1	Cytoskeletal assembly in axonal outgrowth
2	and regeneration analyzed on the nanoscale
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4	Running title: Cytoskeletal assembly in axons
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27 Abstract

28 The axonal cytoskeleton is organized in a highly periodic structure, the membrane-associated 29 periodic skeleton (MPS), which is essential to maintain the structure and function of the axon. 30 Here, we use stimulated emission depletion microscopy (STED) of primary rat cortical neurons 31 in microfluidic chambers to analyze the temporal and spatial sequence of MPS formation at the 32 distal end of growing axons and during regeneration after axotomy. We demonstrate that the 33 MPS does not extend continuously into the growing axon but develops from patches of periodic 34 β-spectrin II arrangements that grow and coalesce into a continuous scaffold. We estimate that 35 the underlying sequence of nucleation, elongation, and subsequent coalescence of periodic β -36 spectrin II patches takes around 15 hours. Strikingly, we find that development of the MPS 37 occurs faster in regenerating axons after axotomy and note marked differences in the 38 morphology of the growth cone and adjacent axonal regions between regenerating and 39 unlesioned axons. Moreover, we find that inhibition of the spectrin-cleaving enzyme calpain 40 accelerates MPS formation in regenerating axons and increases the number of regenerating 41 axons after axotomy. Taken together, we provide here a detailed nanoscale analysis of MPS 42 development in growing axons.

43 Introduction

The cytoskeleton plays a crucial role in the outgrowth and regeneration of axons. It enables the neuron to extend its axon to otherwise unknown lengths for cells. The cytoskeleton must withstand mechanical force and torsion but also needs to facilitate adaption and response to trauma or damage, like axonal lesions. Moreover, it builds the scaffold for the precise distribution of membrane channels and receptors along the axon and the basis for axonal transport, which are both essential for the function of the neuron.

- Recent advances in super-resolution microscopy (SRM) have opened up a door for nanoscale observations of biological structures. In light of these advances, Xu et al. found that the cytoskeletal proteins actin, spectrin, and adducin form highly periodic structures in the axon, the membrane-associated periodic skeleton (MPS) (Xu, Zhong, & Zhuang, 2013). Originally discovered using stochastic optical reconstruction microscopy (STORM), this periodic arrangement was soon confirmed using other SRM-methods like stimulated emission depletion (STED) microscopy (Lukinavičius et al., 2014).
- 57 The MPS is prevalent in a broad range of neuronal cell types across different species (D'Este 58 et al., 2016; He et al., 2016). It has also been shown in brain slices (Xu et al., 2013) and in living 59 neurons (D'Este, Kamin, Göttfert, El-Hady, & Hell, 2015). Several studies have highlighted a 60 broad range of involvements for the MPS in neuronal processes, like stabilization of axons 61 (Hammarlund, Jorgensen, & Bastiani, 2007), electrical conductivity (Costa et al., 2020), latticelike organization of proteins (Li et al., 2020; Vassilopoulos, Gibaud, Jimenez, Caillol, & 62 63 Leterrier, 2019; Xu et al., 2013; Ruobo Zhou, Han, Xia, & Zhuang, 2019), and control of axonal 64 diameter (Costa et al., 2020; Leite et al., 2016). It was established that the MPS consists of actin 65 rings, made up of short actin filaments, capped by adducin, interconnected by α - and β -spectrin tetramers (Leite et al., 2016; Xu et al., 2013). 66
- 67 Several studies have suggested that the development of the MPS within the growing axon 68 occurs in a proximal to distal pattern (Han, Zhou, Xia, & Zhuang, 2017; Lorenzo et al., 2019; 69 Zhong et al., 2014). The MPS starts to form close to the soma at an early stage during axon 70 development, at day in vitro (DIV) 2 in hippocampal rat neurons (Zhong et al., 2014). It then 71 develops and progresses along the growing axon over time.
- The role of the MPS in axonal degeneration has been studied previously (Unsain, Bordenave,
 et al., 2018; Wang et al., 2019). It was shown that axonal degeneration induced by trophic factor
 withdrawal led to a loss of the MPS (Unsain, Bordenave, et al., 2018; Wang et al., 2019), but
- 75 this was independent of caspase apoptotic pathways (Wang et al., 2019). Interestingly,

76 stabilization of F-actin using the drug cucurbitacin-B preserved the MPS and reduced axonal 77 degeneration (Unsain, Bordenave, et al., 2018). In the last decade, mechanical axotomy through 78 vacuum media aspiration in neurons cultured in microfluidic systems has proven to be a reliable 79 model to study axonal degeneration and regeneration in vivo (Park, Vahidi, Taylor, Rhee, & 80 Jeon, 2006; Taylor et al., 2005). 81 Despite this recent interest in the MPS and its development, the exact sequence and mechanism 82 by which the MPS extends to newly formed parts of the outgrowing or regenerating axon 83 remain unknown. Here, we provide a detailed nanoscale analysis of the MPS development in 84 growing axons and the cytoskeletal changes upon regeneration after axotomy. We found that 85 growing axons display a characteristic spatial and temporal increase of spectrin periodicity in a 86 gradient from the distal growth cone towards the proximal axonal regions. We demonstrate that 87 the building blocks in the formation of the MPS are periodic arrangements of β -spectrin II 88 (periodic patches) that grow in size and number and eventually coalesce. In regenerating axons, 89 the MPS develops faster and at more distal axon regions than physiologically outgrowing

90 axons.

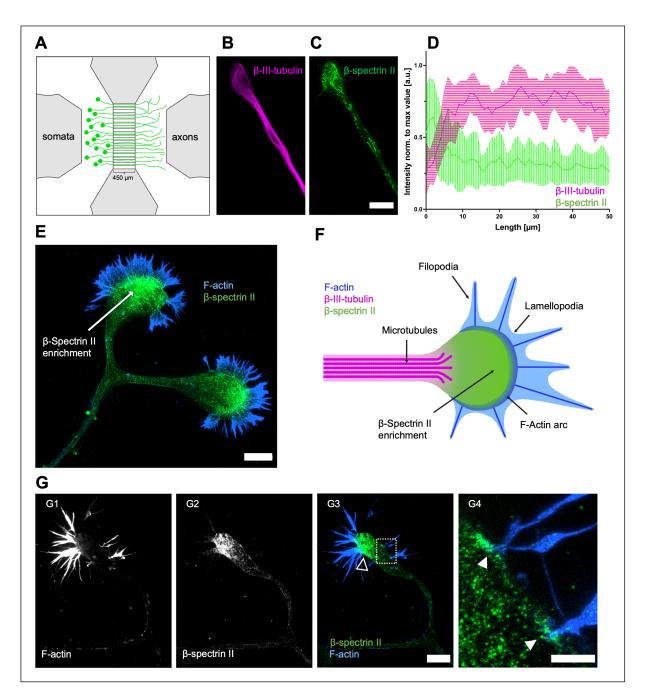
91 Results

92 Distribution of spectrin and tubulin along the axon and the growth cone in rat cortical 93 neurons

To analyze the organization of the developing MPS in physiologically outgrowing (native) and in regenerating (after axotomy) axons, we cultured primary rat embryonic cortical neurons in compartmentalized microfluidic chambers (Taylor et al., 2005) (**Figure 1, A**). The neurons were seeded in the soma compartment and extended their axons through microgrooves (length 450 μ m, width 3 μ m) to the axonal compartment allowing the separate analysis of axons. Cells were fixed on day 11, immunostained against different cytoskeletal proteins, and analyzed using STED microscopy.

101 First, we studied the general spatial distribution of essential components of the axonal 102 cytoskeleton along the native axon. We performed intensity line scans for β -spectrin II and β -103 III-tubulin along the growth cone (GC) and the adjacent axon (Figure 1, B-D, Supplemental 104 Figure 1). The intensity of tubulin was highest along the distal axon shaft and significantly 105 lower in the GC by 40% (0.44 ± 0.04 vs. 0.75 ± 0.02). In contrast, the intensity of spectrin was 106 highest in the GC and was 65% lower along the axon $(0.5 \pm 0.04 \text{ vs. } 0.3 \pm 0.04)$. Both spectrin 107 and actin intensity showed constant values further proximal up to 250 µm from the GC. We 108 thus conclude that spectrin is strongly enriched in the GC and integrated into the axonal 109 cytoskeleton in a lower concentration, while the opposite is true for tubulin.

The spectrin signal was strongest in the most apical central region and the transition region of the growth cone. In GCs co-stained for spectrin and F-actin, spectrin was enriched in the proximity to actin filaments. **Figure 1**, **E** and **F** show the location of the enrichment of spectrin in the GC relative to the F-actin-dominated filopodia. Especially at the base of F-actin bundles, the spectrin signal was enhanced and colocalized with actin. Exemplary micrographs are shown in **Figure 1**, **G**. This suggests a stabilizing or seeding function of spectrin for the outgrowing actin filaments in GCs.



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118 119 120 121 122 123 124 125 126 Figure 1. Distribution of β-spectrin II and β-III-tubulin along the axon and the growth cone in native rat cortical neurons. (A) Schematic drawing of a microfluidic chamber seeded with primary cortical neurons in the somatic department, transduced with AAV.EGFP virus. These neurons extend their axons through the microgrooves (length: 450 µm) into the axonal compartment within 11 days (green cells). (B and C) STED-images of cortical neurons stained for spectrin and tubulin. Note that spectrin is enriched in the growth cone while tubulin concentration is decreased. Scale bar 5 µm. (D) Quantification of line intensity scans along the axon, starting from the growth cone tip. Spectrin fluorescence intensity peaks at the growth cone, whereas the tubulin signal is lower compared to the axon. Data is represented as mean \pm SD. At least 5 biological replicates were analyzed, 11 axons were analyzed in total. (E) Representative STED image of a growth cone stained for spectrin and F-actin. Spectrin is enriched close to the actin filaments. Scale bar 5 µm. (F) Schematic drawing of a growth cone, showing 127 128 129 an enrichment of β -spectrin II in the apical central zone of the GC and microtubules splaying into the GC. (G) Representative STED image of a growth cone stained for F-actin and spectrin. Spectrin is enriched in the growth cone, in particular close to the base of filopodia (open arrowhead). G4 shows an enlarged version of G3; closed arrowheads point to enriched spectrin at 130 the base of filopodia. Scale bar 4 μ m in G1-3 and 1 μ m in G4.

131

132 Development of the membrane-associated periodic skeleton in native rat cortical neurons

133 Zhong et al. reported that the MPS of hippocampal neurons starts to develop near the soma and 134 matures along the axon towards the GC (Zhong et al., 2014). To better understand the process 135 of MPS assembly in a growing axon, we asked how the MPS develops in the youngest part of the axon, i.e., the segment adjacent to GC. We, therefore, analyzed the first 250 µm of the 136 137 axonal segment close to the GC (Figure 2, A). We divided the axon into 10 µm segments and 138 evaluated for each respective segment the total length of the segment showing a periodic 139 arrangement of spectrin with a distance of around 190 µm between spectrin signals. The 140 respective length of axon showing MPS per respective segment was then divided by 10 µm. As 141 depicted in Figure 2, B, there was a consistent, nearly 10-fold increase of axonal spectrin 142 periodicity from the axonal GC base up to 250 μ m towards the soma (first 10 μ m: 7.5% \pm 2.4%, 143 250 μ m from GC: 68.5% ± 16.5%). A periodicity of 50% (5/10 μ m of measured axon showed periodicity) was reached at a mean distance of 220 µm from the GC. With an average growth 144 145 speed of 18.3 µm/hour in our culturing conditions, this distance corresponds to a maturation time of 12 hours for this axonal segment. A fully developed MPS (10/10 µm measured axon 146 147 showed periodicity) was seen beyond distances of $250 - 350 \mu m$ from the GC. This distance 148 corresponds to a maturation time of approximately 14-19 hours until an axonal segment 149 contains a continuous MPS.

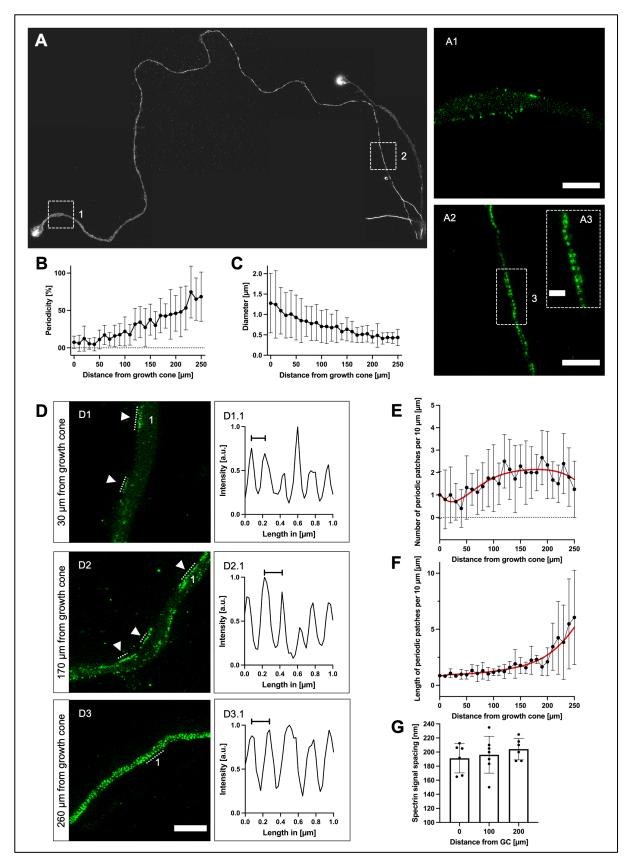




Figure 2. Development of the membrane-associated periodic skeleton along the growing axon. (A) Representative image of an axon with its growth cone, stained for spectrin, stitched from multiple STED images, up to 270 μ m towards soma. A1 shows an enlarged axon section at 10 μ m distance from the growth cone, no periodicity is visible (scale bar 2 μ m). A2 shows an enlarged section of A, 270 μ m from the growth cone (scale bar 2 μ m). A3 shows an enlarged section of A2, MPS is clearly visible (scale bar 0.5 μ m). (B) Graphical display of the development of periodicity over the length of the axon, starting from the growth cone base. Periodicity is increasing linearly towards the soma. Data is represented as mean \pm SD. At least 5 biological replicates were analyzed, 11 axons were analyzed in total. Because of limitations in following axons back for

158 159 160 extended lengths, not all axons could be included at distal points. (C) Reduction of the axonal diameter over the length of the axon, starting from the growth cone base, contrariwise to (B). Data is represented as mean \pm SD. At least 5 biological replicates were analyzed, 11 axons were analyzed in total. Because of limitations in following axons back for extended lengths, not all 161 162 163 axons could be included at distal points. (D) Three sections of an axon stained for spectrin, showing different stages of MPS development, were imaged using STED microscopy. D1 is 30 µm afar from the growth cone, D2 170 µm, and D3 260 µm. The arrowheads point to periodic patches, periodic arrangements of spectrin with at least three signals (scale bar 2 µm). D1.1 shows 164 165 the periodic plot of a line intensity scan along the numbered periodic patch in D1. The marked distance is 150 nm. D2.1 shows a periodic line intensity scan along the numbered periodic patch in D2. The marked distance is 200 nm. D3.1 shows the periodic 166 plot of a line intensity scan along the numbered periodic patch in D3. The marked distance is 200 nm. (E and F) Quantification 167 168 of parameters for the development of MPS along the axon, starting from the growth cone base. The number of periodic patches (E) increases, then decreases beyond 200 µm. The length of periodic patches (F) increases towards soma. Data is represented 169 170 as mean ± SD. Red lines indicate estimated curve fits. At least 5 biological replicates were analyzed, 10 axons were analyzed in total. Because of limitations in following axons back for extended lengths, not all axons could be included at distal points. 171 172 173 174 (G) Development of the spacing of spectrin tetramers in periodic patches in 10 μ m long axonal segments at a distance of 0 μ m, 100 µm, and 200 µm from the growth cone in native axons. Although a slight increase of spectrin tetramer spacing along the axon is visible, no significant changes were detected using ordinary one-way ANOVA, followed by Turkey's multiple comparisons test. Bars represent mean ± SD, at least 5 biological replicates were analyzed, 11 axons were analyzed in total.

175 A role for the MPS in influencing the axonal diameter has been described in previous

176 studies(Costa et al., 2020; Leite et al., 2016). We, therefore, analyzed the development of the

177 axonal diameter along the axon. As **Figure 2**, **C** shows, there was a nearly three-fold decrease

178 of axonal diameter along a distance of 250 μ m from the GC (first 10 μ m: 1.3 μ m \pm 0.2 μ m, 250

179 μ m from GC: 0.4 μ m ± 0.1 μ m). The finding that the youngest part of the axon shows the lowest

180 periodicity, but greatest diameter, supports the idea of a progressive constriction of the axonal

181 diameter by the MPS (Leite et al., 2016).

182 The MPS develops in periodic patches

183 We noticed that spectrin periodicity does not extend continuously from proximal to distal but 184 develops in the distal axon from multiple periodic spectrin patches extending and converging 185 towards the soma (Figure 2, D). A "periodic spectrin patch" was defined as a continuous array 186 of at least three spectrin-fluorescence signals inside the axon with a mean periodic distance of 187 150 – 250 nm. This distance was chosen based on the literature (Brown et al., 2015; Xu et al., 188 2013) and to take possible changes of spectrin tetramer spacing during maturation into account. 189 The number and length of these periodic spectrin patches were measured in 10 µm segments 190 along the growing axon starting at the base of the GC.

191 Short periodic spectrin patches appeared in the first 100 µm adjacent to the GC, with a density

192 of 1 patch per 10 μ m of axon (1.1 \pm 0.1). The density increased to 2 patches per 10 μ m between

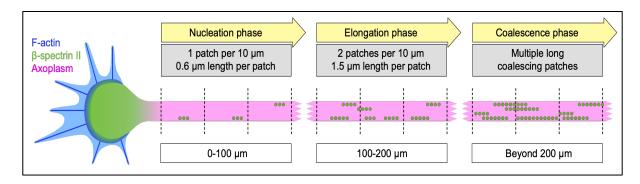
193 $100 - 200 \ \mu m$ from the GC (2.1 \pm 0.1) and then slowly decreased again at distances >200 μm

- 194 from the GC as single periodic spectrin patches extended and coalesced $(1.8 \pm 0.6 \text{ at } 250 \text{ }\mu\text{m})$
- 195 (**Figure 2, E**).

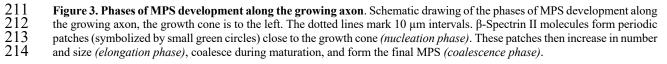
196 The mean length of the periodic spectrin patches in the first 100 μ m adjacent to the GC was 197 0.6 μ m (644 nm \pm 96.4 nm), corresponding to 3 consecutive spectrin tetramers. Between 198 100 – 200 μ m from the GC, the length of the periodic spectrin patches increased gradually

- 199 (1525 nm \pm 14.7 nm) and then exponentially beyond 200 μ m to a mean length of 5.5 μ m
- 200 (5476 nm \pm 179.6 nm) at 250 μ m, corresponding to around 27 spectrin tetramers (**Figure 2**, **F**).
- 201 The average spacing of spectrin tetramers along the axon slowly increased from around 190 nm
- at the GC-base to 200 nm towards the soma (10 μ m: 191.2 nm \pm 8.5 nm, 100 μ m: 196.2 nm \pm
- 203 9.9 nm, and 200 μm from the GC: 204.2 nm \pm 6.2 nm) (Figure 2, G).

Our data supports a model in which the MPS does not extend continuously into the distal outgrowing axon. Instead, the MPS assembles from multiple "MPS seeds" that grow and coalesce into a single scaffold. The formation of the MPS in an outgrowing axon can be divided into a nucleation phase with multilocular periodic spectrin patches developing along the axon, an elongation phase where single patches extend their size and the number of patches further increases, and a final coalescence phase where the MPS is closed (**Figure 3**).



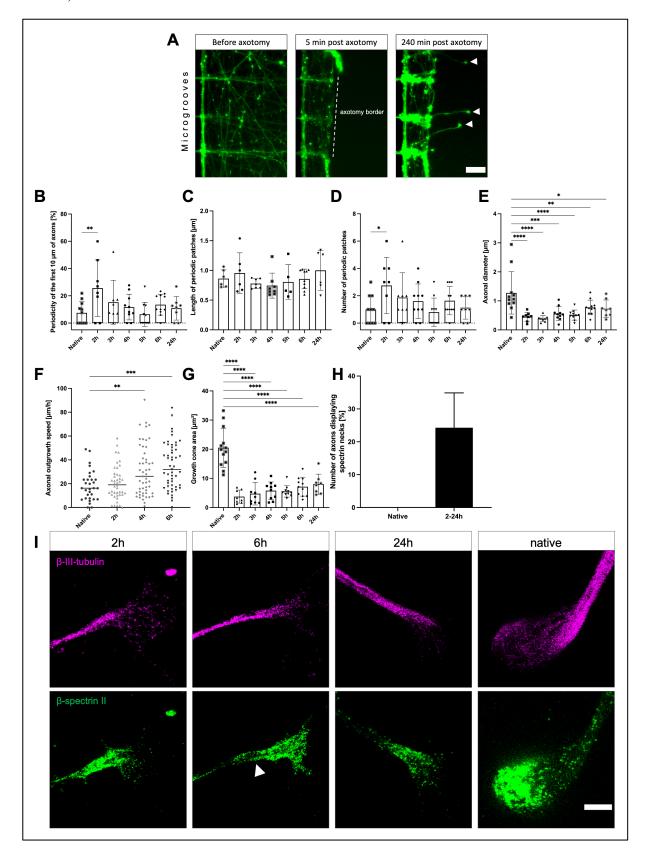
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215 Spectrin periodicity is increased early after regeneration

216 Our next set of experiments aimed to analyze the morphological and cytoskeletal changes in 217 regenerating axons following an axotomy. We performed axotomy by vacuum aspiration in 218 microfluidic chambers and fixed the neurons and regenerating axons at different time points (2, 219 3, 4, 5, 6, and 24 hours) after axotomy. We then analyzed different morphological features of 220 the GC and the axon and analyzed the development of the MPS in the axon. Because of the 221 different lengths of the regenerating axons at different time points, we compared periodicity 222 values only at the first 10 µm close to the GC at all time points. Each analyzed axon was traced 223 over time to confirm its axotomy and to reliably exclude the few native axons growing through 224 the microgrooves from the analysis (Figure 4, A; for further details, see methods section). 225 Regeneration was first noticed at 2 hours after axotomy. Axons extended in length over the 226 examined period.

- 227 Surprisingly, regenerating axons developed thiree MPS much faster than the native axons with
- 228 a significant, almost 3.5-fold higher spectrin periodicity in the first 10 µm at 2 hours after
- axotomy in regenerating axons compared to native axons (Figure 4, B) $(25.7\% \pm 7.3\% \text{ vs. } 7.5\%)$
- 230 ± 2.4%).



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 $\begin{array}{r} 232\\ 233\\ 235\\ 236\\ 237\\ 238\\ 240\\ 242\\ 243\\ 2445\\ 246\\ 247\\ 248\\ 250\\ 251\\ 252\\ 253\\ 255\\ 257\\ 255\\ 257\\ \end{array}$ Figure 4. Comparison of growth cone morphology and β-spectrin II periodicity of the first 10 μm of axon proximal to the growth cone in native and regenerating neurons. (A) Graphical explanation of the method used for marking regenerating axons. Microscopic images of the same section of the axonal compartment were acquired before axotomy, five minutes after, and 2-24 hours after axotomy. For detailed information, see the methods section. White arrowheads indicate regenerating axons. Scale bar is 30 µm. (B) Comparison of periodicity in the first 10 µm of the axon, starting from the growth cone base. Data of different time points after axotomy and native axons are shown. The 2-hour time point shows a significant increase in periodicity. Bars represent mean ± SD. One-way ANOVA, followed by Dunnett's multiple comparisons test, was performed. At least 8 axons were analyzed per condition, n=3, 67 axons were analyzed in total. (C and D) Comparison of the average length and number of periodic patches in the first 10 µm of axon adjacent to the growth cone. Data of different time points after axotomy and native axons are shown. No significant differences were detected for periodic patch length (C). The average number of periodic patches was significantly increased at the 2-hour time point, compared to native axons (D). One-way ANOVA was performed, followed by Dunnett's multiple comparisons test. Bars represent mean \pm SD. At least 8 axons were analyzed per condition, n=3, 67 axons were analyzed in total. (E) Comparison of axonal diameter in the first 10 µm of the axon, starting from the growth cone base. Data of different time points after axotomy and native axons are shown. Reduction of the axonal diameter of the different time points after axotomy is significant, compared to native axons, whereas only the 2hour time point shows a significant increase in periodicity (**B**). Bars represent mean \pm SD, one-way ANOVA, followed by Dunnett's multiple comparisons test was performed. At least 8 axons were analyzed per condition, n=3, 67 axons were analyzed in total. (F) Comparison of axonal outgrowth speed in cortical neurons fixed at DIV 10. The 4-hour and 6-hour time points showed significantly increased outgrowth speed compared to native axons. The line represents mean, one-way ANOVA, followed by Dunnett's multiple comparisons test was performed. At least 31 Axons were analyzed per condition, n=3, 177 axons were analyzed in total. (G), Comparison of growth cone area of different time points after axotomy and native axons, quantified in spectrin-stained axons. The growth cone area of regenerating axons is significantly reduced compared to native axons. Bars represent mean ± SD; one-way ANOVA, followed by Dunnett's multiple comparisons test, was performed. At least 8 axons were analyzed per condition, n=3, 67 axons were analyzed in total. (H) Comparison of regenerating axons of combined time points and native axons displaying spectrin necks. Native axons did not display spectrin necks, whereas regenerating axons showed variating percentages of spectrin necks. A spectrin neck was defined as a visible increase of spectrin $\overline{258}$ signal at the axonal segment directly connected to the growth cone, not longer than 3 µm. At least 8 axons were analyzed per 259 condition, n=3, 67 axons were analyzed in total. (I) Representative STED images of growth cone sizes at different time points 260 261 after axotomy and native. Notice the size difference of native and regenerating axons. White arrowhead points to spectrin neck. Scale bar is $2 \mu m$.

As Figure 4, C shows, the length of periodic patches was not significantly increased in regenerating axons, compared to native axons ($0.96 \ \mu m \pm 0.1 \ \mu m vs. 0.86 \ \mu m \pm 0.06 \ \mu m$). However, the number of periodic patches was increased almost three-fold at the 2-hour time point, compared to native axons ($2.7 \pm 0.7 vs. 1.0 \pm 0.3$; Figure 4, D).

We thus conclude that after axotomy, the axons transiently form more periodic spectrin patches in the outgrowing part of the axon resulting in a higher degree of periodicity and a quicker nucleation phase compared to native axons. Interestingly, at later time points after axotomy, the lesioned axons then resembled more and more the characteristics of the native axons with regards to their periodicity and number of periodic patches.

The axonal diameter of the segment adjacent to the GC was decreased in regenerating axons compared to native axons (**Figure 4**, **E**). At 2 hours after axotomy, the axonal diameter was

- reduced almost three-fold (2 hours: 0.5 μ m ± 0.1 μ m vs. native: 1.3 μ m ± 0.2 μ m). The axonal
- 274 diameter increased in the following hours but was still significantly reduced almost two-fold
- up to 24 hours after the axotomy (24 hours: $0.7 \ \mu m \pm 0.1 \ \mu m$).
- 276 To test whether the outgrowth speed of regenerating axons was faster or not, we performed
- time-lapse microscopy and measured the outgrowth velocity of the GC in the time-lapse video.
- As Figure 4, F shows, we saw a significant increase of outgrowth velocity 4 and 6 hours after
- axotomy but not 2 hours after axotomy (native: 18.3 μ m/h ± 2.3 μ m/h, 2 hours: 20.5 μ m/h ±

280 2.1 μ m/h, 4 hours: 30.8 μ m/h ± 3.0 μ m/h, and 6 hours after axotomy: 34.5 μ m/h ± 2.7 μ m/h). 281 Thus, the outgrowth velocity is not correlated to the higher spectrin periodicity.

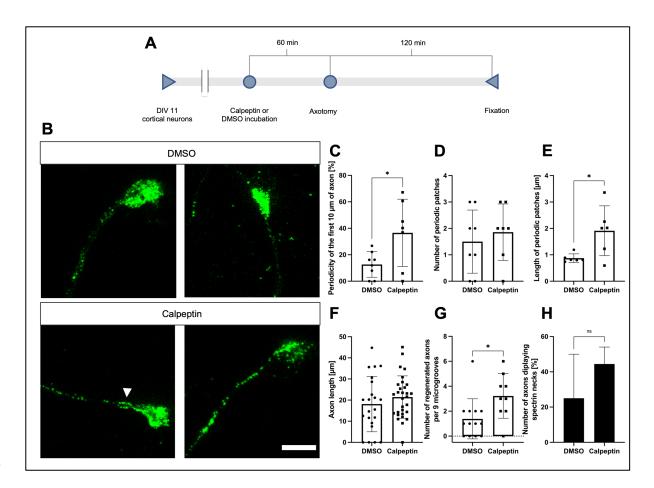
To analyze whether the newly formed GC after axotomy differed morphologically from the native one, we measured the area of native and regenerating GCs of different time points. As **Figure 4, G** shows, the area of regenerating GCs at 2 hours after axotomy was reduced fivefold, compared to native axons. The GC area increased by two times over 24 hours but did not reach native size in this timespan (2 hours: $3.7 \ \mu m^2 \pm 0.8 \ \mu m^2$, 24 hours: $8.0 \ \mu m^2 \pm 1.2 \ \mu m^2$, and native axons: $20.5 \ \mu m^2 \pm 1.9 \ \mu m^2$). We observed a similar reduction regarding the length and width of the GCs; for detailed information, see **Supplemental Figure 2**.

Some axons showed an enrichment of spectrin close to the GC, which we will refer to as "spectrin neck". A spectrin neck was defined as a visible increase of spectrin signal at the axonal segment directly connected to the GC, not longer than 3 micrometers. A fraction of 24,3% of regenerating axons showed spectrin necks compared to none of the native axons (**Figure 4, H** and **I**).

To sum up the results of the axotomy experiments, regenerating axons compared to native axons exhibit a transiently higher degree of spectrin periodicity based on an increased periodic spectrin patch density, a persistent smaller axon diameter, and a reduced growth cone size. Outgrowth velocity is increased at 4 and 6 hours after axotomy in the regenerating axons. A substantial number of regenerating axons show an accumulation of spectrin at the base of the GC.

300 Inhibition of calpain increased periodicity and the number of regenerating axons shortly 301 after axotomy

Since spectrin periodicity was increased early in regenerating axons, we wondered how a more stabilized spectrin cytoskeleton might influence the growth behavior of axons. To stabilize spectrin and putatively further increase periodicity after axotomy, we treated the neurons with the calpain inhibitor calpeptin (10 μ M dissolved in dimethylsulfoxide (DMSO) or DMSO alone as control), performed an axotomy, and analyzed the axons 2 hours after axotomy (**Figure 5, A** and **B**). Calpain is a spectrin cleavage enzyme that is activated early after axotomy leading to a destruction of spectrin structures (Zhang et al., 2016).



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310 311 312 313 314 315 316 317 318 319 320 321 322 Figure 5. Effects of calpeptin treatment on MPS stability and regeneration of axons after axotomy. (A) Schematic display of used test setup. 10 µM of calpeptin or equal amounts of DMSO were used for incubation. STED imaging was performed following fixation. (B) Representative STED images of cortical neurons treated with calpeptin or DMSO before axotomy, fixation after 2 hours, scale bar 2 µm. The arrowhead points to a spectrin neck at the base of the growth cone. (C-E) Quantification of parameters for MPS development along the axon in regenerating axons treated with calpeptin, compared to regenerating DMSO controls. Analyzed were the first 10 µm of axon close to the growth cone. Periodicity (C) and size of periodic patches (E) were increased in calpeptin-treated axons. Bars represent mean \pm SD. Unpaired t test (C and D) or Kolmogorov-Smirnov test (E) was performed. At least 7 axons were analyzed per condition, n=3, 15 axons were analyzed in total. (F and G) Graphical display of axon length of regenerating neurons 2 hours after axotomy. (F) shows no significant increase in axon length, while (G) shows a significant increase in regenerating axons per 9 microgrooves. Bars represent mean ± SD. Non-parametric Mann-Whitney U test was performed. At least 22 axons were analyzed per condition, n=3, 52 axons were analyzed in total. (G) at least 27 microgrooves were analyzed per condition, n=3, 198 microgrooves were analyzed in total. (H) Quantification of axons, treated with DMSO or calpeptin, displaying spectrin necks. Calpeptin-treated axons $\overline{3}\overline{2}\overline{3}$ displayed a higher percentage of growth cones with spectrin necks. A spectrin neck was defined as a clearly visible increase of 324 325 spectrin signal at the axonal segment directly connected to the growth cone, not longer than 3 µm. At least 7 axons were analyzed per condition, n=3, 15 axons were analyzed in total.

326 The calpeptin treatment indeed led to a more than 3-fold increase in spectrin periodicity at 2

- hours after axotomy in the treated axons compared to control $(36.6\% \pm 9.6\% \text{ vs. } 12.7\% \pm 3.4\%)$
- 328 (Figure 5, C). Interestingly, the axonal diameter did not differ between both groups
- 329 (Supplemental Figure 3).
- 330 The number of periodic spectrin patches was not significantly changed in calpeptin-treated
- neurons compared to control $(1.9 \pm 0.4 \text{ vs. } 1.5 \pm 0.4)$ (Figure 5, D). The length of the periodic
- 332 spectrin patches, however, was increased by 2.5-times in calpeptin treated neurons, compared
- to DMSO treated neurons (1.6 μ m ± 0.4 μ m vs. 0.7 μ m ± 0.2 μ m) (Figure 5, E). The number

334 of periodic patches was decreased in both conditions compared to the previous experiment.

- This inconsistency could be due to toxic effects of the DMSO.
- 336 The length of the regenerating segment of the axons was not significantly different between
- calpeptin or DMSO treated neurons (21.4 μ m ± 1.8 μ m vs. 18.1 μ m vs. 2.8 μ m) (Figure 5, F).
- 338 Interestingly, the number of regenerating axons per 9 microgrooves (9 microgrooves are in the
- 339 viewfinder at 20x magnification) was significantly higher in analyzed images of calpeptin-
- treated chambers, compared to DMSO control group $(3.2 \pm 0.6 \text{ vs. } 1.4 \pm 0.4)$ (Figure 5, G).
- 341 Also, axons treated with calpeptin showed spectrin necks more often than in the control group
- 342 (44.4% vs. 25%), although the difference was not significant (Figure 5, H).
- 343 Inhibition of spectrin cleavage thus significantly enhances the elongation phase of MPS
- 344 formation, leading to increased spectrin periodicity without influencing the nucleation phase.
- 345 This does not increase outgrowth velocity but enables more axons to regenerate within the
- analyzed time period.

347 **Discussion**

In this study, we analyzed the formation and development of the MPS in the outgrowing axon. We report that the developing MPS is organized in periodic patches that grow in size and number and coalesce over time during its maturation. We provide further evidence that the MPS develops in a proximal to distal pattern. We also show that regenerating axons after axotomy acquire periodicity faster over a transient timespan.

- We found an enrichment of spectrin inside the axonal GC in line with previous studies (Galiano et al., 2012; Levine & Willard, 1981; Matsuoka, Li, & Bennett, 2000; Tian et al., 2012). This enrichment was present in native GCs and regenerated GCs at different time points after axotomy. Our data show that within the GC, spectrin is often localized in close proximity to the base of filopodia. Actin filaments could therefore be anchored or stabilized in the spectrin cytoskeleton of the GC. Elevated spectrin levels in the GC could also resemble a pool of spectrin tetramers that are later incorporated into the growing axon.
- 360 The MPS has been found along the axon, in dendrites, the AIS, and the neck of dendritic spines 361 (Unsain, Stefani, & Cáceres, 2018). In outgrowing axons, the MPS has been suggested to 362 develop in a proximal to distal pattern (Han et al., 2017; Lorenzo et al., 2019; Zhong et al., 363 2014). We can specify that growing axons display a characteristic spatial and temporal increase 364 of spectrin periodicity in a gradient from the distal growth cone towards the proximal axonal 365 regions. We show that the MPS does not extend in a continuous pattern along the axon. Instead, 366 multiple periodic patches are nucleated along the growing axon. This nucleation happens close 367 to the GC, at a frequency of about one patch per 10 μ m. These periodic patches double in 368 number over a timespan of about 11 hours, then increase in size and conflate as maturation 369 continues to form the mature MPS finally. This maturation happens gradually as the axon 370 grows, resulting in a gradient from immature MPS at the GC base towards mature MPS in the 371 mid-axon.

372 We suggest the GC or its neck as a possible nucleation site for three reasons: First, the GC sits 373 on top of the growing axon where spectrin nucleation naturally starts. Second, our data shows 374 periodic patches in the axonal segment adjacent to the GC. Third, spectrin is highly enriched 375 within the GC. It could therefore function as a pool for the implementation of spectrin into the 376 growing axon. Interestingly, the number of periodic patches also increases at distances of 377 $100 - 200 \,\mu\text{m}$ from the GC. This increase could be explained by nucleation molecules that stay 378 on the way as the axon extends after integration into the axon at the GC. Zhong et al. 379 demonstrated that formation of the MPS in dendrites is dependent on the local concentration of 380 spectrin and the presence of ankyrin B (Zhong et al., 2014). Surprisingly, we did not detect 381 periodic patches or periodicity inside the GC, despite higher concentrations of spectrin and 382 ankyrin B (Galiano et al., 2012). This points to other nucleation factors that might not be present 383 in the GC but in the axonal segment close to the GC. Our findings, therefore, shine a new light 384 on the role of the GC and its neck in the formation of the axonal MPS.

- 385 A role of the MPS in maintaining axonal diameter has been described in previous studies (Costa 386 et al., 2020; Leite et al., 2016). We show here that the axonal diameter decreases while the MPS 387 is formed. The finding that the youngest part of the axon shows the lowest periodicity, but 388 greatest diameter, supports the idea of a progressive constriction of the axonal diameter by the 389 MPS (Leite et al., 2016). It has also been shown that the diameter of actin rings decreased over 390 time in culture (Leite et al., 2016). Furthermore, knockout of a-adducin, also distributed 391 periodically along the axon, increased the axonal diameter, without changes in the spacing of 392 actin rings (Leite et al., 2016). Previous studies found recruitment of adducin into the MPS after 393 DIV 8 (Zhong et al., 2014). It is also well known that non-muscle myosin II (NMII) has a role 394 in regulating axonal diameter (Fan, Tofangchi, Kandel, Popescu, & Saif, 2017) and is 395 distributed throughout the MPS (Costa et al., 2020; Vassilopoulos et al., 2019; R. Zhou et al., 396 2020). We found that stabilization of the MPS using calpeptin did not affect the axonal diameter 397 in young, regenerating axons. This suggests that the role of both adducin and NMII in axonal 398 diameter regulation is not directly linked to spectrin periodicity abundance and that spectrin 399 plays an inferior role in diameter regulation. Interestingly, degeneration of the axon and its MPS 400 through trophic factor withdrawal did also not lead to an increase in axonal diameter (Unsain, 401 Bordenave, et al., 2018).
- The role of the MPS in axonal degeneration has been studied previously (Unsain, Bordenave, et al., 2018; Wang et al., 2019). However, the MPS has not been studied in regenerating axons so far. Neuronal regeneration is an important process that needs to be activated in many neurological conditions in order to compensate for the disease-associated damage of nerve tissue, e.g., in spinal cord injury, stroke, or neurodegenerative diseases (Mahar & Cavalli, 2018).
- We show here that axon morphology and the formation of the MPS in regenerating axons differ significantly from native axon outgrowth. Most strikingly, the axon diameter and growth cone size were significantly reduced in regenerating axons compared to native axons at all examined time points within 24h after axotomy. This reduction correlated with an increased outgrowth speed of the regenerating axons at 4 and 6 hours after axotomy.

413 At 2 hours after axotomy, spectrin periodicity and the number of spectrin patches were 414 significantly increased in the newly formed axon but reached "normal" values at later time 415 points. This suggests that the nucleation phase and most likely also the coalescence phase of 416 MPS formation are enhanced early after axotomy. Since we accurately checked that all 417 analyzed axons had been axotomized and evaluated only the newly formed part of the axons, 418 we are confident that the observed alterations of spectrin MPS formation are characteristic of 419 the regenerating axon and not confounded by residual proximal parts of the axotomized axon. 420 A plausible explanation for the observations is that the molecular machinery necessary for 421 nucleation and coalescence of spectrin patches is already in place in the more proximal part of 422 the axon that remains after axotomy and can thus more quickly be employed for MPS formation. 423 The decrease (i.e., normalization) of the number of periodic patches at later time points could 424 be due to the increased outgrowth speed in the following hours, which would dilute a putatively 425 still increased number of patches.

Also, spectrin itself is more abundant in the distal regenerating axon as compared to native axons. In around 25% of all regenerating axons, we saw an accumulation of spectrin adjacent to the GC, which we termed spectrin neck. Spectrin necks were never observed in axons without axotomy and thus specific for regenerating axons. The underlying mechanism for spectrin neck development might be increased local translation of spectrin RNA (Yoon, Zivraj, & Holt, 2009), altered axonal transport (Lorenzo et al., 2019), or local cleavage with increased spectrin reutilization (Wang et al., 2019).

433 Although hypothesized in a review by Leite and Sousa (2016), our finding of increased spectrin 434 levels and periodicity in regenerating axons is particularly surprising, considering the 435 degradation of the MPS in degenerating axons following axotomy or NGF-withdrawal, 436 presented in previous studies (Unsain, Bordenave, et al., 2018; Wang et al., 2019). A possible 437 explanation could be increased levels of free spectrin molecules after the axotomy. It has been 438 shown that overexpression of spectrin increases periodicity in dendrites (Zhong et al., 2014). 439 Since only a fraction of the axotomized axons regenerates while another fraction degenerates 440 and only the regenerating axons were selected in our analysis, it can also be speculated that the 441 observed features of the cytoskeleton are structural prerequisites that enable lesioned axons to 442 regenerate after a lesion quickly.

443 It is well known that axotomy leads to a rapid calcium influx into the axoplasm and activation 444 of the calcium-dependent protease calpain that induces the consecutive destruction of the 445 axonal cytoskeleton (George, Glass, & Griffin, 1995; Knöferle et al., 2010). Inhibition of 446 calpain, e.g., via calpeptin, has been shown to inhibit axonal degeneration after lesion, at least 447 within the first hours after axotomy (George et al., 1995; Zhang et al., 2016). We thus analyzed 448 the effects of calpeptin treatment on the spectrin cytoskeleton on the nanoscale. Calpeptin 449 treatment indeed led to a significant increase of spectrin periodicity in regenerating axons. 450 Interestingly, upon calpeptin treatment, the size of the periodic patches was increased, but not 451 their number. Thus, calpain inhibition does not interfere with spectrin nucleation but enhances 452 the elongation of periodic spectrin patches. The most likely explanation for this finding is a 453 decreased degradation of spectrin molecules in the distal axon caused by calpain inhibition.

The increased and more rapidly evolving spectrin periodicity did not translate into an increased length of the regenerating axons. This is in line with previous studies that reported adverse effects of calpeptin on growth cone stability and axonal regeneration (Gitler & Spira, 1998). It is proposed that a constant turnover of the cytoskeletal molecules is essential for the highly dynamic growth cone and axonal elongation. However, we found an increased number of regenerating axons in the calpeptin-treated neurons. Thus, enhanced MPS-formation might set a basis for more axonal regeneration.

461 Taken together, we demonstrate here that the MPS is formed in the region 250 µm proximal 462 from the GC in three phases: nucleation of periodic patches adjacent to the GC, subsequent 463 elongation of periodic patches, and finally coalescence of periodic patches into a mature MPS. 464 After a lesion, the outgrowth of regenerating axons is characterized by a greater speed, thinner 465 axon diameter, smaller growth cones, and at least transiently an increased number of periodic 466 patches and higher periodicity. Inhibition of the spectrin-severing protease calpain stabilized 467 the MPS and enhanced the elongation phase. Future research should explore the underlying 468 molecular mechanisms and test different substances that alter MPS dynamics.

469 Materials and Methods

470 Cell culture, viral transduction, and axotomy

The use of E18 Wistar rat embryos for isolation of cortical tissue was conducted according to the approved experimental animal licenses (33.9-42502-04-11/0408) issued by the responsible animal welfare authority (Niedersächsisches Landesamt für Verbraucherschutz und Lebensmittelsicherheit) and controlled by the local animal welfare committee of the University Medical Center Göttingen, Germany.

476 Primary cortical neurons were isolated from E18 Wistar rat embryos as described before (Sahu, 477 Nikkilä, Lågas, Kolehmainen, & Castrén, 2019). In short, the dissected tissue was incubated 478 with 1 ml trypsin (Sigma Aldrich, St. Louis, MO, USA; #T9935, 25.000 U/ml) for 15 min at 479 37 °C. Approximately 1 minute before the trypsinization was complete, 50 µl of DNase I 480 (Roche, Basel, Swiss; #11284932001, 2000 U/mg) were added. The suspension was 481 centrifuged shortly, and the supernatant was removed. Pellet was then resuspended in 1 ml fetal 482 bovine serum (FBS; Biochrom, Berlin, Germany) and carefully triturated. If tissue clumps were 483 still present, more DNase was supplemented. The supernatant was then transferred into a 15 ml 484 falcon tube, and the remaining tissue was again triturated in 1 ml of media. The supernatant 485 was transferred again, and the falcon tube was centrifuged. Neurons were resuspended in 500 486 µl Neurobasal medium (Gibco, Waltham, MA, USA), supplemented with 10% of FBS, 2% of 487 50x B27 Supplement (Gibco), 1% Penicillin-Streptomycin-Neomycin (Gibco), 0,5% holo-488 Transferrin human (Applichem, Darmstadt, Germany), and 0.25% L-glutamax (Gibco). 489 Neurons were seeded at a density of 60,000 cells per chamber into the somatic compartment of 490 a microfluidic chamber (MFC) (XONA microfluidics, Research Triangle Park, NC, USA). 491 MFCs were mounted onto 35 mm FluoroDishes (World Precision Instruments, Inc, Sarasota, 492 FL, USA), precoated with 0.1 mg/ml poly-L-ornithine (Sigma Aldrich, P3655) in borate buffer 493 solution (Sigma Aldrich, P8638), and 0,1% laminin (Sigma Aldrich, L2020) as described 494 previously (Zhang et al., 2016).

All cell cultures used for STED microscopy were transduced with an AAV 1/2, expressing EGFP (GenBank HQ416702.1), kindly gifted by Uwe Michel, Göttingen, as described before (Koch, Barski, Lingor, Bähr, & Michel, 2011). The stock solution had a titer of 5.6 x 10⁷ transduction units (TU). Expression was driven by a human synapsin-1 promoter and a human h1 promoter. The virus was employed with a titer of 0.5×10^7 TU per chamber and added to the medium on DIV 4. Usually, a transduction rate of > 90% was achieved, with no significant toxicity. The medium was augmented from DIV 4 – 11 with 10 ng/µl of ciliary neurotrophic

factor (CNTF, Pepro Tech, Rocky Hill, NJ, USA) and brain-derived neurotrophic factor (BDNF, Pepro-Tech). 10% of FBS was added to the medium in the first week but was omitted after DIV 6 to prevent the overpopulation of astrocytes. Cultured neurons were maintained with frequent medium exchange (every 2 to 3 days) and kept at 37 °C and 5% CO₂. Cells looked healthy, and no signs of apoptosis were observed until DIV 11 following this protocol.

507 Axotomy was performed on DIV 11 by vacuum aspiration of the media in the axonal 508 compartment as described elsewhere (Park et al., 2006). In short, the medium was aspirated 509 using a cell culture pump (Schütt Labortechnik, Göttingen, Germany) until an air bubble passed 510 through the axonal compartment, dissecting the axons. This procedure was repeated 2-4 times, 511 and successful axotomy was assessed using a standard cell culture microscope (Zeiss, Jena, 512 Germany). For treatment of neurons with calpeptin (Sigma Aldrich, #C8999), axons were 513 incubated with 10 µM of calpeptin in DMSO (Applichem) or equal amounts of DMSO for 514 1 hour prior to the axotomy, administered only into the axonal compartment.

515 Immunocytochemistry

516 All neuronal cultures were fixed on DIV 11. Axotomized neurons were fixed either 2, 3, 4, 5,

517 6, or 24 hours post-axotomy. Cells were washed with PBS (Applichem) and fixed with pure

518 methanol (Applichem) for 10 minutes at -20°C. Afterward, fixed cells were washed three times

519 with PBS and stored at 4°C in PBS if not immediately processed. Neuronal cultures used for

520 staining actin were fixed for 1 hour using 4% paraformaldehyde (PFA; Applichem) in PHEM

521 buffer, with 0.5% glutaraldehyde (Applichem), adapted from (Dent & Kalil, 2001).

For immunocytochemistry (ICC), MFCs were washed with PBS. If neurons were fixed with 522 523 PFA, quenching with glycine (Applichem) and ammonium chloride (Applichem; both 100 mM 524 in PBS) was performed to eliminate free aldehyde groups. After quenching (in case of PFA 525 fixation) or initial washing (upon methanol fixation), cells were permeabilized with Triton 526 X100 (0.3% in PBS; Applichem) for 5 min at room temperature (RT), MFCs were then 527 incubated in blocking solution containing 2% bovine serum albumin (BSA) in PBS for 20 min 528 at RT (Jackson Immuno Research, West Grove, PA, USA, #001-000-162). The blocking 529 solution was removed, and primary antibodies were added for 1 hour at RT.

After the primary antibody reaction, suitable secondary antibodies or phalloidin STAR RED were added. Incubation time was set to 45 minutes and was performed at 37°C. If not immediately imaged, stained neurons were stored light-protected in PBS at 4°C.

533 Monoclonal primary antibodies directed against β -spectrin II were obtained from BD (BD 534 Bioscience, Franklin Lakes, NJ, USA; mouse, #612563, 1:200), and rabbit β -III-tubulin 535 antibodies were from Cell Signaling (Cambridge, UK; #5568S, 1:100). Polyclonal secondary 536 antibodies goat anti-mouse Alexa594 were obtained from Invitrogen (Waltham, MA, USA; #A-537 11005, 1:100), and goat anti-rabbit STAR635p was from Abberior (#2-0012-007-2, 1:100, 538 Göttingen, Germany) as was Phalloidin-STAR RED (#2-0205-011-7, 1:100). Calpeptin was 539 obtained from Sigma (#C8999, 10 μ M).

540 Identification of regenerated axons

541 To identify and mark regenerated axons for STED microscopy, fluorescence live microscopy 542 of neuronal cultures was performed. Cultures were imaged using an inverted fluorescence 543 microscope (Axio Observer.Z1, Zeiss), equipped with an incubation chamber (Incubator PM 544 2000 RBT, PeCon, Erbach an der Donau, Germany), set to 37°C and 5% CO₂. MFCs were 545 placed inside the incubation chamber using a custom-made, 3D-printed template to seal the chamber against loss of CO₂. Images were acquired at 20x magnification before axotomy, 546 547 directly after, and at respective time points during regeneration (2, 3, 4, 5, 6, and 24 hours after 548 axotomy). Using the "multiview" function of the Zen 2.5 software (blue edition, Zeiss), images 549 were arranged side by side and aligned. After confirmation of a sufficient axotomy, the 550 regenerating GCs that were not present before or shortly after the axotomy were marked and 551 sorted out. Segments of the MFCs, showing neuronal somata on the axonal side were excluded. 552 The microgrooves were numbered, and STED microscopy was performed only with axons 553 previously identified as regenerating axons.

554 Analysis of axonal outgrowth speed

Axonal outgrowth speed was assessed using time-lapse microscopy. In short, cells were cultured as described above. Several 10-minute-long time-lapse videos of the axonal compartment were captured before axotomy and 2, 4, and 6 hours after axotomy, at 40x magnification, using the setup described above. The position of the GC was marked in the first and last image of the video, and the distance traveled by the GC was then measured using the open-source freeware Fiji (Schindelin et al., 2009).

561 Stimulated Emission Depletion (STED) microscopy

562 STED microscopy was performed to analyze MPS development in regenerating and native 563 axons. At least two regenerated axons were recorded per condition and chamber, starting at the 564 GC moving retrogradely towards the axotomy border. At 24 hours post-axotomy, axons were

565 too long to be followed back to the axotomy border. Therefore, at least three images were 566 acquired along the axon, starting from the GC. Native axons were imaged likewise. Two-color 567 STED microscopy was performed using a quad scanning STED microscope (Abberior 568 Instruments, Göttingen, Germany) equipped with an UPlanSApo 100x/1,40 Oil objective 569 (Olympus, Tokyo, Japan). An IX83 Olympus microscope with 4-color LED illumination source 570 and a monochrome widefield camera with a ¹/₂" CCD chip and 1280x960 pixels were used as a 571 platform. Pixel size was set to 20 - 25 nm with a pinhole size of 0.7 airy units. Dwell times 572 were set to 10 µs. Lasers with wavelengths of 561 nm and 640 nm were used for excitation of 573 Alexa 594 or STAR 635p / STAR RED, respectively. Depletion was performed using a 775 nm 574 pulsed STED laser. Images were acquired using the Imspector software (Abberior, Göttingen, 575 Germany) (Schönle, 2006).

576 Analysis of STED images

577 The brightness and contrast of the obtained STED images were adjusted using a custom-made 578 macro and Fiji, mimicking the "auto-function" of the brightness and contrast tool, and 579 converted into an 8-bit tagged image file format. If necessary, images of axon segments were 580 stitched manually with MosaicJ (Thévenaz & Unser, 2007) upon deactivated smart color 581 conversion, rotation, and blending. Images of the different cytoskeletal components were 582 analyzed independently using Fiji. As for the evaluation of GC morphology (length, width, and 583 area), the margin was outlined by applying the "polygon selection" tool. The length was 584 determined by drawing a straight line from the GC base to the farthest point of the previously 585 constructed polygon. The width was set to the longest straight within the polygon, being 586 orthogonal to the length-line. A spline-fitted segmented line, with a thickness covering the 587 axons width at the narrowest part, was drawn from the tip of the GC along the axon towards 588 the axotomy border to plot intensity curves. To reduce data size and avoid outliers, we 589 summarized the measurements for every 1 µm and normalized each dataset to its maximum. 590 The axon itself was further evaluated regarding diameter and β -spectrin II periodicity in 10 μ m 591 segments, using the self-programmed "Segmenter" macro for the segmentation of nonlinear 592 structures. The diameter was measured along the axon, starting at the GC neck, for every 10 µm. 593 We calculated β -spectrin II periodicity by measuring every periodic patch in a segment and 594 dividing the total length by 10µm. The number of periodic patches per segment and the average 595 patch size were also assessed. One periodic patch was defined as at least three individual β-596 spectrin II-signals inside the axon with a mean distance of 150 - 250 nm between spectrin 597 signals. This was assessed by performing intensity line scans along periodic patches,

598 confirming the periodic distance of intensity peaks. Primary axons were analyzed likewise, but 599 due to the absence of an axotomy border, the first 250 μ m of the axon, starting at the GC base, 600 were considered. To analyze the average spacing of spectrin tetramers, line intensity scans were 601 performed in spectrin stains along periodic patches at distances of 0, 100, and 200 μ m from the 602 GC in all native axons. The distances between periodic signal peaks of spectrin were then 603 measured. In co-stainings of actin and spectrin (**Figure 1, e** and **g**), actin is pseudo-colored 604 using the CET-L15 colormap by Peter Kovesi (Kovesi, 2015) (colorcet.com).

605 Statistics

606 Statistical evaluation was performed using GraphPad PRISM 9.1 (GraphPad Software, Inc, San

607 Diego, CA, USA). Two groups were compared using unpaired Student's t test (two-tailed).

608 Non-parametric Mann-Whitney U or Kolmogorov-Smirnov tests were used if the assumptions

609 for the t test were not met. Ordinary one-way ANOVA was conducted to compare two or more

610 groups, followed by Dunnett's multiple comparisons test. In general, mean values \pm SEM are

611 described unless otherwise noted, and at least three independent experiments were analyzed.

612 Significant differences between compared groups are exhibited as follows: * p < 0.05, ** p <

 $613 \qquad 0.01, \, {}^{***} \, p < 0.001, \, {}^{****} \, p < 0.0001.$

614

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619 **Competing Interests**

620 All authors declare that they do not have any competing interests.

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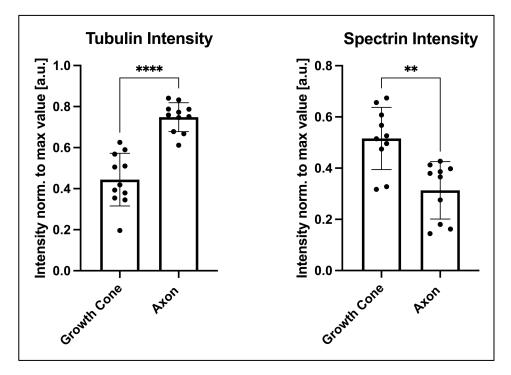
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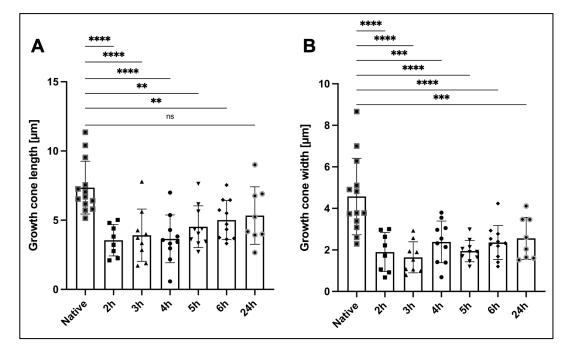
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757 Supplemental Figures

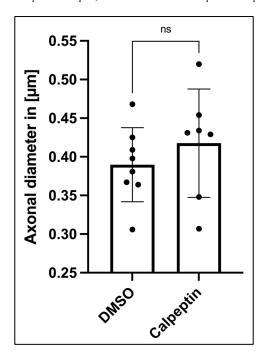


 Supplemental Figure 1. Distribution of β -spectrin II and β -III-tubulin along the axon and the growth cone in native rat cortical neurons. Quantification of 50 µm long line intensity scans along the axon, starting from the growth cone base. Due to the mean growth cone length, the first 7 µm of the line intensities represent the GCs, whereas the other 7 – 50 µm represent the axons. β -Spectrin II-fluorescence intensity peaks at the growth cone, whereas the β -III-tubulin signal is lower compared to the axon. Data is represented as mean ± SD. At least 4 biological replicates were analyzed, 11 axons were analyzed in total.



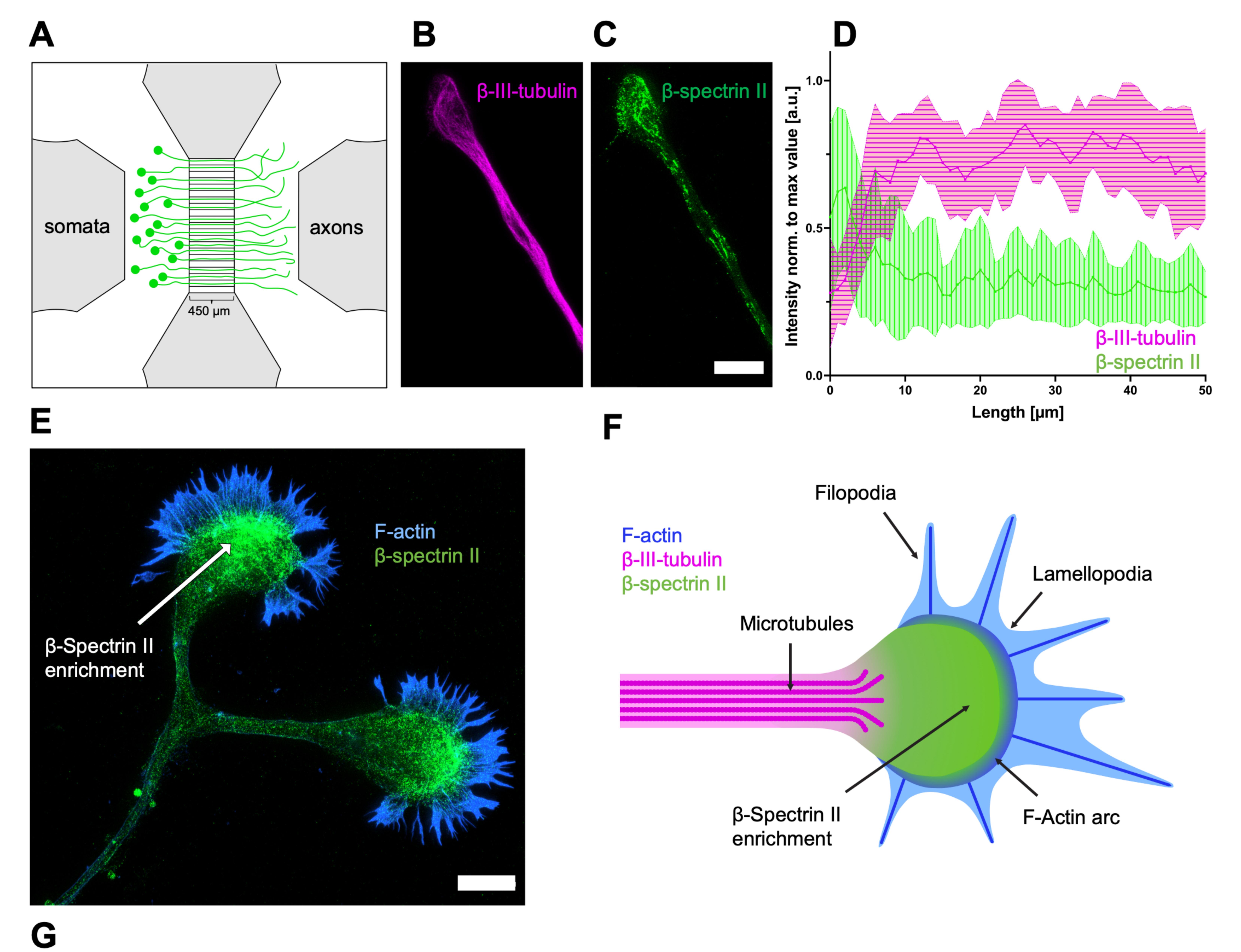
766Supplemental Figure 2. Difference of GC length and width of native and regenerating axons. Comparison of growth767cone length (A), and growth cone width (B) of different time points after axotomy and native axons, quantified in spectrin-768stained axons. The growth cone length and width of regenerating axons are significantly reduced compared to native axons,769except for the growth cone width of the 24-hour time point. Bars represent mean \pm SD; one-way ANOVA, followed by770Dunnett's multiple comparisons test, was performed. At least 8 axons were analyzed per condition, n=3, 67 axons were771analyzed in total. (A) The length of regenerating GCs at 2 hours after axotomy was decreased two-fold compared to native772axons. GC length was still significantly shorter up to 6 hours, but not 24 hours after axotomy (2 hours: $3.6 \ \mum \pm 0.4 \ \mum, 6$ 773hours: $5.0 \ \mum \pm 0.4 \ \mum, 24 \ hours: 5.3 \ \mum \pm 0.7 \ \mum, and native axons: 7.4 \ \mum \pm 0.5 \ \mum$). (B) The width of regenerating GCs

774 775 776 was reduced by almost 2.5-times at 2 hours after axotomy, compared to native axons. This trend did not normalize over 24 hours, where GC width was still decreased by 1.8-times compared to native group (2 hours: $1.9 \ \mu\text{m} \pm 0.3 \ \mu\text{m}$, 24 hours: 2.6 μ m \pm 0.4 μ m, and native axons: 4.6 μ m \pm 0.5 μ m).

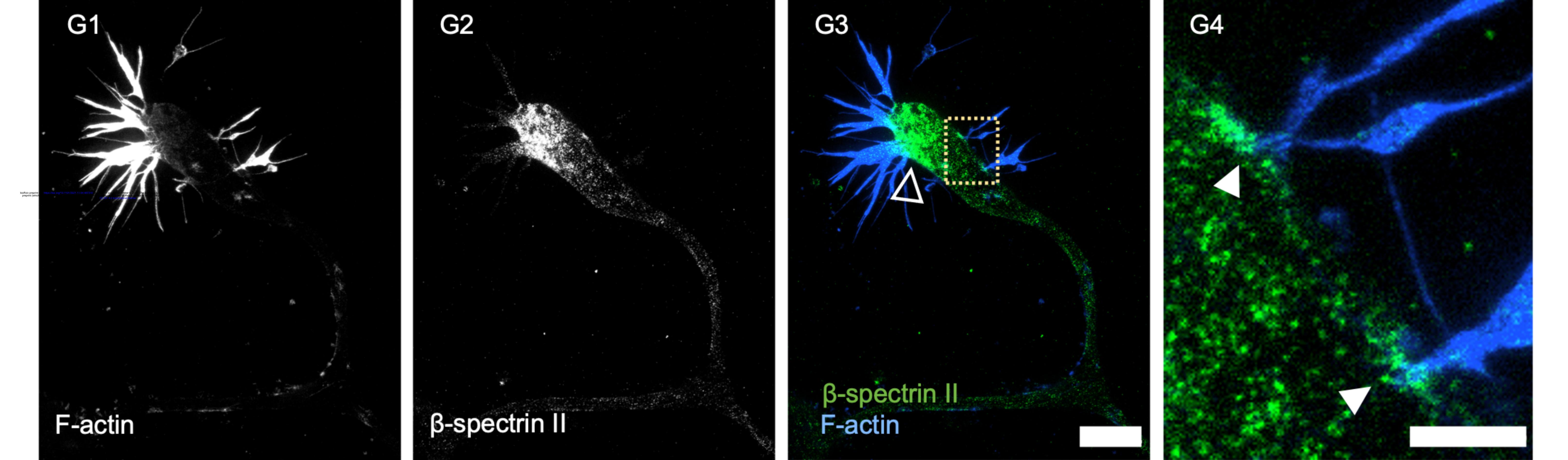


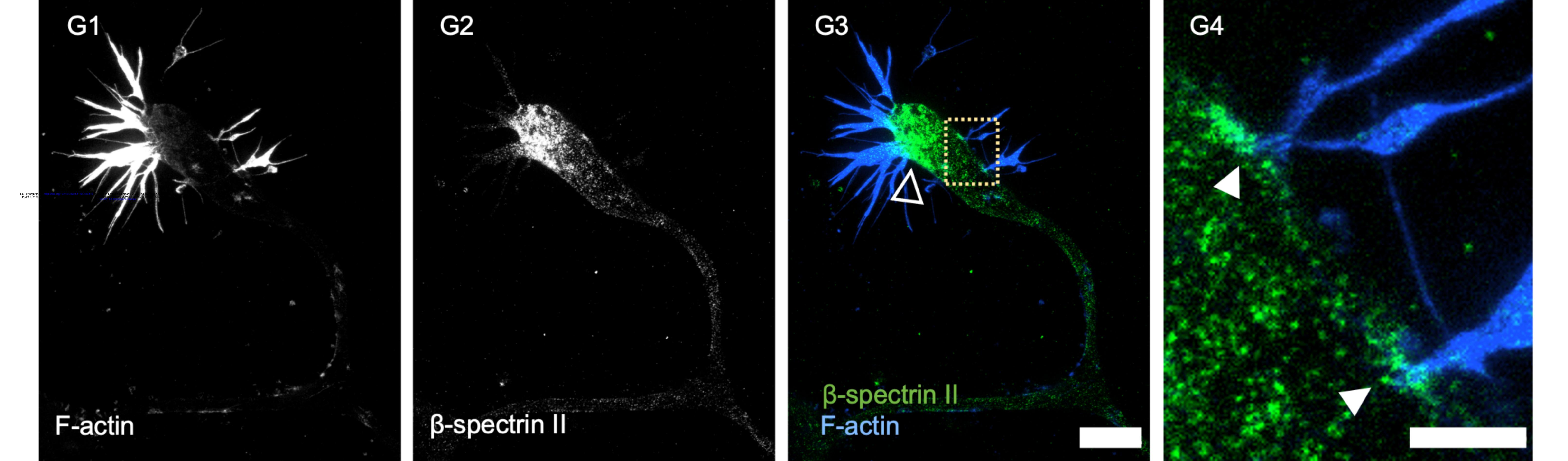
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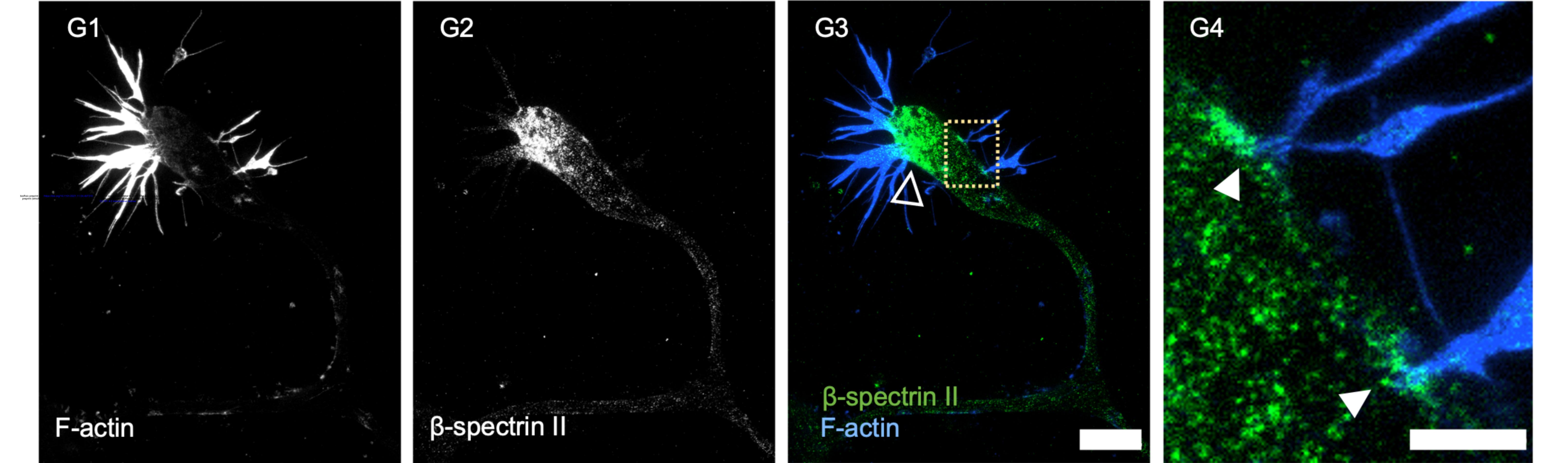
778 779 780 Supplemental Figure 3. Diameter of the first 10 µm of axon adjacent to the growth cone in calpeptin or DMSO treated axons. Cells were cultured for 11 days. On DIV 11, calpeptin or DMSO was administered into the axonal compartment for 1 hour and axotomized afterward. Regenerating axons were fixed 120 minutes later and stained for spectrin and tubulin. 781 782 Axonal diameter was then measured at 10 μ m from the GC. Data is given as mean \pm SD. At least 7 Axons were analyzed per condition, n=3, 15 axons were analyzed in total. No significant difference was detected according to unpaired t test.

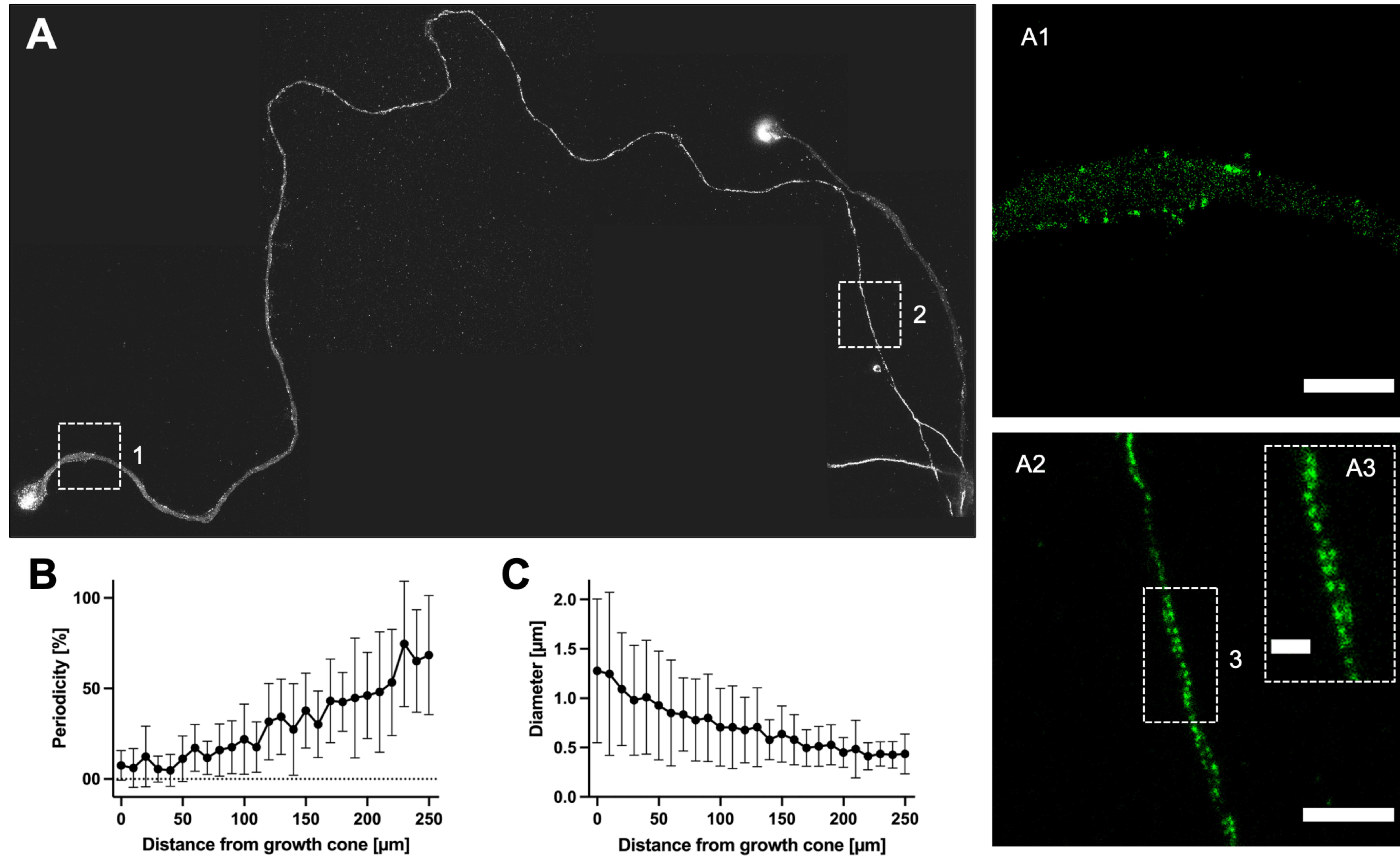


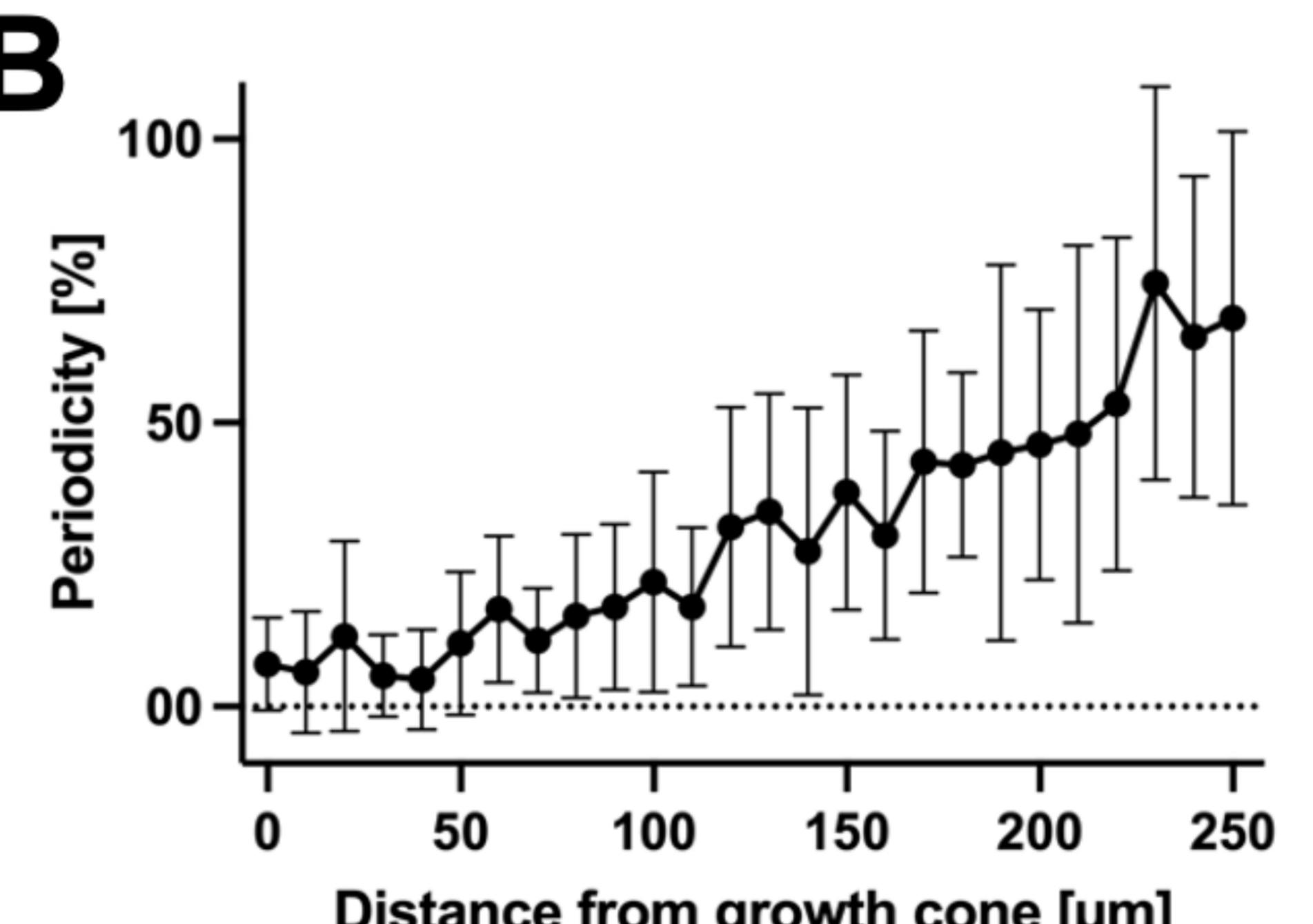


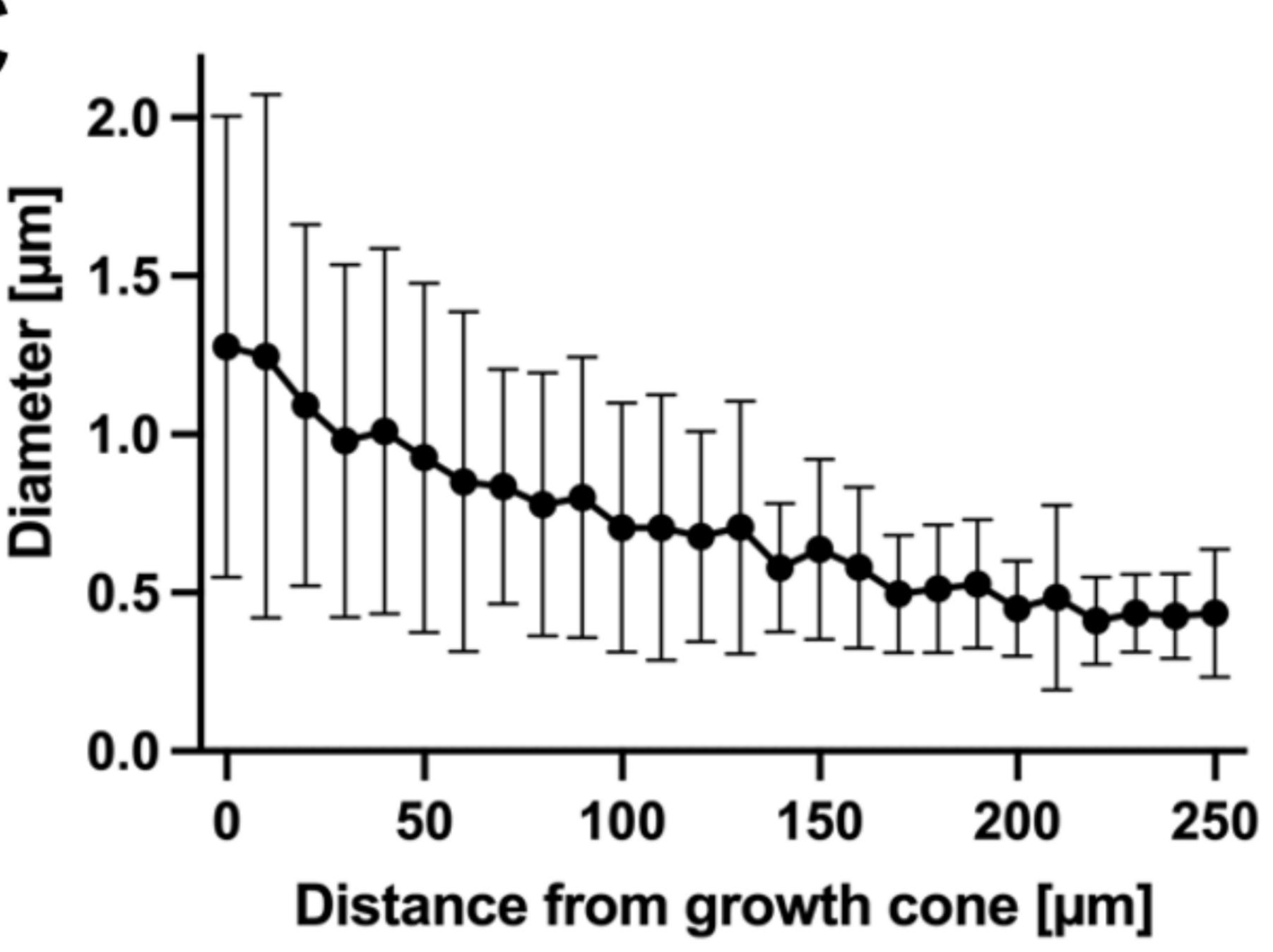


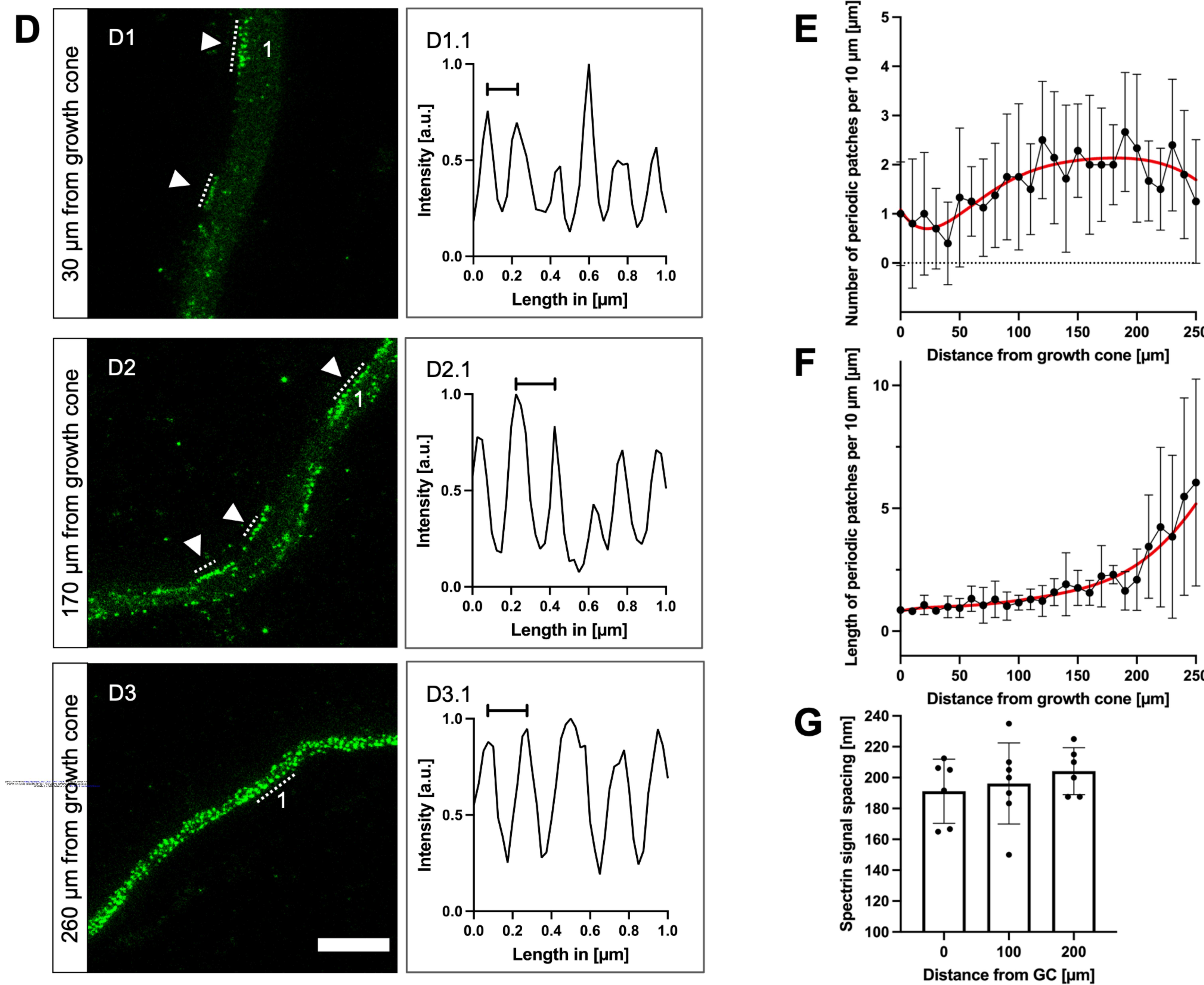


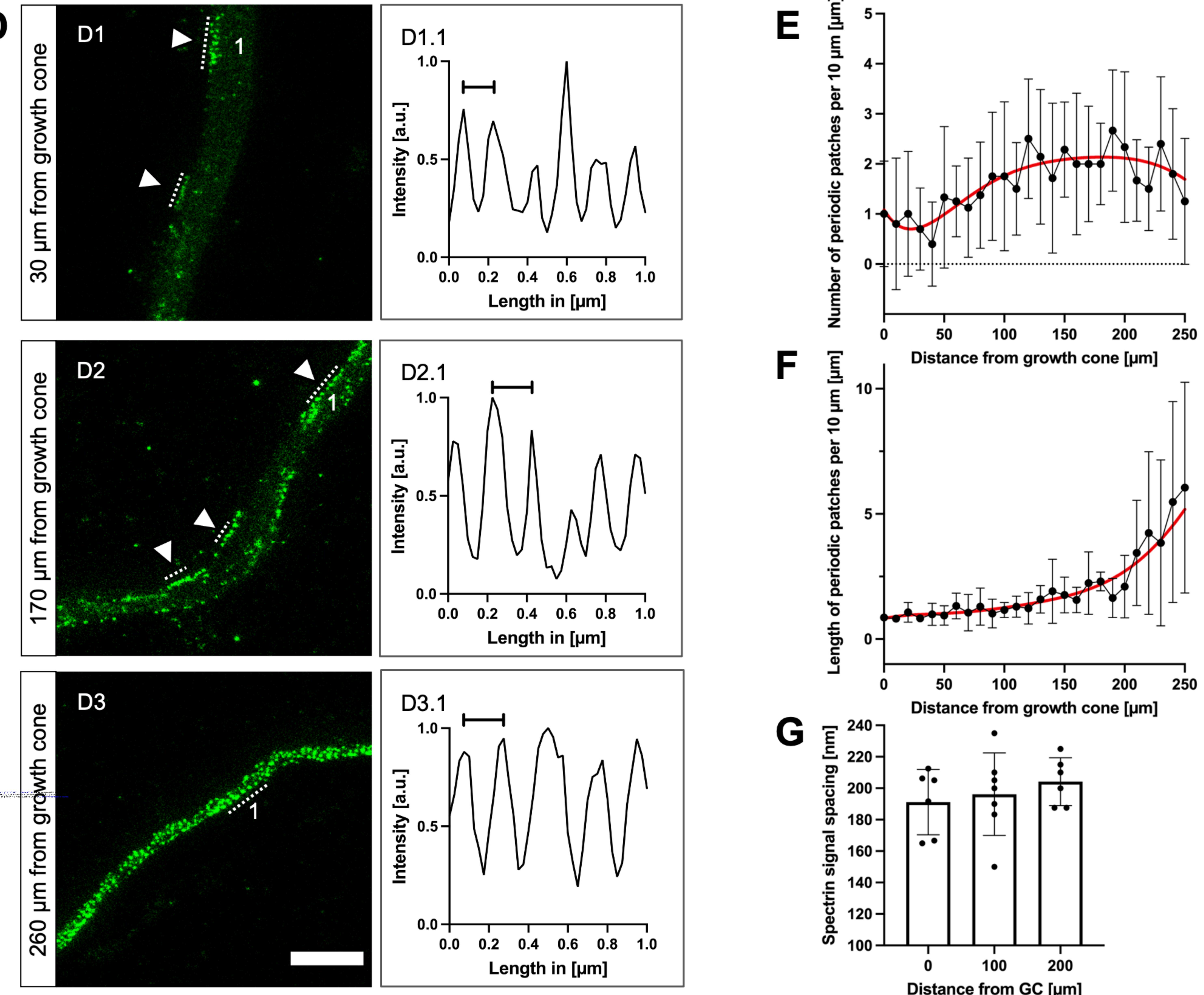


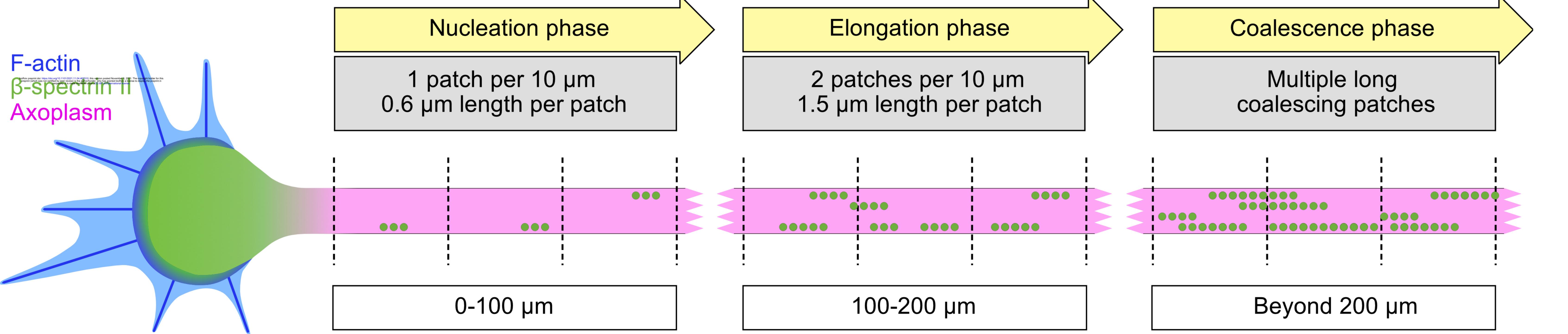





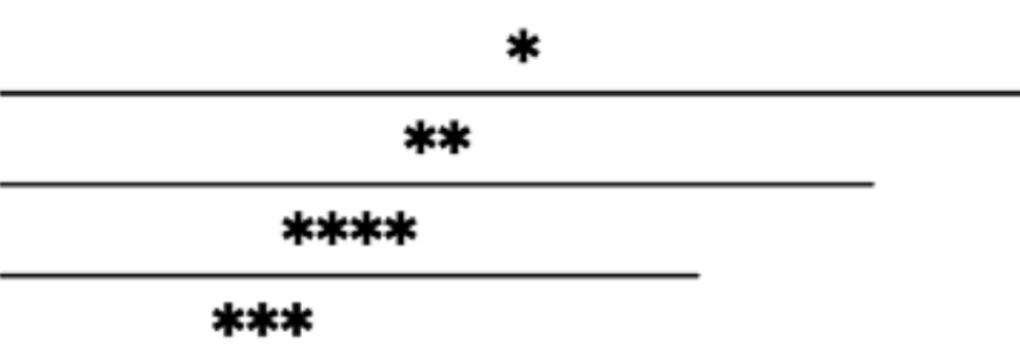


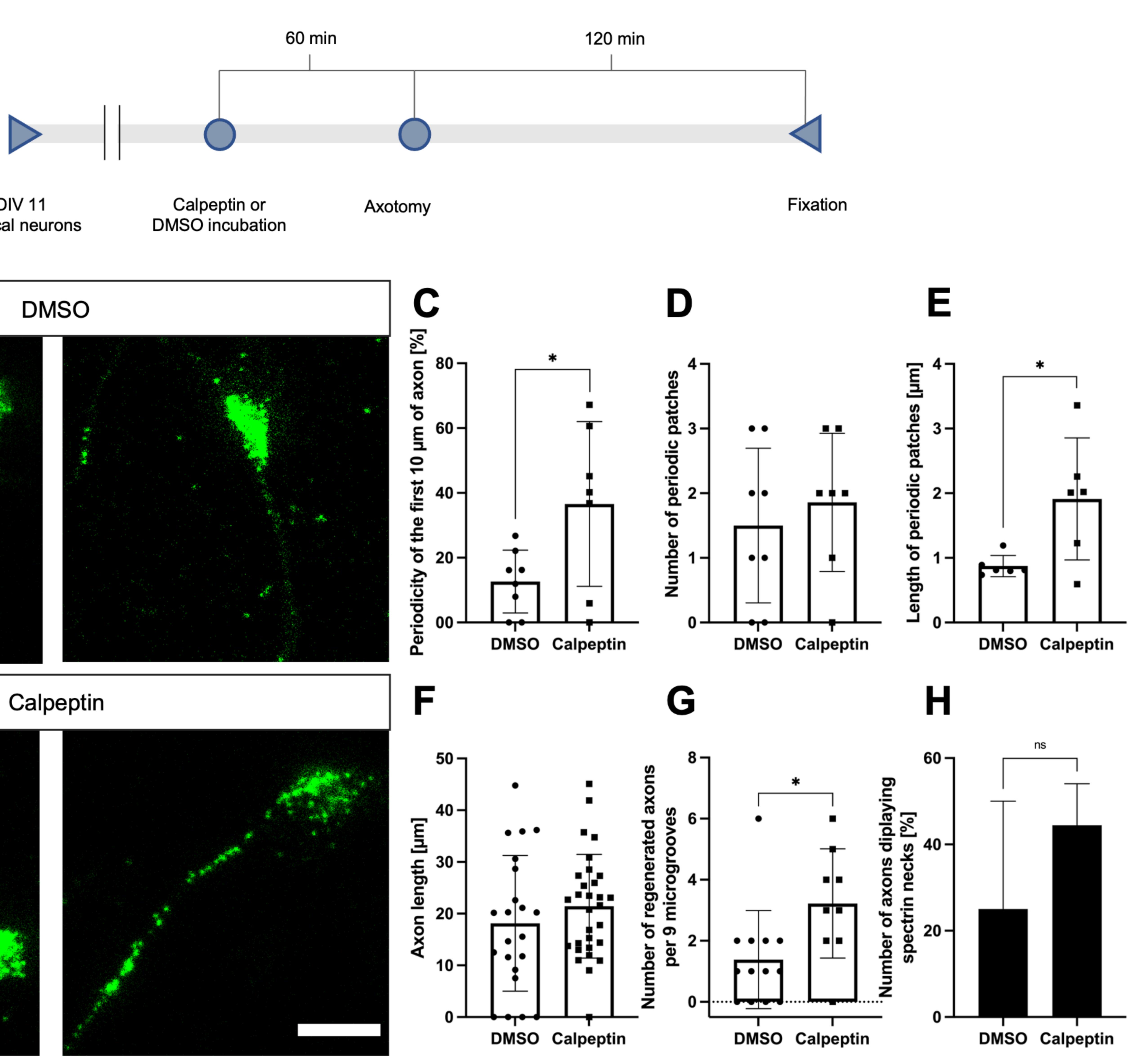




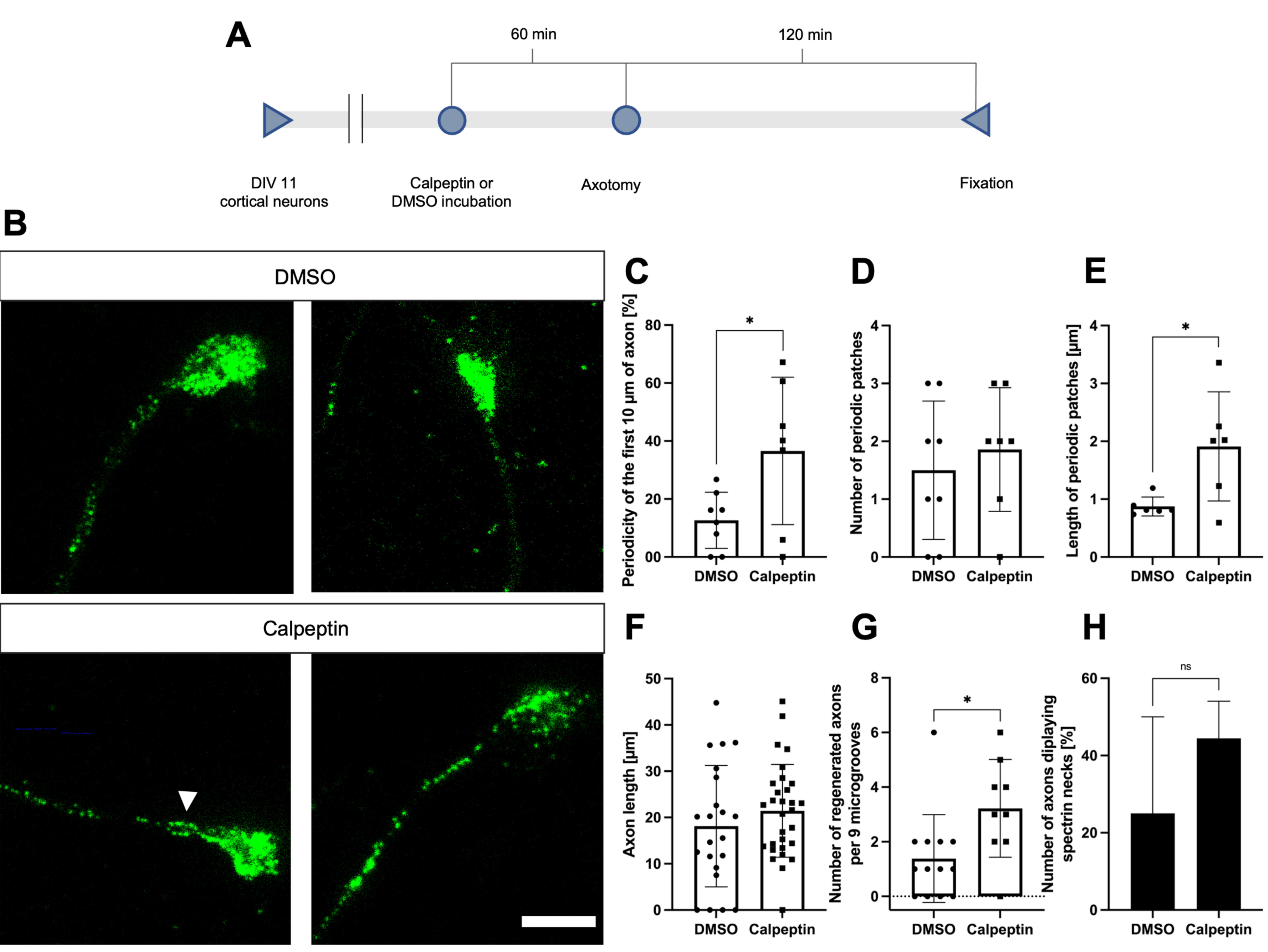


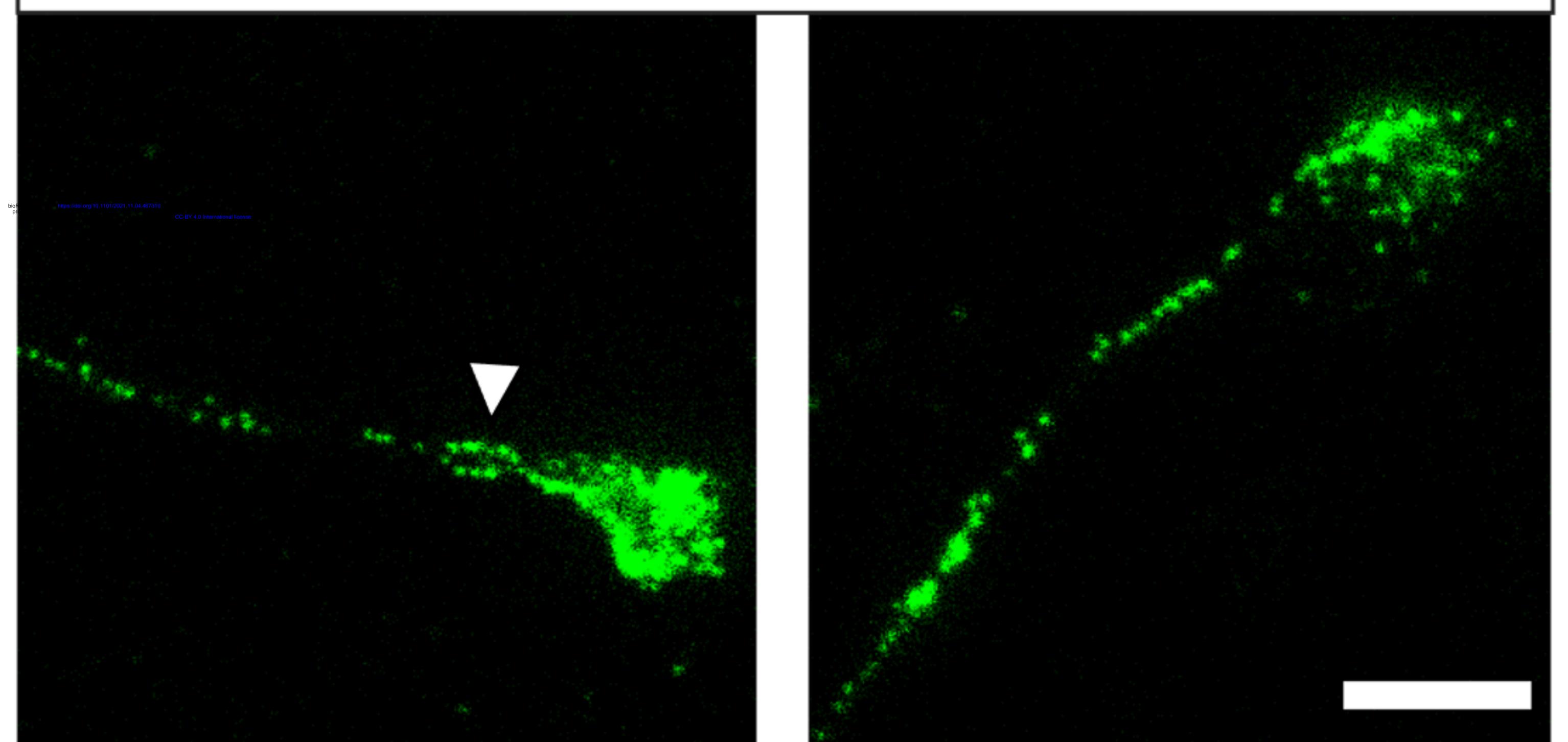




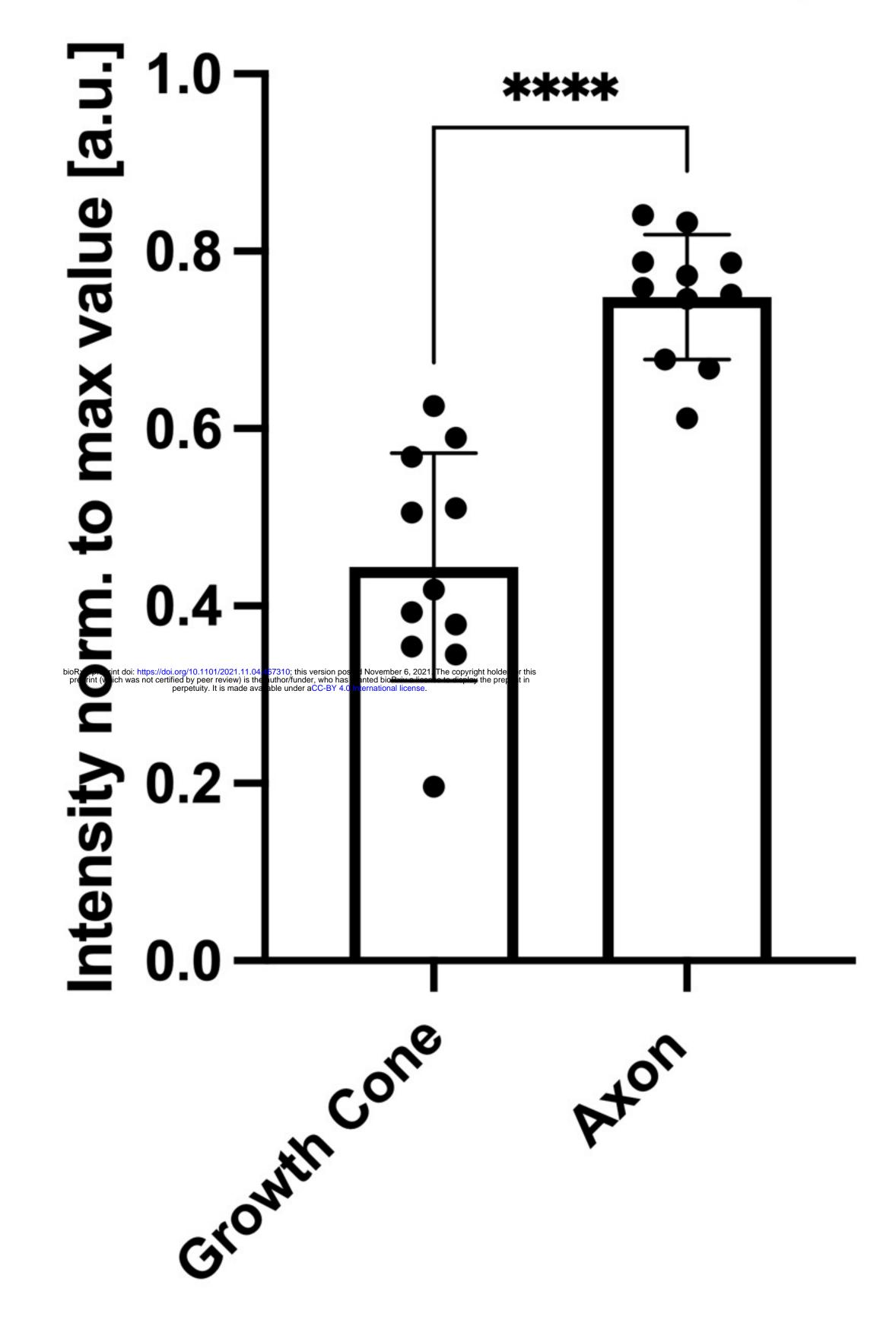


DIV 11

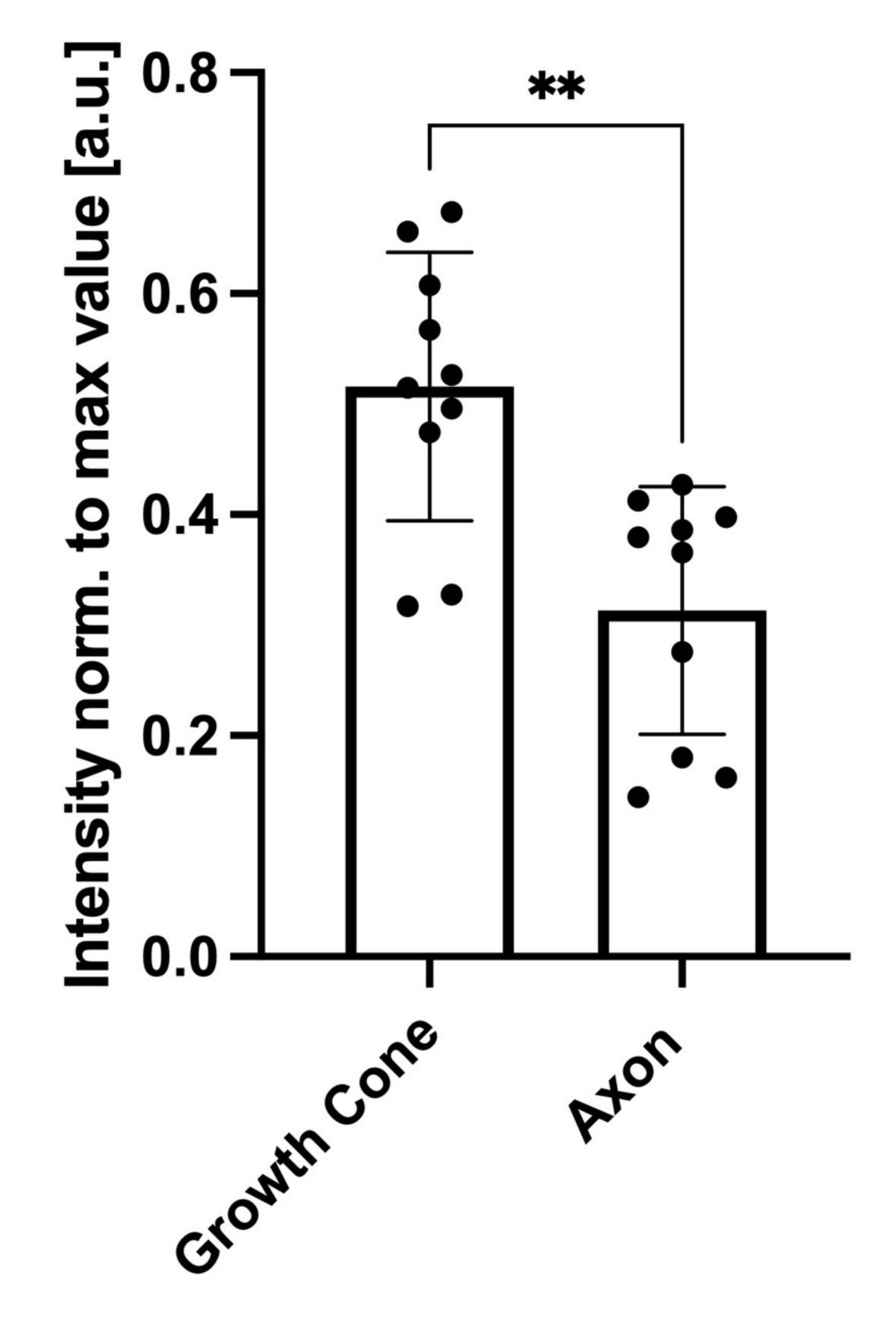




Tubulin Intensity

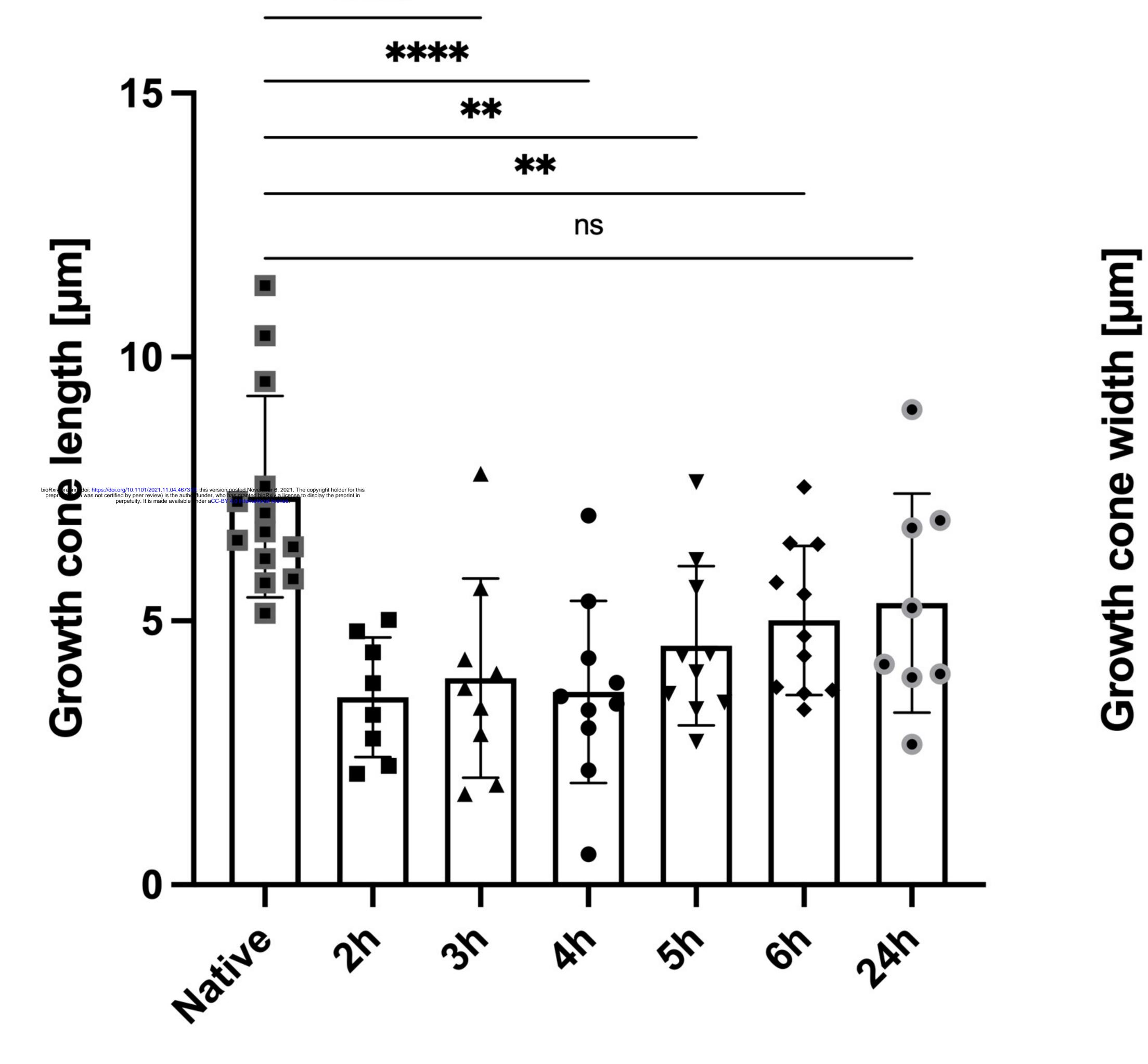


Spectrin Intensity









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