmission throughout the body at high speed.

The enormous length of axons, which can extend over several meters in some vertebrate species,

poses substantial logistical challenges. Proteins and genetic material originating in the cell body

need to be actively transported down the axon. Transport occurs along microtubules, which are

long, polarized polymers that undergo stochastic cycles of growth and shrinkage (**FIGURE 1A**).

Motor proteins transport cargo either towards a microtubule's dynamic (i.e., growing or

shrinking) +end, or the stabilized -end.

In immature axons, microtubule orientation is mixed, with 50-80% of all microtubules having

their +end out. During early neuronal development, the fraction of +end out axonal microtubules

increases (*1*, *2*). In mature axons, ~95% of all microtubules point in the same direction (+end out)

(*3*, *4*), enabling polarized transport (*5*). Deficits in polarized transport have been associated with

human neurodegenerative diseases, such as Alzheimer's and Parkinson's disease (*5*). Despite the

importance of polarized transport in neuronal axons, the mechanism that breaks microtubule

symmetry is still poorly understood.

Microtubules in post-mitotic neurons are not attached to the centrosome (*6*). Nucleation of new

microtubules occurs from somatic Golgi (*7*) through elongation of severed pieces (*8*) or *de novo*

polymerization alongside existing microtubules (*9, 10*). These newly formed microtubules often

orient in the same direction as existing ones, enforcing any pre-existing orientation bias (*11, 12*),

which, however, by itself cannot explain the +end out orientation of microtubules in mature

axons. Furthermore, in axons, short microtubules pointing with their -end out can be transported

towards the cell body (i.e., away from the tip) by cytoplasmic dynein, a process likely assisted by

the microtubule bundling protein TRIM46 (*2, 13, 14*). However, also this mechanism is not

sufficient to explain how axonal microtubules orient with their +ends pointing away from the cell

body (see below).

To address how microtubule orientation in neuronal axons is achieved, we first quantified

microtubule growth in *Drosophila melanogaster* axons by labelling growing microtubule +ends

with the fusion protein EB1-GFP, which labels +ends with bright 'comets' (**FIGURE 1A-C**) (*15*). The

distance over which a comet moves in the axon is equal to the overall length d_g that is added to

a microtubule between the start of its growth cycle and the catastrophe leading to microtubule

shrinkage (**FIGURE 1A**). The direction of growth highlights whether a microtubule is oriented with

its +end away from (+end out) or towards (-end out) the cell body. Time-lapse movies of EB1-GFP

comets were converted into kymographs and analyzed using *KymoButler* (*16*) (**FIGURE 1C-D**).

The fraction of +end out microtubules increased over time and with increasing axonal length

(**SUPPLEMENTAL FIGURE 1**), confirming that microtubule orientation increases during development

(*1, 2*). Most microtubule growth events (~66%) were found within the first 20 μm from the

advancing axon tip (**FIGURE 1A,E**). Microtubule growth lengths per cycle, d_g , were significantly

higher near the axon tip compared to further away from it (**FIGURE 1F**). Furthermore, +end out

microtubules added significantly more length per growth cycle than -end out microtubules, with

the highest difference between +end out and -end out microtubules (~0.5 µm/cycle) found within

the first 10 µm from the axon tip (**FIGURE 1G**). Hence, the orientation of microtubules that added

more length per growth cycle (i.e., +end out microtubules) became the dominant microtubule

orientation during neuronal development, indicating a correlation between increased +end out

microtubule growth and the fraction of +end out microtubules in axons.

On long time scales, differences in microtubule growth length per growth cycle should primarily

affect average microtubule lengths l_{MT}. To test whether the rather small differences in d_g of ~0.5

µm/cycle lead to biologically meaningful differences in the average expected microtubule lengths

between +end out and -end out oriented microtubules, we formulated a 2-state Master equation

model of microtubule growth and shrinkage (see (*17*) & supplemental methods for details The

model distinguishes two regimes (**FIGURE 2A**):

$$l_{MT} = \begin{cases} d_g d_s/(d_s - d_g) & d_s > d_g & \text{('bounded' growth)} \\ \infty & d_s \leq d_g & \text{('unbounded' growth)} \end{cases} \qquad (1)$$

where d_s = lost length per shrinkage cycle.

When $d_s \leq d_g$, the average length added to the microtubule exceeds the average shrinkage length

per cycle so that a microtubule will exhibit net growth and elongate as long as physically possible

in its confined environment (called 'unbounded' growth). For $d_s > d_g$, however, growth is

'bounded' and average microtubule lengths follow an exponential distribution with a mean of

$d_s d_g/(d_s-d_g)$. In practice, this means that microtubules with $d_g \geq d_s$ would grow until encountering

a physical barrier (for example the axon tip), while microtubules with $d_s > d_g$ remain finite

(approximately 2 μm with d_s = 2.2 μm and d_g = 1 μm).

We determined the microtubule shrinkage per cycle d_s by co-expressing a Jupiter-mCherry fusion

protein (a tubulin marker) together with EB1-GFP in *D. melanogaster* axons. Microtubules

stopped growing when the GFP signal disappeared from their +end, indicating a catastrophe

event. Subsequent microtubule shrinkage was visualized by simultaneously imaging tubulin

(Jupiter-mCherry), and quantified by tracing the resulting tubulin edges in the dual colour

kymographs (**FIGURE 2B-D**). Axonal microtubule shrinkage lengths were d_s = 2.2 ± 0.2 μm / cycle

(mean ± s.e.m., **FIGURE 2E**). With this value for d_s, our model predicted the divergence of l_{MT} at

around d_g = 2.2 μm (**FIGURE 2F).** The measured values of d_g *and* d_s hence suggested that +end out-

oriented microtubules within 10 μm from the axon tip exhibit a considerably higher likelihood of

unbounded growth than -end out microtubules within that range and any microtubule further

away from the tip (**FIGURE 2F**). The enhanced growth of +end out microtubules near the axon tip

implied a higher chance of survival for +end out microtubules while leaving -end out microtubules

labile, thus establishing and maintaining overall axonal +end out microtubule orientation.

To test this hypothesis, we decreased microtubule growth using Nocodazole, a drug that disrupts

microtubule polymerization. We also decreased microtubule growth by increasing the osmolarity

of the cell culture medium (*18*, *19*), potentially by decreasing the space for microtubules to

polymerize into (*20*). In both approaches, axons exhibited significantly decreased +end out

microtubule growth at the axon tip (< 10 μm), with d_g < 1.7μm / cycle (**FIGURE 3A-E**). In agreement

with our prediction, microtubules within these axons were overall significantly less uniformly

oriented (**Figure 3D-E**), confirming an important role of the enhanced growth lengths per cycle

of +end out microtubules in establishing axonal microtubule orientation.

However, why do +end out microtubules grow longer in the vicinity of the axon tip? Local

gradients of microtubule growth-promoting factors could lead to an increase in +end microtubule

growth in that region (**Figure 1G**). Axon tips contain a multitude of different proteins and are the

site of protein synthesis (*21*). Locally synthesized microtubule growth-promoting factors include

microtubule stabilizing proteins, such as p150 (*22*, *23*), CRMP-2 (*24*, *25*), and TRIM46 (*13*, *14*), as

well as free tubulin and others (*26*).

Drosophila has a p150 homologue which, similar to murine neurons(*22*), is enriched in axon (but

not in dendritic) tips (**Figure 3F, H and Supplementary Figure 2**). These dendritic processes

exhibited no +end out microtubule orientation (**Supplementary Figure 3**). Inserting the measured

axonal p150 gradient profile into a mathematical model indicated that the observed gradient

should, in theory, be able to cause different growth behaviors for +end out and -end out

microtubules in axonal tips (**Supplementary Figure 4**). Accordingly, +end out microtubule growth

lengths per cycle d_g were significantly higher in axons than in dendritic processes -which exhibited

mixed microtubule polarities (**Supplementary Figure 3**).

To test this prediction further, we assessed microtubule dynamics in *wild-type*, *p150-RNAi*

expressing neurons, and in neurons from *p150[1]/+* mutant flies (**Figure 3G-K**). *p150[1]* (also known

as *Gl[1]*) mutants express a truncated p150-RNA transcript, which results in a dominant negative

phenotype (*27*). Both the expression of *p150-RNAi* and of dominant negative *p150[1]/+* led to a

significant decrease in +end out microtubule growth within 10 μm from the axon tip (**FIGURE 3L**).

In agreement with our model, microtubule orientation was significantly decreased in both *p150-*

RNAi and *p150[1]/+* axons compared to controls (**FIGURE 3M**), indicating that growth-promoting

protein gradients at the axon tip do indeed have an important role in regulating the overall

polarization of the axonal microtubule network.

Previous work showed that p150 accumulation at axon tips depends on the activity of the

microtubule-specific molecular motor protein kinesin 1 (*22*), which preferentially enters axons

over dendrites (*28*). Accordingly, disruption of kinesin 1 function with an RNAi treatment led to

considerable mislocation of p150 along axons (**SUPPLEMENTARY FIGURE 5A-D**). Again, the absence

of a p150 gradient in these neurons led to significantly decreased +end out microtubule growth

near the axon tip, and hence to an overall decrease in axonal microtubule orientation

(**SUPPLEMENTARY FIGURE 5E-I**). In addition to unbounded +end out microtubule growth identified

here, other mechanisms have previously been suggested to be involved in regulating the overall

axonal microtubule orientation, including dynein-mediated sliding of -end out microtubules

towards the cell body (*2, 13*) and augmin-mediated microtubule templating, that is nucleating

microtubules tend to have the same orientation as existing ones in the same cellular region) (*9,*

10).

Computer simulations of microtubule sliding and/or templating led to a steady state in which a

fixed fraction of the axon length (close to the cell body), remained enriched with -end-out

microtubules (**FIGURE 4A,B AND SUPPLEMENTARY FIGURE 6**). Similarly, the unbounded growth of

+end out microtubules alone (**SUPPLEMENTARY FIGURE 6C**) also failed to capture the high fractions

of +end out microtubules observed experimentally at the axon entry (**Figure 4B**). However,

integrating sliding, templating, and unbounded +end out microtubule growth in the simulation

(assuming that the likelihood of exhibiting unbounded growth corresponds to a successful

nucleation event) led to axons with overall +end out microtubule orientation recapitulating our

experimental results (**Figure 4C-D and Supplementary Figure 6**). This highlights the necessity of

multiple mechanisms to establish and maintain axonal microtubule orientation. Similarly, a

combination of different microtubule growth-promoting proteins (e.g., tubulin, TRIM46, CRMP2,

and/or p150) is likely to contribute to unbounded microtubule growth into the axon tip.

p150, which we investigated here as an example, was concentrated at axon tips but not enriched

at dendritic tips (**Supplementary Figure 2),** confirming that differences in the localization of

microtubule growth-promoting proteins correlate with differences in microtubule growth. p150

is mainly known for of its role in the dynactin complex, which is an important cargo adapter

protein for the molecular motor protein dynein (*29*). Since p150 and dynein are functionally

related, it is difficult to separate their individual contributions to microtubule growth and cell

body-directed sliding of -end out microtubules. However, it remains unclear whether p150 is

required for dynein-mediated microtubule sliding (*30*, *31*), and the *Drosophila melanogaster*

oocyte also contains a biased microtubule cytoskeleton whose orientation is, presumably,

maintained by p150 (*32*). Hence, while p150 is unlikely to induce unbounded microtubule growth

alone, it emerges as a key contributor to the establishment of microtubule orientation.

We propose the following model explaining the spontaneous establishment of microtubule

orientation in developing axons. Growth-promoting proteins accumulate at the axon tip due to

microtubule +end-directed transport by kinesin motors (**SUPPLEMENTARY FIGURE 5**), rendering

+end out microtubule growth into the axon tip unbounded. In contrast, short -end out

microtubules are more prone to depolymerization and/or transport away from the tip by dynein-

mediated cell body-directed sliding (*1*, *2*), thus contributing to the orientation bias of

microtubules in the axon. This bias is further enhanced by augmin-mediated templating to

establish and maintain a fully organized microtubule cytoskeleton (see **FIGURE 4E** for a schematic

summary). Future work will reveal whether other cellular systems use similar mechanisms to

organize their cytoskeleton.

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Supplementary Materials:

Materials and Methods

Figures S1-S5

Figure 1: Axonal microtubule orientation and localization regulate microtubule growth. (A) Schematic depicting

the microtubule growth and shrinkage cycle. Microtubules grow until they undergo a catastrophe, which initiates

microtubule shrinkage and can start growing again after a rescue event. During growth (but not shrinkage) EB1

localizes to microtubule tips. **(B)** First frame of a live cell imaging movie of axonal EB1-GFP dynamics. Bright dots

represent individual EB1-GFP puncta, which label growing microtubule +ends. **(C)** Maximum intensity projection of

a 200 seconds long movie depicting EB1-GFP dynamics in a *D melanogaster* axon. EB1-GFP density is increased

towards the tip. **(D)** Schematic showing how EB1-GFP live imaging movies were visualized and analyzed using

kymographs. The growing tips of +end out microtubules (blue) and -end out microtubules (red) were fluorescently

labelled with EB1-GFP (green tear drop shaped 'comets'). The same axon is shown at three different time points;

microtubules grow at their +end, where EB1-GFP is located. The axonal intensity profiles of all time points are plotted

underneath each other, resulting in a space-time grid called 'kymograph'. Connecting puncta between consecutive

kymograph lines with blue/red lines yields the overall displacement d_g for individual microtubule growth events.

Note that the red -end out microtubule stops growing in the 2nd frame and shrinks in the 3rd frame **(E)** Kymograph

showing EB1-GFP dynamics analyzed with *KymoButler* (*16*). Lines with a positive slope (blue, left to right upwards)

are microtubules growing with their +end towards the axon tip, lines with negative slope (red, left to right

downwards) are microtubules growing away from the tip. Horizontal bars indicate the growth lengths (d_g) for

individual microtubule growth cycles. **(F)** EB1-GFP comet density as a function of the distance from the axon tip.

Most microtubule polymerization occurred near the advancing axon tip. (N = 346, 20 biological replicates, $p < 10^{-20,}$

Kruskal Wallis test, $p < 10^{-7}$ for pairwise comparisons between bin 1-4, Dunn-Sidak post hoc test). **(G)** Microtubule

added length per cycle d_g as a function of distance from the axon tip. Microtubules grew longer in the vicinity of the

axon tip ($p < 10^{-20}$, Kruskal Wallis test, $p < 10^{-7}$ for pairwise comparisons of either bin 1 or 2 with any other bin, Dunn-

Sidak post hoc test). **(H)** d_g for +end out (blue) and -end out (red) microtubules grouped for growth in the distalmost

10 µm of the axon tip, and further away than 10 µm from the axon tip. Each dot represents the average of one axon

in the respective region, grey lines indicate median values. With d_g = 2.24 ± 0.00 µm / cycle (mean ± SEM), +end out

microtubules near the axon tip grew significantly longer than -end out microtubules (d_g = 1.68 ± 0.00 µm / cycle) and

microtubules located further away from the tip (N = 346 axons, 20 biological replicates; $p < 10^{-30}$, Kruskal Wallis test

followed by Dunn-Sidak post hoc test; *** $p < 10^{-4}$). Scale bars: 3µm.

Figure 2: Microtubule length depends on growth length per cycle. (**A**) Schematic highlighting the assumptions of

our two-state master equation model. Microtubules were assumed to occupy either a growing or shrinking state.

During a growth cycle, the average microtubule length increases by d_g, during a shrinkage cycle, microtubule length

decreases by d_s. Additionally, microtubules were able to stochastically switch between the two states as shown in

Figure 1A. (**B-D**) Kymographs from a *D. melanogaster* axon that expressed (**B**) EB1-GFP (green) and (**C**) Jupiter-

mCherry, a tubulin label (magenta). Individual microtubule shrinkage events, visible as (**C**) fluorescent edges and (**D**)

dashed white lines in the kymograph, yielded microtubule shrinkage lengths per cycle d_s. Yellow and blue markers

in **B** & **C** indicate start and end points of an individual shrinkage event and the inset in **D** highlights an individual

shrinkage event. Scale bars: 3µm. (**E**) Average d_s values for N=47 axons (3 biological replicates; mean: 2.20±0.20 µm

(± s.e.m). (**F**) A plot of the estimated overall microtubule length l_{MT} as a function of d_g with d_s = 2.20±0.20 µm. The

two solid black lines indicate the lower and upper bounds of the average microtubule lengths for a given d_g with d_s

= 2.0 or 2.4 µm. One can separate two regimes; "unbounded" and "bounded" growth, separated by a dashed line.

Blue and red horizontal lines represent median ± lower and upper quantiles for +end out and -end out microtubules,

353   respectively. The top two lines represent microtubules found at the distalmost 10 µm of the axon, the bottom lines

354   microtubules found further than 10 µm away. +end out microtubules close to the tip were considerably more likely

355   to exhibit unbounded growth than +end out microtubules further away from the tip and -end out microtubules.

Figure 3: Decreased microtubule growth decreases axonal microtubule orientation. (A-C) Representative

kymographs analyzed with *KymoButler* (*16*) from axonal processes treated with (**A**) 0.025% DMSO (control) for 8h,

(B) 5uM Nocodazole for 8h, **(C)** medium with increased osmolarity ("osmo+") for 22h. Growth of +end out

microtubules is shown as blue lines, -end microtubules are red. Scale bars = 3μm. **(D)** Added growth lengths per cycle

d_g of +end out microtubules at the distalmost 10μm from the axon tip for control (N = 107 axons from 5 biological

replicates), nocodazole-treated axons (N = 116 axons from 3 biological replicates), and axons cultured in osmo+

medium (N = 30, 2 biological replicates). Microtubule lengths increased significantly less per growth cycle in axons

treated with Nocodazole or osmo+ media than controls ($p < 10^{-4}$, Kruskal Wallis test, * $p < 0.05$ for pairwise

comparisons, Dunn-Sidak post hoc test). **(E)** The fraction of +end out microtubules, in the different groups.

Microtubule orientation was calculated by counting all microtubules that grew away from the cell body (blue lines

in kymographs) and dividing them by all growing microtubules (blue and red). This way, a kymograph with only blue

lines gives a value of 1 while an equal number of blue and red lines yields a value of 0.5. Microtubules in axons

treated with Nocodazole or osmo+ media were significantly less uniformly oriented than those in the control group,

i.e., they contained a larger fraction of microtubules pointing with their +ends toward the cell body (red lines in **(A-**

C)) ($p < 10^{-4}$, Kruskal Wallis test, ** $p < 10^{-2}$ for pairwise comparisons, Dunn-Sidak post hoc test). **(F-G)** Tubulin (top)

and normalized p150 (bottom) immunostaining of cultured *D. melanogaster* larvae axonal processes of **(F)** controls

and **(G)** neurons expressing *elav-gal4* UAS driven *p150-RNAi*. Large p150 puncta were found clustered around the

axon tip (arrow) in controls **(F)** but not in *p150-RNAi* axons **(G)**. Scale bars = 2μm **(H)** Normalized p150 fluorescence

intensity as a function of distance from the axon tip for *wild-type* axons (N = 33) and *p150-RNAi* axons (N = 32, 2

biological replicates). Lines represent mean ± s.e.m. for *wild-type* (grey) and *p150-RNAi* (magenta). P150

fluorescence intensities changed along the axon ($p < 10^{-70}$; Kruskal Wallis test). In *wild-type* axons, p150 was enriched

at the axon tip (* $p < 0.05$ between bin 1 and bin 3 or 4; pairwise comparisons with Dunn-Sidak post hoc test), but

not in *p150-RNAi* expressing axons ($p > 0.05$ for all pairwise comparisons). Overall, p150 expression levels were

diminished in *p150-RNAi* axons compared to *wild-type* (*** $p < 10^{-7}$ for any pairwise comparison between conditions).

(I-K) *KymoButler* output for kymographs of EB1-GFP expressed in **(I)** a *wild-type* axon, **(J)** an axon expressing p150-

RNAi, and **(K)** an axon in a *p150^1/+* mutant background. Scale bars = 3 μm. Blue/red lines represent microtubules

with +/-end out orientation, respectively. **(L)** +end out microtubule added length per cycle d_g for *wild-type* (N = 85,

9 biological replicates), *p150-RNAi* (N = 34, 3 biological replicates), and *p150^1/+* (N = 83, 6 biological replicates).

Microtubule growth lengths were significantly decreased in both *p150-RNAi* and *p150^1* conditions compared to

controls ($p < 10^{-9}$, Kruskal Wallis test, **$p < 0.001$, *$p < 0.05$, Dunn-Sidak post hoc test). **(M)** Microtubule orientation

for *wild-type*, *p150-RNAi*, and *p150^1/+*. Microtubules were less uniformly oriented in both *p150-RNAi* and *p150^1*

axons ($p < 10^{-9}$, Kruskal Wallis test, *** $p < 10^{-5}$ for pairwise comparisons with Dunn-Sidak post hoc test). Overall,

axonal microtubule orientation was decreased after chemical, physical, and genetic perturbations of microtubule

growth.

Figure 4: Unbounded microtubule growth into the advancing axon tip is required to establish axonal microtubule orientation. (A) Schematic showing the microtubule sliding simulation and its relevant parameters. **(B-D)** Simulation of microtubule dynamics. **(B)** Snapshot of a simulated axon with dynein-based sliding of microtubules. -end out microtubules accumulated within the proximal axon. **(C)** Snapshot of an axon simulated with sliding, augmin templating (new microtubules were likely oriented into the same direction as their surrounding ones), and higher likelihood of unbounded growth for +end out microtubules. Much like in real axons, most microtubules were oriented with their +end out throughout the axon. **(D)** Microtubule orientation along the normalized axon length for the simulations with sliding only (orange trace), with sliding, templating, and unbonded growth (green), and our experimental data (blue). Lines represent 30% to 70% quantiles. There was excellent agreement between the experimental and *in silico data.* **(E)** Summary of proposed mechanism for establishing microtubule orientation in axons. Red and blue lines represent -end out and +end out microtubules, respectively. Green drop shapes indicate unbounded microtubule growth into the axon tip for +end out microtubules. Kinesin 1 deposits microtubule growth

405   promoting proteins, such as p150, at axon tips (**SUPPLEMENTARY FIGURE 4**), leading to local unbounded growth of +end

406   out microtubules. Augmin templating and cell body-directed sliding of -end out microtubules further amplifies this

407   bias. All three mechanisms together lead to a +end out microtubule cytoskeleton.