

Calcineurin-dependent cofilin activation and increased retrograde actin flow drive 5-HT-dependent neurite outgrowth in *Aplysia* bag cell neurons

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ABSTRACT Neurite outgrowth in response to soluble growth factors often involves changes in intracellular Ca^{2+} ; however, mechanistic roles for Ca^{2+} in controlling the underlying dynamic cytoskeletal processes have remained enigmatic. Bag cell neurons exposed to serotonin (5-hydroxytryptamine [5-HT]) respond with a threefold increase in neurite outgrowth rates. Outgrowth depends on phospholipase C (PLC) \rightarrow inositol trisphosphate \rightarrow Ca^{2+} \rightarrow calcineurin signaling and is accompanied by increased rates of retrograde actin network flow in the growth cone P domain. Calcineurin inhibitors had no effect on Ca^{2+} release or basal levels of retrograde actin flow; however, they completely suppressed 5-HT-dependent outgrowth and F-actin flow acceleration. 5-HT treatments were accompanied by calcineurin-dependent increases in cofilin activity in the growth cone P domain. 5-HT effects were mimicked by direct activation of PLC, suggesting that increased actin network treadmilling may be a widespread mechanism for promoting neurite outgrowth in response to neurotrophic factors.

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INTRODUCTION

Soluble neurotrophic factors play an important role in development (Kennedy *et al.*, 1994; Ming *et al.*, 1997; Campbell and Holt, 2001; Briancon-Marjollet *et al.*, 2008). However, basic cytoskeletal mechanisms by which soluble factors affect rates of neuronal outgrowth remain poorly understood. Serotonin (5-hydroxytryptamine [5-HT]) is a soluble ligand that can signal through G(q)-coupled receptors in *Aplysia* neurons (Li *et al.*, 1995, 2005; Cai *et al.*, 2008) and trigger phospholipase C (PLC)- and inositol trisphosphate (IP_3)-dependent Ca^{2+} release from intracellular endoplasmic reticulum (ER) stores in bag cell neuron growth cones (Zhang and Forscher, 2009). The

efficacy and spatial characteristics of Ca^{2+} release can be modulated by activity of the small GTPase Rac1. Rac1 activity promotes microtubule assembly and microtubule-dependent ER Ca^{2+} store transport into peripheral regions of the growth cone. Rac1 activity also promotes reactive oxygen species production, which sensitizes IP_3 -dependent Ca^{2+} release (Gordeeva *et al.*, 2003; Zhang and Forscher, 2009).

Here we investigate how release of Ca^{2+} from intracellular stores affects actin filament dynamics involved in neurite outgrowth. We show that 5-HT application results in increased rates of neurite outgrowth, accompanied by increased rates of retrograde F-actin network flow. 5-HT-evoked growth involves Ca^{2+} release from IP_3 -gated stores and calcineurin (protein phosphatase 2B)-dependent activation of the actin-recycling protein cofilin. Of interest, the resulting increases in actin network flow were independent of myosin II activity, whereas increases in neurite outgrowth were myosin II dependent.

RESULTS

5-HT induces neurite outgrowth on laminin substrates

Several lines of evidence suggest that laminin-integrin interactions activate Rac1 and such activity is correlated with growth cone

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Abbreviations used: CN-AIP, calcineurin autoinhibitory peptide; CsA, cyclosporin A; FSM, fluorescent speckle microscopy; 5-HT, serotonin; MLCK, myosin light-chain kinase.

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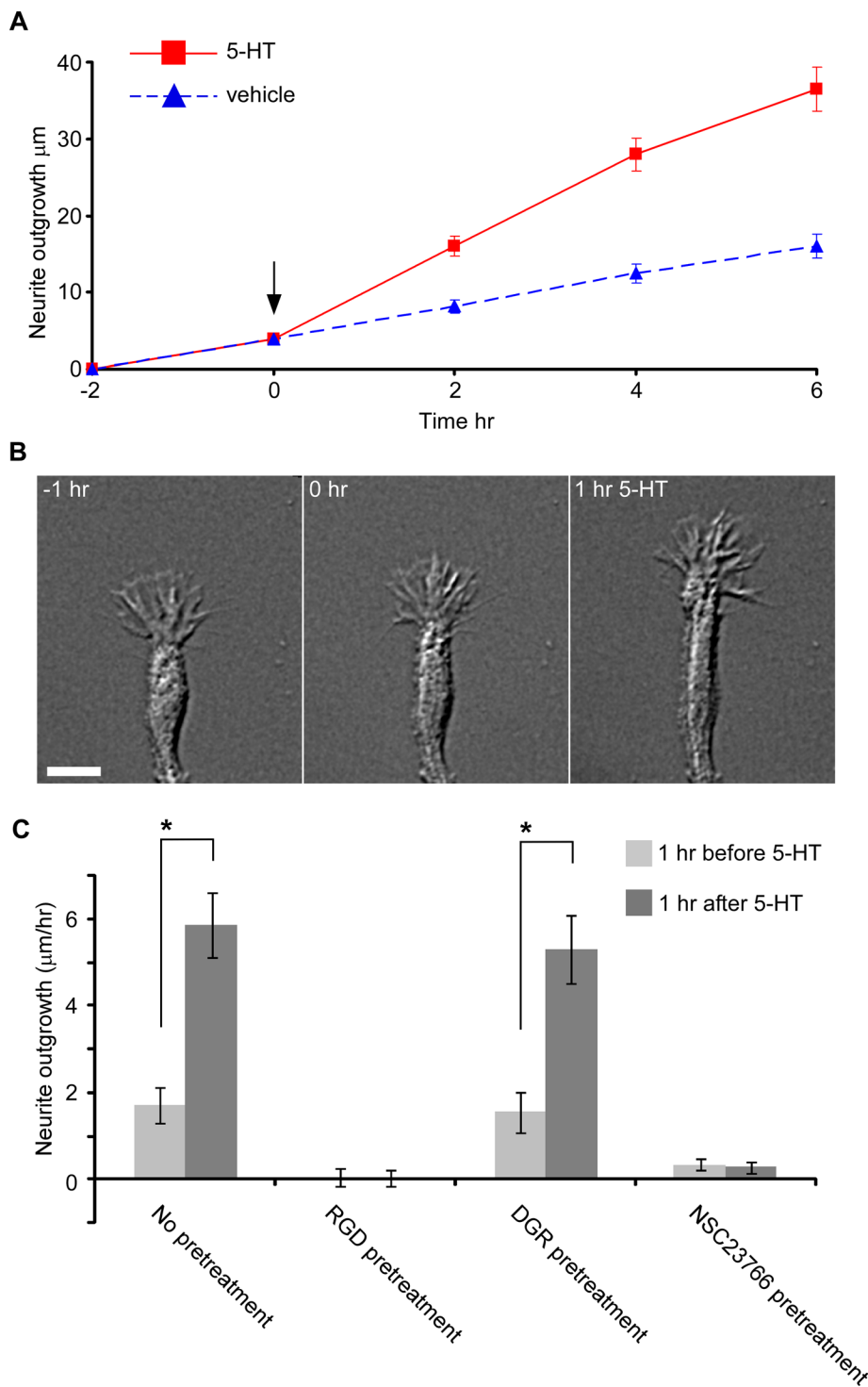


FIGURE 1: 5-HT induces neurite outgrowth on laminin substrates. (A) Neurite outgrowth on laminin substrates over time under control conditions (number of neurites tested [N] = 48) or before and after addition of 10 μ M 5-HT (N = 46). Black arrow, 5-HT or vehicle addition. Data points are averages \pm SEM. (B) A representative example of 5-HT (10 μ M, 1 h) effect on neurite outgrowth. DIC image; bar, 10 μ m. (C) Summary of neurite outgrowth rates 1 h before and after 5-HT (10 μ M) addition under these conditions: control (number of growth cones tested [N] = 58), RGD (50 μ M, 1-h pretreatment; N = 55), DGR (50 μ M, 1-h pretreatment; N = 55), NSC23766 (0.1 mM, 1-h pretreatment; N = 58). * p < 0.001. Values are mean \pm SEM. Statistical analysis by two tailed paired t test.

advance (Kuhn *et al.*, 1998; Grabham *et al.*, 2003; Matsuo *et al.*, 2003; Laforest *et al.*, 2005). On the basis of our previous finding that Ca^{2+} release from IP_3 stores was Rac dependent, we hypothesized

that culturing neurons on laminin substrates might increase basal Rac activity to a level at which 5-HT would elicit Ca^{2+} release without the use of constitutively active Rac constructs (Zhang and Forscher, 2009).

Bag cell neurons were cultured on laminin substrates and neurite outgrowth assessed by differential interference contrast (DIC) time-lapse imaging for 2 h before and 6 h after 5-HT (10 μ M) or vehicle addition (Figure 1A). Under these conditions, 5-HT treatment resulted in approximately three-fold increase in average neurite outgrowth rate (Figure 1A). Because neurite outgrowth rates were more or less constant for up to 6 h in 5-HT, we chose to analyze population responses before and after 1 h of 5-HT exposure (Figure 1, B and C) when a \sim 3.5-fold increase in neurite outgrowth rate was typically observed (Figure 1C).

Both basal and 5-HT-dependent outgrowth rates were strongly attenuated by RGD peptide, which competitively inhibits laminin-integrin interactions (Gruenbaum and Carew, 1999; Tucker *et al.*, 2005). In contrast, the reverse sequence DGR peptide control had no effect. These results confirm the specificity of the permissive role laminin plays in supporting both basal (Turney and Bridgman, 2005) and 5-HT-evoked (Figure 1C) neurite outgrowth. To test whether growth on laminin depended on Rac GTPase activity, we used a Rac1-specific small-molecule inhibitor, NSC23766 (Gao *et al.*, 2004). NSC23766 inhibited both basal and 5-HT-stimulated growth, consistent with the reported role of Rac activity in integrin function (Figure 1C; Kuhn *et al.*, 1998; Grabham *et al.*, 2003; Matsuo *et al.*, 2003; Laforest *et al.*, 2005).

5-HT increases peripheral retrograde F-actin flow rate

Next we investigated actin filament dynamics before and during 5-HT-evoked growth responses. Neurons were injected with either Alexa 568-G-actin or Alexa 594-phalloidin at trace levels to generate F-actin speckles for kinetic analysis. Features of Alexa 568-G-actin speckles incorporated into F-actin were tracked over time using a previously reported quantitative cross-correlation approach (Ji and Danuser, 2005; Burnette *et al.*, 2007; Hu *et al.*, 2007). Resulting retrograde actin flow velocities were pseudocolor coded and corresponding vectors overlaid on images to illustrate actin translocation.

Figure 2A is a representative example of Alexa 568-G-actin fluorescent speckle microscopy (FSM) data (top) and corresponding flow maps (bottom) from a growth cone before and after 5-HT treatment. 5-HT exposure elicited $6.6 \pm 1.0\%$ and

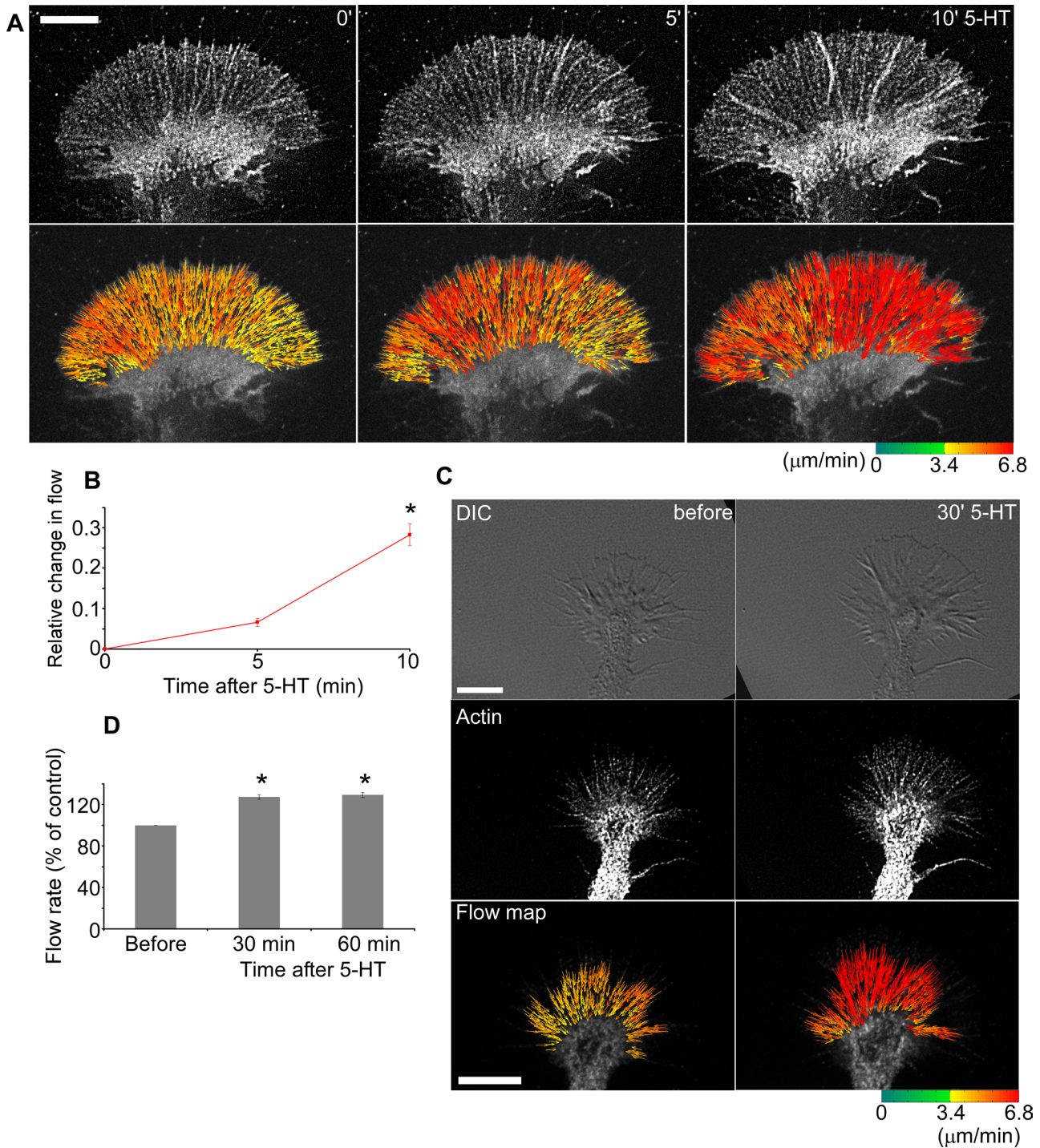


FIGURE 2: 5-HT increases peripheral retrograde actin filament flow rate. (A) Representative G-actin FSM images (top) and corresponding flow maps (bottom) from a growth cone before and after 5 or 10 min in 5-HT. Bar, 10 μm . Flow map colors encode speed (see scale bar), and arrows indicate flow direction. (B) Summary of relative changes in retrograde actin flow rates in response to 5-HT. Images acquired every 5 s with 1-min elapsed recording time and flow rates assessed as in A. Number of growth cones tested ($N = 3$). * $p < 0.01$ vs. before 5-HT addition. (C) DIC (top), Alexa 594-phalloidin FSM (middle), and corresponding flow map (bottom) of a growth cone before and after 30 min in 5-HT. Bar, 10 μm . Images acquired every 5 s with 2-min elapsed recording time. (D) Summary of P domain retrograde flow rates in response to 5-HT (10 μM , 30 min, and 60 min). Data normalized to rates before 5-HT addition. Number of growth cones evaluated ($N = 25$). Values are mean \pm SEM. * $P < 0.001$ vs. before 5-HT addition. Statistical analysis by two-tailed paired t test.

28.3 \pm 2.7% increases in F-actin flow rates at 5- and 10-min time points, respectively (Figure 2B). To investigate whether growth cones maintained accelerated F-actin flow rates during prolonged periods

of evoked outgrowth, we assessed F-actin dynamics 30 and 60 min after 5-HT treatment. Figure 2C illustrates representative growth cone structures (DIC), F-actin distributions, and corresponding actin

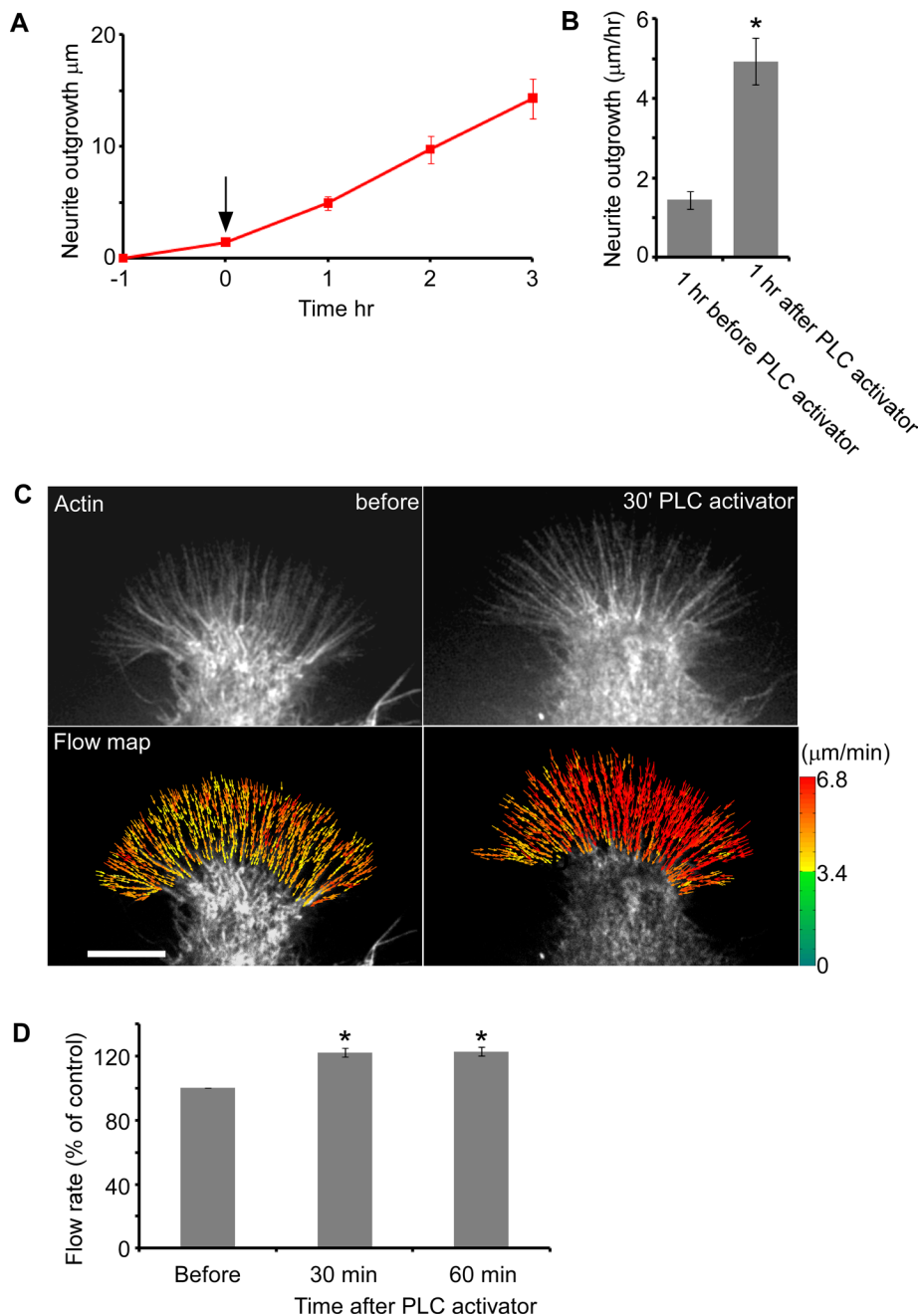


FIGURE 3: PLC activator increases peripheral retrograde actin filament flow rate and promotes neurite outgrowth. (A) Neurite outgrowth on laminin substrates over time before and after addition of 25 μM PLC activator m-3M3FBS (number of neurites tested [N] = 32). Black arrow, m-3M3FBS addition. Data points are averages \pm SEM. (B) Comparison of neurite outgrowth rates 1 h before and after m-3M3FBS (25 μM) addition. Data is taken from A. * $p < 0.001$. Values are mean \pm SEM. Statistical analysis by two-tailed paired t test. (C) Alexa 594–phalloidin FSM (top) and corresponding flow map (bottom) from a growth cone before and after 30 min in m-3M3FBS (25 μM). Bar, 10 μm . Flow map generated as described. Images acquired every 5 s with 2-min elapsed recording time. (D) Summary of normalized P domain retrograde flow rates in response to 25 μM m-3M3FBS. Data normalized to rates before m-3M3FBS addition. $N = 12$ growth cones. * $p < 0.001$ vs. before m-3M3FBS addition. Values are mean \pm SEM.

flow maps before and after 30 min in 5-HT. Rates of growth cone advance were correlated with increased F-actin flow rates (Figure 2C, DIC vs. flow map; see Supplemental Movie S1). On average, F-actin flow rates significantly increased by $27.2 \pm 2.1\%$ and $29.3 \pm 2.5\%$ after 30- and 60-min 5-HT exposures, respectively (Figure 2D).

S3). Finally, pretreatment with the Rac inhibitor NSC23766 abolished 5-HT-dependent Ca^{2+} responses (Supplemental Figure S2). Collectively, these data show that 5-HT triggers sustained Ca^{2+} increases in parallel with increased retrograde F-actin flow when Rac levels are sufficiently elevated.

Taken together, these results indicate that 5-HT treatment triggers a persistent increase in peripheral F-actin flow that is well correlated with the observed accelerated rate of growth cone advance.

Direct phospholipase C activation increases neurite growth and retrograde flow rates

To investigate the generality of this response, we bypassed the 5-HT receptor and directly activated PLC—the downstream effector of receptor signaling through trimeric Gq proteins (Figure 3). PLC activation with m-3M3FBS (25 μM ; Bae et al., 2003; Li et al., 2009) resulted in sustained ~ 3.5 -fold increases in neurite outgrowth rates (Figure 3, A and B) accompanied by increases in retrograde actin flow (Figure 3, C and D) essentially identical to those observed after 5-HT treatments (Figure 2).

5-HT-induced F-actin flow increases and outgrowth depend on Ca^{2+} release from IP_3 -gated stores

Ca^{2+} is known to participate in 5-HT function in neurons (Dropic et al., 2005; Li et al., 2005; Cai et al., 2008). Previously we reported that 5-HT evoked Ca^{2+} release in *Aplysia* growth cones in the presence of constitutively active (but not dominant negative) Rac1 when cells were plated on PLL substrates (Zhang and Forscher, 2009). Given that laminin has been widely reported to increase Rac activity (Kuhn et al., 1998; Grabham et al., 2003; Matsuo et al., 2003; Laforest et al., 2005) and the robust Rac dependence of the outgrowth responses under study (Figure 1C), we tested whether 5-HT would elicit Ca^{2+} release in growth cones on laminin substrates. Supplemental Figure 1A shows that 5-HT treatments indeed evoked rapid and sustained Ca^{2+} increases in the entire growth cone (see also Supplemental Movie S2). These responses depended on the PLC \rightarrow IP_3 signaling cascade since the PLC inhibitor U73122 (Jin et al., 1994; Zhou et al., 1999) or the IP_3 receptor blocker xestospongine C (XeC; Gafni et al., 1997) abolished 5-HT-evoked Ca^{2+} responses (Supplemental Figure S1, A and B). During more prolonged 5-HT exposure (1 h), Ca^{2+} levels remained elevated $\sim 22\%$ above baseline. These Ca^{2+} changes were also inhibited by pretreatment with PLC or IP_3 receptor antagonists (Supplemental Figure 1, C and D, and Supplemental Movie

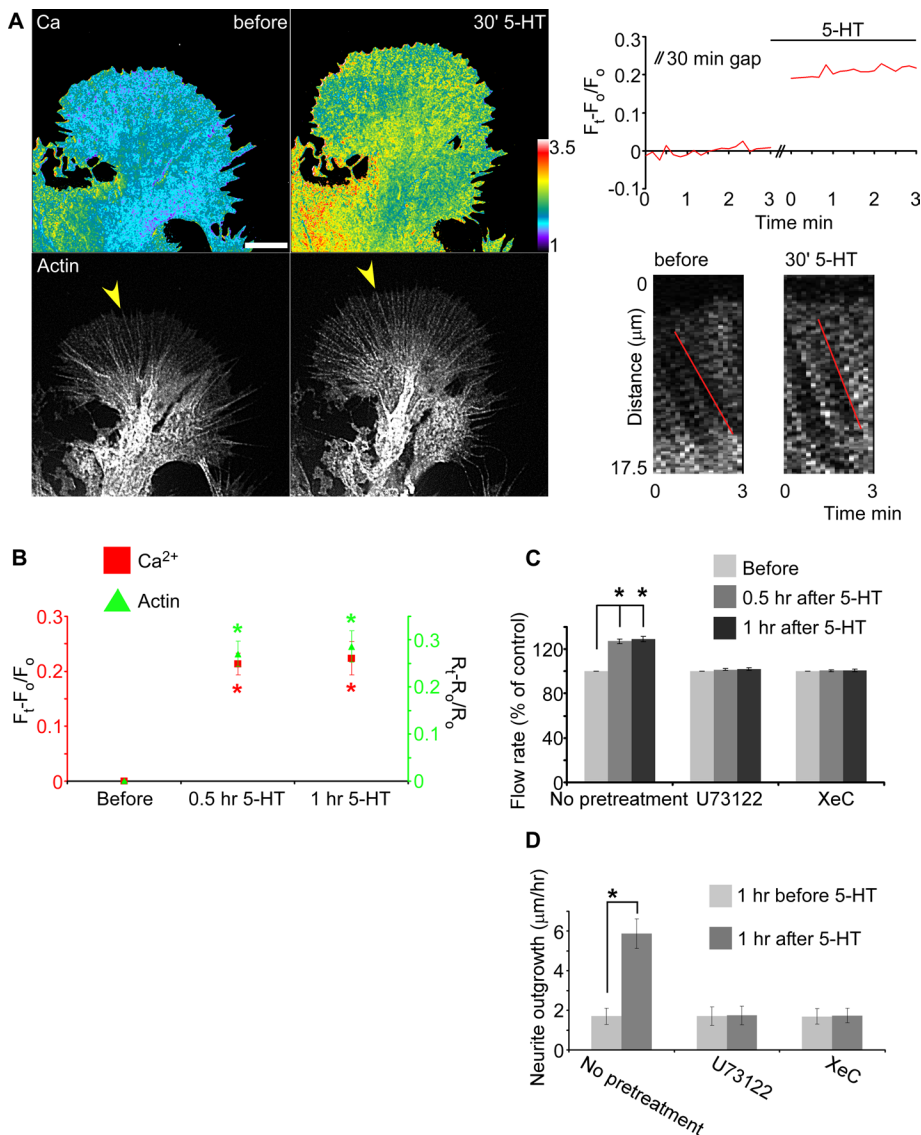


FIGURE 4: 5-HT-evoked Ca²⁺ release and peripheral retrograde actin flow increases are correlated and necessary for induced neurite outgrowth. (A) Tandem Ca²⁺ ratio imaging and actin FSM. Ca²⁺ ratio image (top) and Alexa 594-phalloidin FSM (bottom) of a growth cone before and after 30 min in 5-HT. Bar, 10 μm. The Ca²⁺ ratio is encoded by a linear pseudocolor scale. Images acquired every 10 s with 3-min elapsed recording time. Top right, growth cone Ca²⁺ response to 5-HT over time. F₀, average Ca²⁺ level before 5-HT addition. Bottom right, kymographs sampled from area of interest indicated in actin panel (yellow arrowheads). Retrograde flow rate before and after 30 min of 5-HT exposure: 4.88 ± 0.21 and 6.12 ± 0.19 μm min⁻¹, respectively (mean ± SD, N = 5 measurements). (B) Comparison of Ca²⁺ levels and P domain flow rates before and after 5-HT. The Ca²⁺ ratio imaging and FSM were carried out simultaneously. Images acquired every 5 or 10 s with 2- to 3-min elapsed recording time. F₀, average Ca²⁺ level before 5-HT addition. R₀, average retrograde flow rate before 5-HT addition. N = 14 growth cones. *p < 0.001 vs. before 5-HT addition. (C) Retrograde flow rates in response to 5-HT in various conditions normalized to before 5-HT addition. N = 25 growth cones (control), N = 18 (U73122, 2 μM, 30-min pretreatment), and N = 21 (XeC, 20 μM, 30-min pretreatment). Control from Figure 2D included for comparison. *p < 0.001 vs. before 5-HT addition. (D) Summary of neurite outgrowth 1 h before and after 5-HT addition in various conditions. Control (N = 58 growth cones), U73122 (2 μM, 1-h pretreatment, N = 34), and XeC (20 μM, 1-h pretreatment, N = 37). Control from Figure 1C is included for comparison. *p < 0.001 vs. before 5-HT addition. Values are mean ± SEM. Statistical analysis was done by two-tailed paired t test.

To directly address the relationship between Ca²⁺ levels and F-actin dynamics, we performed simultaneous ratiometric Ca²⁺ imaging and F-actin FSM (see Supplemental Movie S4, A and B, for

multichannel fluorescence imaging). Figure 4A shows Ca²⁺ levels and FSM records from a growth cone before and after a 30-min 5-HT treatment. Ca²⁺ levels and peripheral F-actin flow rates increased in parallel by 20.9 and 25.4%, respectively. There was a strong temporal correlation between average 5-HT-evoked Ca²⁺ elevations and increases in F-actin flow (Figure 4B). When Ca²⁺ release from intracellular stores was blocked by inhibiting PLC or IP₃ using U73122 or XeC, respectively, no changes in baseline actin flow or growth rates were observed; however, 5-HT-evoked increases in F-actin flow and concomitant increases in neurite outgrowth were completely suppressed (Figure 4, C and D). Taken together, these results indicate that IP₃-dependent Ca²⁺ mobilization is upstream of, and necessary for, the increased F-actin flow rates and growth-promoting effects of 5-HT observed.

5-HT effect on F-actin flow is independent of myosin light-chain kinase activity

Given that 5-HT-induced increases in F-actin flow were Ca²⁺ dependent and myosin II activity is known to affect F-actin flow rates (Lin *et al.*, 1996; Medeiros *et al.*, 2006), we investigated whether activation of myosin light-chain kinase (MLCK), which is a Ca²⁺/calmodulin-dependent regulator of myosin II activity (Kamm and Stull, 2001; Schmidt *et al.*, 2002), could be the Ca²⁺ effector mediating flow increases. We used ML-7, a well-characterized MLCK inhibitor (Ruchhoeft and Harris, 1997; Zhou and Cohan, 2001; Zhang *et al.*, 2003) effective in our system (Zhang *et al.*, 2003). In control experiments, we found that exposure to ML-7 alone (15–20 min) did not alter Ca²⁺ levels (Supplemental Figure S3A; also see Figure 5B, inset vs. left); however, F-actin flow rates decreased by ~15% (Supplemental Figure S3B; also see Figure 5D, inset vs. left) consistent with previous reports that MLCK activity is involved in setting basal F-actin flow rates (Zhang *et al.*, 2003).

We next investigated the effect of MLCK inhibition on 5-HT-evoked Ca²⁺ release and F-actin flow. After MLCK inhibition, 5-HT continued to evoke Ca²⁺ release (+17%; Figure 5B), as well as increases in peripheral F-actin flow (+29%; Figure 5, C and D). Indeed, statistical analysis indicated that MLCK inhibition did not significantly affect the depth of 5-HT-evoked Ca²⁺ responses (Table 1) or increases in F-actin flow (Table 2). In summary, although MLCK plays a role in setting basal F-actin flow

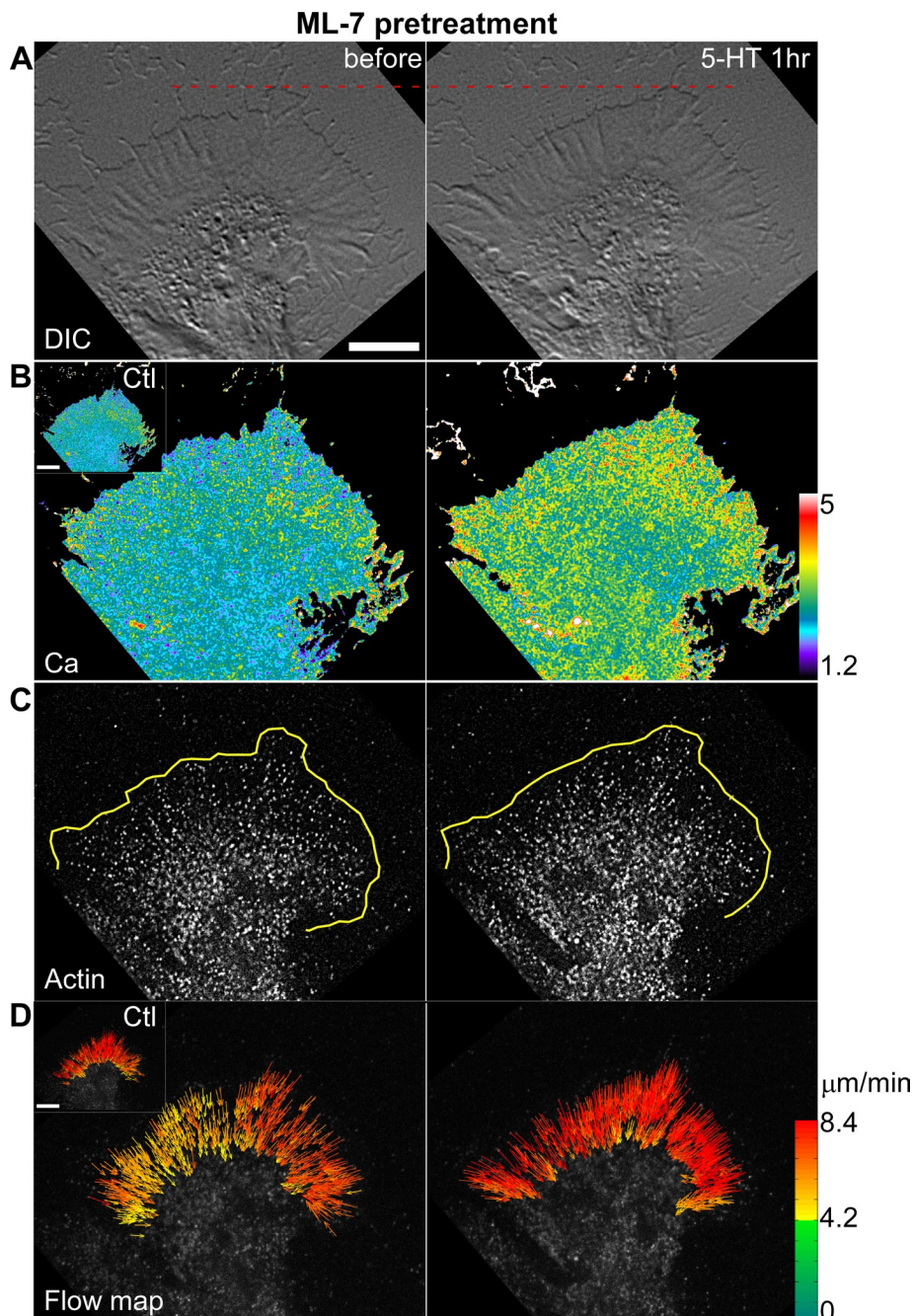


FIGURE 5: 5-HT-induced increase in actin flow is independent of MLCK activity. (A–D) A representative example of growth cone responding to 5-HT in ML-7 background. (A) DIC, (B) Ca^{2+} ratio, (C) Alexa 594–phalloidin FSM, (D) corresponding flow map. Cells pretreated with ML-7 (10 μM) for 15 min and ML-7 present throughout 1 h in 5-HT. Bar, 10 μm . Data acquired every 5 s with 2-min elapsed recording time. The Ca^{2+} level is encoded by a linear pseudocolor scale as before. Yellow lines demarcate leading edge in C. Insets in B and D are controls before ML-7 addition.

rates, 5-HT effects on actin network flow were essentially independent of MLCK activity.

Of interest, MLCK inhibition did appear to inhibit 5-HT-evoked neurite outgrowth (Figure 5A; dotted red line), suggesting that 5-HT effects on actin dynamics versus neurite growth might have differential myosin II dependences. To test this possibility, we further investigated the role of myosin II in 5-HT-mediated growth.

5-HT effects on F-actin flow are independent of myosin II activity

Previous evidence suggested that myosin II sets the maximum rate of retrograde F-actin flow in growth cones (Lin *et al.*, 1996; Medeiros *et al.*, 2006; Burnette *et al.*, 2007). To directly investigate the role myosin II plays in 5-HT effects, we used blebbistatin, a specific nonmuscle myosin II ATPase inhibitor (Straight *et al.*, 2003; Allingham *et al.*, 2005) that has been extensively characterized in our system (Medeiros *et al.*, 2006). Similar to what was previously observed for growth cones plated on PLL substrates, 10- to 20-min blebbistatin exposures promoted elongation of filopodial actin bundles, resulting in rearward expansion of the peripheral cytoplasmic domain (Supplemental Figure S4A and Supplemental Movie S5). During blebbistatin treatment peripheral F-actin flow rates typically decreased by ~20% (Supplemental Figure 4, A bottom, and B), consistent with nonmuscle myosin II playing a role in setting basal rates of retrograde F-actin flow on laminin as previously observed on PLL substrates (Medeiros *et al.*, 2006).

We then investigated a role for myosin II in 5-HT-induced increases in F-actin flow. Neurons were pretreated with blebbistatin for 20 min to maximally inhibit myosin II activity (Medeiros *et al.*, 2006) and then challenged with 5-HT. Remarkably, after blebbistatin pretreatment, 5-HT exposure continued to trigger increases in retrograde flow (~25.5% increase after 10 min; Figure 6A) at levels similar to that observed under control conditions (Figure 6B) that were sustained for up to 60 min (Figure 6C; kymographs show flow increases of 22.0 and 25.1% at 30- and 60-min time points, respectively). The average magnitude of 5-HT-induced actin flow increases was not significantly different between control and blebbistatin-pretreated growth cones (Figure 6D). These results strongly suggest that changes in myosin II activity are not involved in the 5-HT-induced increases in actin network flow and corresponding increased rates of actin filament turnover observed.

We then investigated whether myosin II activity is necessary for the more global effect of 5-HT on neurite outgrowth. To be consistent with the aforementioned experiments, outgrowth rates were assessed for 1 h under control conditions and during 5-HT exposure. We noted that in ~75% of the growth cones ($n = 90$), ~1.5 h of exposure to blebbistatin alone resulted in neurite branching accompanied by enhanced rates of neurite outgrowth. Similarly, long-term blebbistatin treatment has been reported to promote neurite outgrowth in chicken retina explants, medulla, and spinal cord neurons (Rosner *et al.*, 2007). Given the foregoing

	Control (no pretreatment)	ML-7 pretreatment
Before 5-HT: ($F_0 - F_0$)/ F_0	0 ± 0	0 ± 0
5-HT 30 min: ($F_{30} - F_0$)/ F_0	0.205 ± 0.023*	0.201 ± 0.026*
5-HT 60 min: ($F_{60} - F_0$)/ F_0	0.215 ± 0.025*	0.205 ± 0.027*
N	17	9

Summary of ($F_t - F_0$)/ F_0 recorded in the entire growth cone area quantifying Ca^{2+} response to 5-HT with or without MLCK inhibition. For MLCK inhibition, cells were pretreated with ML-7 (10 μ M) for 15–20 min and ML-7 was present throughout. Records acquired every 5 or 10 s with 2- to 3-min elapsed recording time. F_0 , average Ca^{2+} level before 5-HT addition. F_{30} and F_{60} , average Ca^{2+} level 30 and 60 min after 5-HT addition, respectively. N denotes number of growth cones evaluated. Values are expressed as mean ± SEM. Statistical analysis was done by two-tailed paired t test.

* $p < 0.001$ vs. before 5-HT addition. In addition, there was no significant difference in the magnitude of 5-HT responses in control vs. ML-7 group.

TABLE 1: MLCK inhibition does not affect 5-HT-evoked Ca^{2+} increases.

considerations, neurons were pretreated with blebbistatin for 10 min to establish maximal inhibitory effects on actin flow (Medeiros *et al.*, 2006) and then exposed to 5-HT for only 1 h in the continued presence of blebbistatin, at which time outgrowth rates were assessed. Blebbistatin treatment completely blocked 5-HT-evoked neurite outgrowth (Figure 6E; see also DIC in C), consistent with effects seen after MLCK inhibition (Figure 6E). Taken together, these results suggest that 5-HT treatment leads to increases in actin network flow that are independent of myosin II activity; however, myosin contractility appears to be essential to functionally couple increased actin network flow rates to the process of neurite advance.

5-HT-induced increase in F-actin flow depends on calcineurin activity

Calcineurin, or Ca^{2+} /calmodulin-dependent protein phosphatase 2B (PP2B), is enriched in growth cones and has been implicated in Ca^{2+} -dependent regulation of neurite extension and filopodium dynamics (Ferreira *et al.*, 1993; Chang *et al.*, 1995; Lautermilch and Spitzer, 2000; Cheng *et al.*, 2002; Arie *et al.*, 2009). Although cytoskeletal proteins have often been implicated in calcineurin actions, to our knowledge calcineurin effects on actin dynamics have never been directly assessed. Thus we first looked at potential effects of calcineurin inhibition alone on actin dynamics. To this end, we treated cells with the cell-permeable calcineurin inhibitor FK-506 or cyclosporin A (CsA; Liu *et al.*, 1991; Fruman *et al.*, 1992). The Ca^{2+} levels and F-actin flow rates did not differ significantly from controls after 20–40 min of FK-506 or CsA exposure alone (Supplemental Figure S5, A and B). As an alternative approach, we injected a specific calcineurin inhibitor consisting of the conserved calcineurin autoinhibitory peptide domain (CN-AIP; Hashimoto *et al.*, 1990; Perrino *et al.*, 1995) into cells. CN-AIP injection also had no effect on F-actin flow rates (unpublished data). These observations indicate that basal levels of retrograde actin flow do not depend on calcineurin activity.

We next investigated effects of calcineurin inhibition on 5-HT-evoked Ca^{2+} release. In CN-AIP-injected or FK-506- or CsA-pretreated neurons, 5-HT continued to evoke typical levels of Ca^{2+}

	No pretreatment	ML-7 pretreatment
Flow rate before 5-HT	100 ± 0	100 ± 0
Flow rate 30 min after 5-HT	127.2 ± 2.1*	124.8 ± 2.7*
Flow rate 60 min after 5-HT	129.3 ± 2.5*	126.1 ± 2.7*
N	25	16

Data normalized to flow rates before 5-HT addition (percentage of before 5-HT) to quantify changes of P domain retrograde flow rates in response to 5-HT with or without MLCK inhibition. For MLCK inhibition, cells were pretreated with ML-7 (10 μ M) for 15–20 min and ML-7 was present throughout. Data were acquired every 5 or 10 s with 2- to 3-min elapsed recording time. N denotes the number of growth cones tested. Values are expressed as mean ± SEM. Statistical analysis was done by two-tailed paired t test.

* $p < 0.001$ vs. before 5-HT addition. In addition, there was no significant difference in the magnitude of 5-HT responses in control vs. ML-7 group.

TABLE 2: MLCK inhibition does not affect 5-HT-evoked actin flow increases.

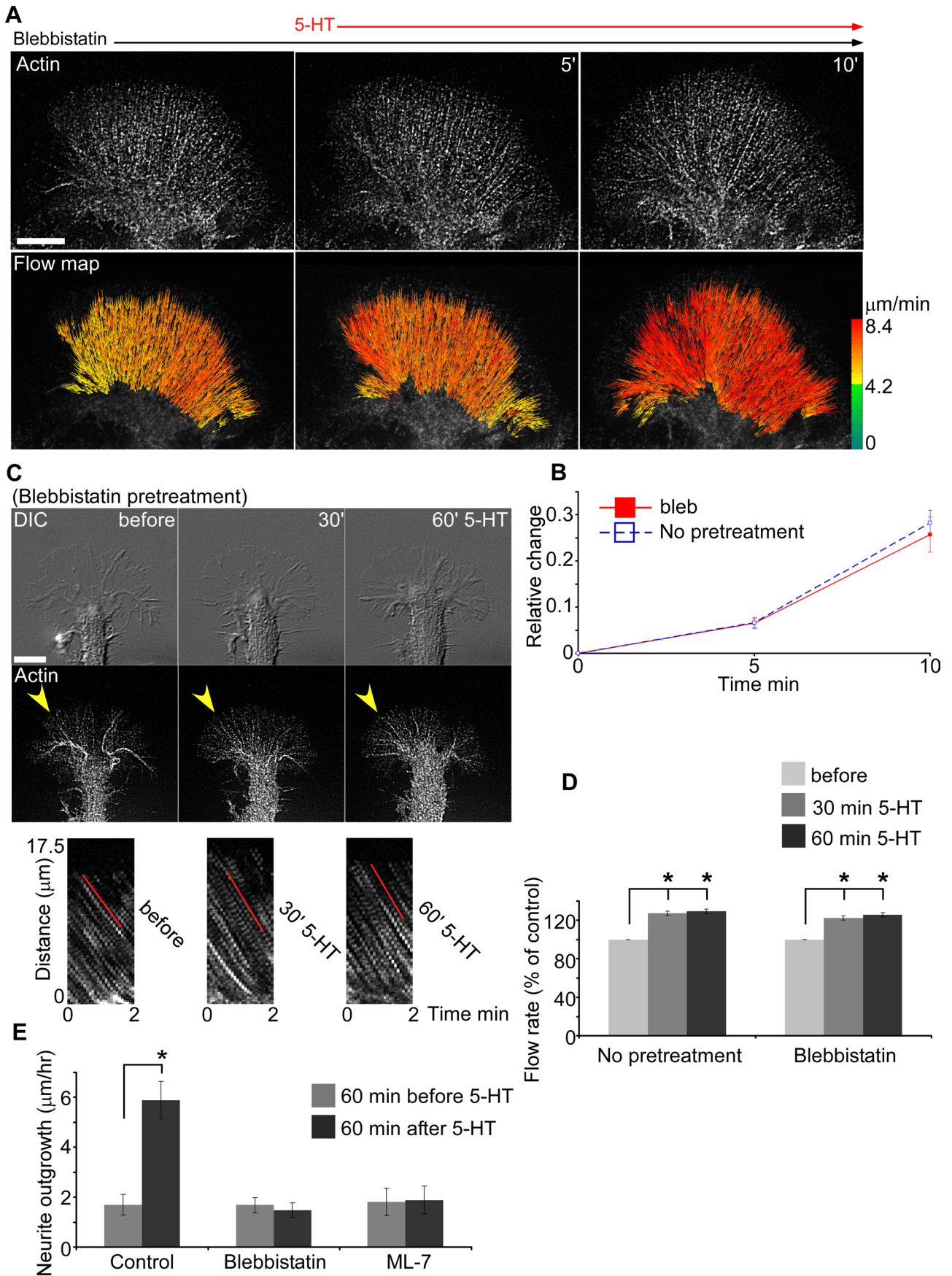
release (Supplemental Figure S6 and Figure 7, A and B); however, increases in F-actin flow rate were completely suppressed (Figure 7, A and C). These results strongly suggest that 5-HT-induced increases in F-actin flow depend on calcineurin activation downstream of Ca^{2+} release from intracellular stores.

To investigate the role of calcineurin in 5-HT-evoked neurite outgrowth, we assessed growth rates for 1 h in the presence of CN-AIP, FK-506, or CsA alone and after 5-HT exposure. Calcineurin inhibition had no effect on basal neurite outgrowth rates but completely blocked 5-HT-dependent neurite outgrowth (Figure 7D; see also DIC in Supplemental Figure S6). Taken together, these results support a mechanism by which 5-HT treatment promotes neurite outgrowth through a calcineurin-dependent increase in actin network flow and turnover.

5-HT-dependent calcineurin activation increases apCofilin1 activity

Cofilin is highly expressed in neuronal growth cones, and increased cofilin activity has been implicated in neurite extension (Meberg and Bamberg, 2000; Endo *et al.*, 2003; Ng and Luo, 2004; Tahirovic and Bradke, 2009). Cofilin is inhibited by phosphorylation by LIM kinase and activated by the cofilin phosphatase Slingshot (DesMarais *et al.*, 2005). It has been reported that intracellular Ca^{2+} elevation leads to calcineurin-dependent Slingshot activation and cofilin dephosphorylation (Wang *et al.*, 2005). Therefore we investigated a potential link between calcineurin and cofilin activity in 5-HT-induced neurite outgrowth. To generate cofilin activity probes, we identified and cloned two *Aplysia californica* cofilin homologues (apCofilin1 and apCofilin2, which share only 25.1% sequence identity). Antibodies were generated using recombinant proteins encoding both full-length apCofilins. We successfully generated phospho-specific antibodies against apCofilin1 and used this antibody in combination with total anti-apCofilin1 to assess apCofilin1 activity patterns (see Table 3 and Supplemental Figures S7 and S8).

To assess 5-HT effects on the spatial profile of apCofilin1 activity, we generated ratiometric P-apCofilin1/Total-apCofilin1 images. 5-HT treatments significantly increased, that is, disinhibited apCofilin1 activity in the entire growth cone, reflected by a decrease in measured P-apCofilin1/Total-apCofilin1 levels (Figure 8, A and B). Pretreatment with the calcineurin inhibitor FK-506 completely



abolished 5-HT-dependent cofilin activation, resulting in activity patterns very similar to those of controls (Figure 8, A, right, and C). Population analysis demonstrated that 5-HT treatment markedly increased cofilin activity in the distal one-third and the proximal one-third of the P domain by ~15 and ~22%, respectively, compared with controls (Figure 8D). However, in FK-506 backgrounds, 5-HT treatment did not cause any significant changes in cofilin activity (Figure 8D). Treatment with FK-506 alone did not significantly change apCofilin1 activity. Taken together, these observations suggest that 5-HT triggers calcineurin-dependent apCofilin1 activation, which supports increased rates of growth cone and neurite advance.

DISCUSSION

In *Aplysia* neurons, 5-HT induces actin polymerization essential for synaptic remodeling associated with long-term facilitation (Hatada et al., 2000; Udo et al., 2005). Facilitation also depends on Ca^{2+} release from postsynaptic stores (Li et al., 2005; Cai et al., 2008). These findings suggest roles for Ca^{2+} release and actin dynamics in synaptic plasticity; however, mechanisms by which these processes contribute to 5-HT-dependent neurite outgrowth are not well understood. Here we describe a novel mechanism of neurite growth by which exposure to a soluble factor, 5-HT, triggers Ca^{2+} release from intracellular stores, which in turn promotes increased rates of retrograde actin network flow accompanied by calcineurin-dependent apCofilin1 activation (Figure 9A).

Calcineurin, or PP2B, is a Ca^{2+} /calmodulin-dependent serine-threonine phosphatase, which plays a role in coupling Ca^{2+} signals to many neuronal responses (Groth et al., 2003; Nguyen and Di Giovanni, 2008; Bodmer et al., 2011) and has been implicated in promoting neurite outgrowth (Ferreira et al., 1993; Chang et al., 1995; Sotogaku et al., 2007; Arie et al., 2009; Figge et al., 2011). We found that 5-HT continued to evoke Ca^{2+} release after calcineurin inhibition; however, accompanying increases in F-actin flow and neurite outgrowth were completely absent (Figure 7). Our results indicated that calcineurin activation was necessary for the observed changes in actin dynamics and raised the question of the calcineurin effector. Calcineurin-dependent activation of Slingshot phosphatase can activate cofilin (Wang et al., 2005; Pandey et al., 2007; Zhao et al., 2008). In agreement, we found that 5-HT treatments resulted in calcineurin-dependent activation of apCofilin1 in regions in which increased F-actin flow were observed (Figure 8).

The cytoskeletal mechanism of this 5-HT growth response contrasts with what was observed during acute transitions from nonpermissive to permissive extracellular growth substrates (Lin and Forscher, 1995) or after application of apCAM-coated beads, for which increased rates of advance were correlated with decreased retrograde actin flow rates (Suter and Forscher, 2000, 2001). The rate of retrograde flow (V_r) in the P domain is determined by the balance of forces on actin networks (Craig et al., 2012). Actin network assembly near the leading edge and nonmuscle myosin II contractility in the T zone generate pushing and pulling forces, f_{poly} and f_{motor} , respectively, which drive network flow (Figure 9B, red arrows; Lin et al., 1996; Henson et al., 1999; Mogilner and Oster, 2003; Medeiros et al., 2006). Constant actin polymer turnover is necessary to prevent buildup of compressive forces in the T zone (Figure 9B; f_{break}), which resist actin filament flow (Van Goor et al., 2012). In addition to these internal forces, adhesion to extracellular substrates can generate traction forces (Figure 9B; $f_{adhesion}$), which tend to oppose retrograde flow (Lin and Forscher, 1995) and are the basis of the "molecular clutch hypothesis" for regulation of neurite growth (Mitchison and Kirschner, 1988; Suter and Forscher, 1998; Suter and Forscher, 2001; Schaefer et al., 2008). Adhesion and network compression can be modeled as viscous drags ($\epsilon_{adhesion}$ and ϵ_{break} , respectively), which result in forces that scale with actin flow velocity. In summary, actin polymerization and myosin II contractility tend to increase retrograde flow rates, whereas network compression in the T zone and/or adhesion to extracellular substrates tend to decrease it (Figure 9C; see Craig et al., 2012).

We found that 5-HT treatments (or PLC activation) resulted in acceleration of V_r without a significant change in P domain width. This means that increases in actin assembly had to be matched by increases in filament turnover. In line with this finding, recent related studies from our group support a key role for actin filament turnover in determining retrograde flow rates and P domain geometry (Van Goor et al., 2012; Yang et al., 2012). The importance of F-actin turnover for axon extension has also been reported (Bradke and Dotti, 1999; Gallo et al., 2002).

Cofilin activation promotes actin filament turnover, thereby reducing network density and decreasing ϵ_{break} , which would support the faster F-actin flow rates observed. Moreover, actin filament disassembly tends to increase G-actin concentration. Together these processes would increase polymerization rates and facilitate the accelerated actin treadmill rates observed. In addition,

FIGURE 6: The 5-HT-induced increase in actin flow is independent of myosin II activity. (A–C) Cells were pretreated for 20 min with blebbistatin (60 μ M) and drug maintained throughout experiments. (A) Alexa 594-phalloidin FSM (top) and corresponding flow map (bottom) from a growth cone before and after 5 and 10 min in 5-HT. Bar, 10 μ m. Flow map generated as described. (B) Summary of changes in retrograde F-actin flow rates in response to 5-HT. Images acquired every 5 s with 1-min elapsed recording time. No pretreatment (control) from Figure 2B is shown for comparison. $N = 3$ growth cones for each condition. Values are mean \pm SEM. (C) DIC (top) and phalloidin–Alexa 594 FSM (middle) of a blebbistatin-treated growth cone before and after 30 and 60 min in 5-HT. Bar, 10 μ m. Images were acquired every 5 s with 2-min elapsed recording time. Bottom, kymographs sampled across the P domain near yellow arrowheads showing rates of retrograde flow before and after 30 and 60 min of 5-HT exposure. Before, 4.55 ± 0.25 ; 30-min 5-HT, 5.55 ± 0.27 ; 60-min 5-HT, $5.69 \pm 0.25 \mu\text{m min}^{-1}$ (mean \pm SD, five measurements each). (D) Summary of normalized P domain retrograde flow rates in response to 10 μ M 5-HT with or without myosin II inhibition. Data normalized to rates before 5-HT addition. No pretreatment control: $N = 25$ growth cones from Figure 2D shown; blebbistatin ($N = 22$, 60 μ M pretreatment for 10–20 min). * $P < 0.001$ vs. before 5-HT addition. (E) Summary of neurite outgrowth sampled 60 min before and 60 min after 5-HT addition. Control conditions ($N = 58$ growth cones), blebbistatin pretreatment (60 μ M, 10 min, and presence throughout, $N = 36$); ML-7 pretreatment (10 μ M, 60 min, and presence throughout, $N = 34$). Control is from Figure 1C for comparison. In blebbistatin background, data were excluded from growth cones undergoing branching. Values are mean \pm SEM. Statistical analysis by two-tailed paired t test. * $p < 0.001$.

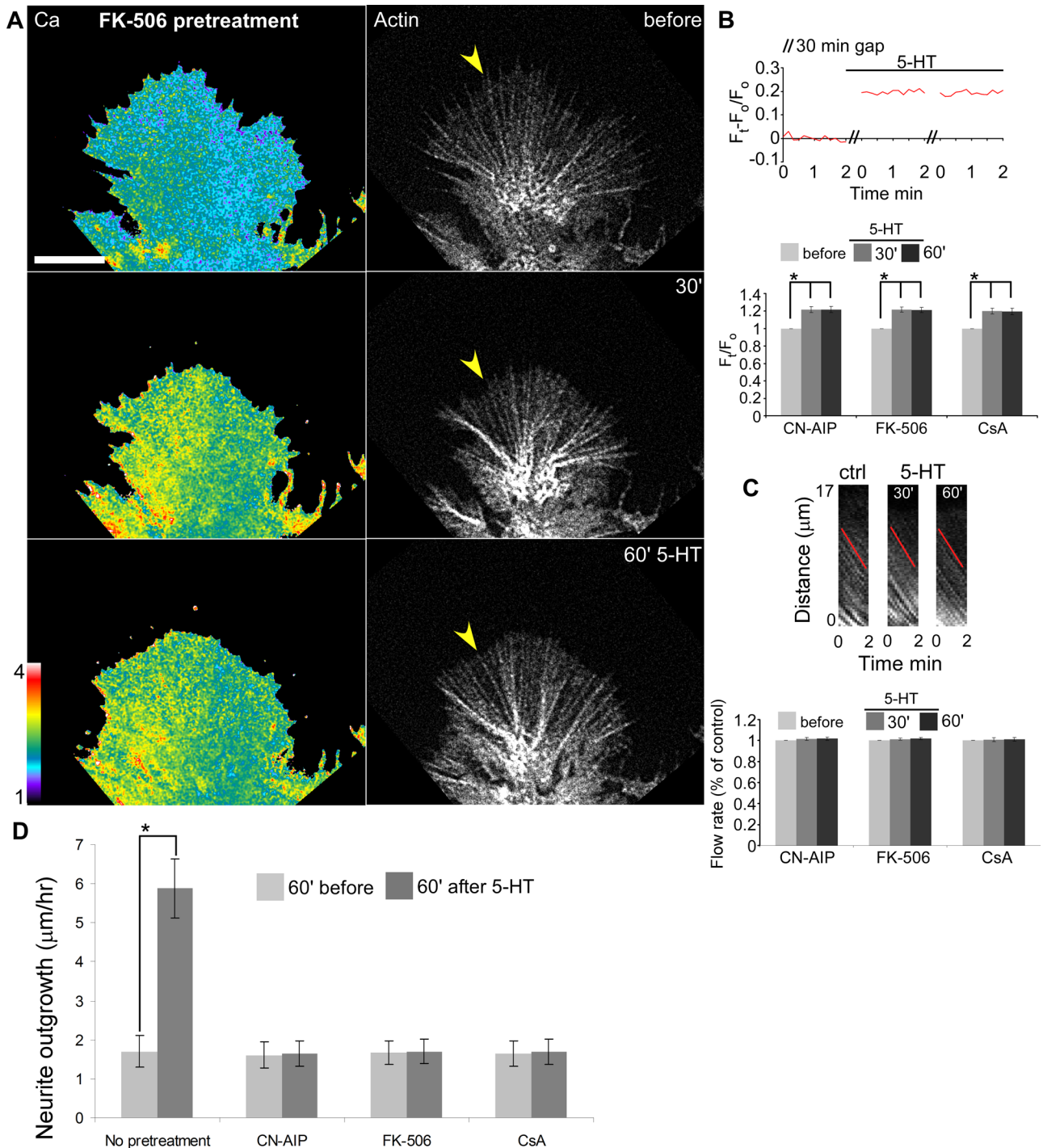


FIGURE 7: The 5-HT-induced increases in actin flow rates and neurite outgrowth depend on calcineurin activation downstream of Ca^{2+} release. (A) The Ca^{2+} ratio image (left) and phalloidin–Alexa 594 FSM (right) of a growth cone before and after 30 and 60 min in 5-HT. The cell was pretreated with FK-506 (2.5 μM) for 30 min. Note that FK-506 was present throughout. Bar, 10 μm . The Ca^{2+} ratio image is coded by pseudocolors in the linear scale (see scale bar). Data were acquired every 10 s with 2-min elapsed recording time. (B) Top, plot of Ca^{2+} response to 5-HT. Data obtained from the entire growth cone in A. F_0 , the average Ca^{2+} level before 5-HT addition. (B) Bottom, summary of F_t/F_0 plot recorded in the entire growth cone area quantifying Ca^{2+} response to 5-HT after calcineurin inhibition. Data were acquired every 10 s with 2- to 3-min elapsed recording time. F_0 , the average Ca^{2+} level before 5-HT addition. Number of growth cones tested ($N = 6$ (CN-AIP, injected, 0.2 mM in needle), 8 (FK-506, 2.5 μM pretreatment for 30 min and presence throughout), or 6 (CsA, 1 μM pretreatment for 30 min and presence throughout). Values are expressed as mean \pm SEM. Statistical analysis was done by two-tailed paired t test. $*P < 0.005$. (C) Top, kymographs created from area of interest as indicated in A (yellow arrowhead), showing rates of P domain retrograde flow before and after 30 and 60 min of 5-HT exposure. Before, 3.22 ± 0.23 , vs. 30-min 5-HT, 3.24 ± 0.24 , vs. 60-min 5-HT, $3.21 \pm 0.26 \mu\text{m min}^{-1}$ (mean \pm SD, five measurements). (C) Bottom, summary of normalized P domain retrograde flow rates in response to 5-HT after

Name	Sequence
apCofilin1	MASGKIADTVKEVYSRISLNSVKQTKLKYGVFK-FADDGASIVVETTATNADAMSDELISGLPKD-DVRYIAYDFDLSKDNVKTSEIVLVSWAPEKSAI-KRKMMCASTFNSLKMALAVSKNVLQGDSFDEVDV-VAALDKVGGKPLP
NP1	Ac-AMASGKIADTV-Camide
PP1	Ac-AMA(pS)GIKIADTV-Camide

ApCofilin1 contains 147 amino acids and has a predicted molecular weight of 15.985 kDa. Homologue-specific antibody to apCofilin1 was generated against the full-length protein of apCofilin1; phospho-specific anti-apCofilin1 antibody was raised against the first 11 amino acids of apCofilin1. Ac, acetyl group added to remove positive charge of the amine group; Camide, amide group added to remove negative charge of the carboxylic acid group; pS, phosphorylated serine.

TABLE 3: The full-length apCofilin1 protein sequence and peptide sequences used for homologue-specific antibody and phosphospecific antibody generation.

increased cofilin activity leads to barbed-end production, which increases the density of actin assembly sites and promotes leading-edge protrusion during cell migration in response to soluble growth factors in nonneuronal cells (Pollard and Borisy, 2003; Ghosh *et al.*, 2004; Kiuchi *et al.*, 2007) and to neurotrophic factors (BDNF, NGF, or netrin-1) in embryonic DRG and retinal neuron growth cones (Gehler *et al.*, 2004; Marsick *et al.*, 2010). In line with these findings, we recently reported a high density of free barbed ends in a band along the leading edge of bag cell neuron growth cones that were sensitive to treatments that inhibit cofilin activity (Van Goor *et al.*, 2012).

How can these observations be reconciled with the original molecular clutch hypothesis, which predicts that accelerated neurite outgrowth is correlated with decreased rates of retrograde actin flow? As per the foregoing discussion, retrograde flow rates depend on four parameters: 1) actin filament assembly, 2) myosin II contractility, 3) actin filament severing/recycling, and 4) cell adhesion (Figure 9, B and C). Assembly and turnover facilitate retrograde flow, whereas network compression and adhesion tend to oppose it. The present results suggest that a revision of the original molecular clutch hypothesis is in order. In particular, a less constrained multi-level slip clutch model as suggested by Giannone *et al.* (2009) can explain the results we observed. Indeed, with a viscous slip clutch-type model, faster flow rates *alone* without a change in adhesion strength ($\epsilon_{\text{adhesion}}$) would result in increased traction force due to increased viscous drag (Figure 9C). Alternatively, $\epsilon_{\text{adhesion}}$ could increase, but the resulting increase in drag force (f_{adhesion}) would have to be matched by increased polymerization and/or motor force to account for the flow rate acceleration observed. It is also possible that flow rates are increasing in 5-HT as a result of decreased $\epsilon_{\text{adhesion}}$. It will be of interest to measure traction forces in parallel with actin dynamics to address these outstanding issues. It should

be noted that the present results are not without precedent: immune system dendritic cells have been shown to migrate under conditions of extremely low adhesion by increasing their retrograde actin flow rates (Renkawitz *et al.*, 2009).

Myosin II activity can also promote actin turnover in growth cones (Medeiros *et al.*, 2006) and nonneuronal cells (Guha *et al.*, 2005; Murthy and Wadsworth, 2005; Haviv *et al.*, 2008). Here we investigated potential roles for myosin light-chain kinase and myosin II activity in 5-HT growth responses. Of interest, 5-HT-dependent increases in network turnover and retrograde flow persisted even after MLCK inhibition or direct inhibition of myosin II (Figures 5 and 6, A–D, and Table 2). In contrast, neurite outgrowth depended on myosin II activity (Figures 5, 6E, and 9), as previously observed in vertebrate neurons (Bridgman *et al.*, 2001; Turney and Bridgman, 2005). Myosin II activity might regulate adhesion site dynamics and maturation (Papusheva and Heisenberg, 2010) involved in generating traction forces that promote neurite extension (Zheng *et al.*, 1991; Heidemann and Buxbaum, 1994; Heidemann *et al.*, 1995). Although our results are consistent with myosin II inhibition reducing point contact consolidation (Woo and Gomez, 2006), further studies involving measurement of growth cone traction force are necessary to address this outstanding issue.

MATERIALS AND METHODS

Cell culture and chemicals

Primary culture of *Aplysia* bag cell neurons was as previously described (Forscher *et al.*, 1987). Coverslips were pretreated by 20 $\mu\text{g}/\text{ml}$ poly-L-lysine (Sigma-Aldrich, St. Louis, MO) for 15 min, then incubated in a 50 $\mu\text{g}/\text{ml}$ laminin (Sigma-Aldrich) solution for 2 h and rinsed in L15-ASW. 5-HT, U-73122, xestospongion C (XeC), Gly-Arg-Gly-Asp-Ser (RGD), and Ser-Asp-Gly-Arg-Gly (DGR) were from Sigma-Aldrich. Blebbistatin, ML-7, calcineurin autoinhibitory peptide (CN-AIP), FK-560, and CsA were from Calbiochem (La Jolla, CA). Calcium green-1 dextran, potassium salt, 10,000 MW (CG-1), Alexa Fluor 568 dextran, 10,000 MW (Alexa 568), Alexa Fluor 647 dextran, 10,000 MW (Alexa 647), Alexa Fluor 488 dextran, 10,000 MW (Alexa 488), Alexa Fluor 568–rabbit skeletal muscle G-actin (Alex 568 G-actin), and Alex Fluor 594 phalloidin were purchased from Molecular Probes (Eugene, OR).

Solutions

Artificial seawater (Na-ASW) contained (in mM) 400 NaCl, 10 KCl, 15 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES), 10 CaCl_2 , and 55 MgCl_2 at pH 7.8. Na-AWS was supplemented with 3 mg/ml bovine serum albumin (BSA), 0.5 mM vitamin E (Sigma-Aldrich), and 1 mg/ml carnosine (Sigma-Aldrich) before experiments. The Ca^{2+} injection buffer consisted of (in mM) 100 potassium aspartate and 10 HEPES at pH 7.4.

Microinjection

Microinjection protocol as described previously (Lin and Forscher, 1995). For Ca^{2+} imaging, neurons were injected with CG-1 and Alexa 647 or Alexa 568 in Ca^{2+} injection buffer (needle concentration

calcineurin inhibition. Data normalized to rates before 5-HT addition. Data were acquired every 5 or 10 s with 2- to 3-min elapsed recording time. Number of growth cones tested ($N = 10$ (CN-AIP, injected, 0.2 mM in needle), 11 (FK-506, 2.5 μM pretreatment for 30 min and presence throughout), or 9 (CsA 1 μM pretreatment for 30 min and presence throughout). Values are expressed as mean \pm SEM. (D) Summary of neurite outgrowth 1 h before and after 5-HT addition in various conditions. No pretreatment (control) from Figure 1C is shown for comparison. Control ($N = 58$), CN-AIP ($N = 45$), FK-506 ($N = 52$), and CsA ($N = 50$). Values are expressed as mean \pm SEM. Statistical analysis was done by two-tailed paired t test. * $p < 0.001$.

A Phosphorylated / Total Cofilin

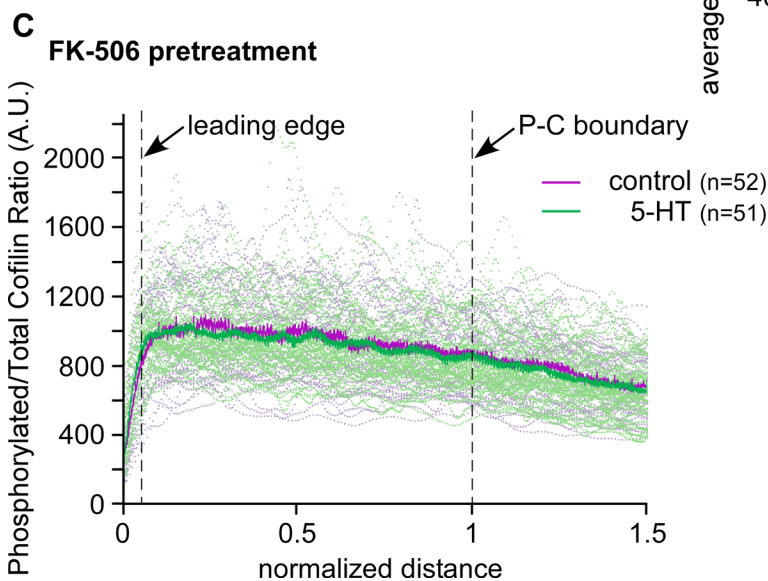
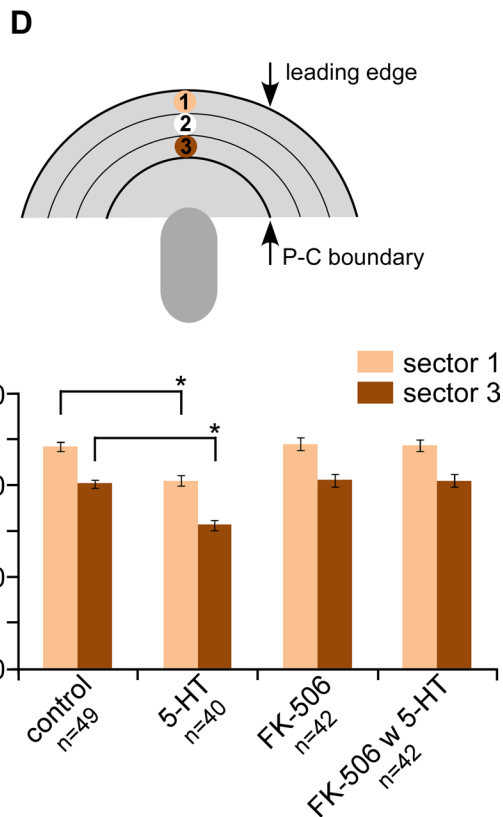
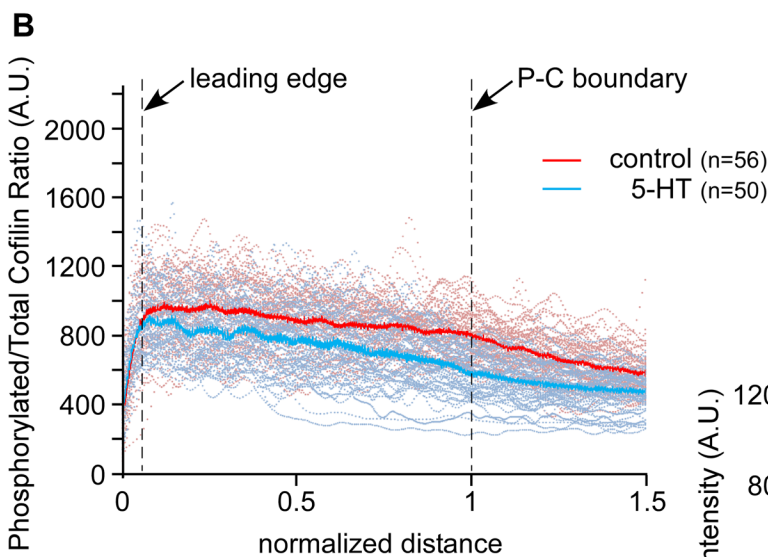
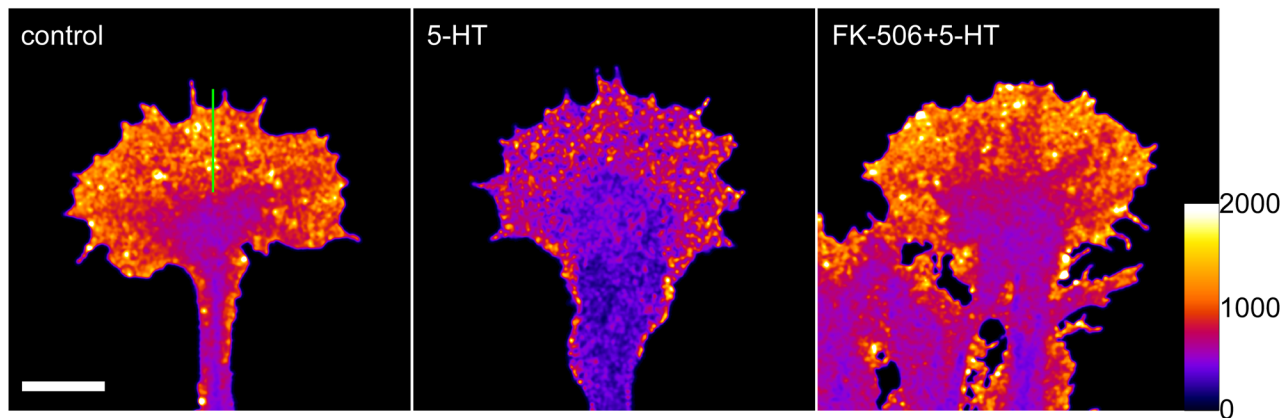


FIGURE 8: 5-HT increases calcineurin-dependent cofilin activity. (A) Ratio images of phosphorylated vs. total cofilin of growth cones. Cells were treated with vehicle (Na-ASW, left), 5-HT (10 μ M, middle) for 30 min or pretreated with FK-506 (2.5 μ M) for 30 min, followed by 30 min in 5-HT (10 μ M) with the continuous presence of FK-506 (right). Dual labeling of total and phosphorylated apCofilin1 was assessed with R- α -apCofilin1 (1:1000) and Sh- α -P-apCofilin1 (1:100) primary antibodies and Alexa 488 D- α -R (1:100) and Alexa 594 D- α -Sh (1:100) secondary antibodies. Bar, 10 μ m. Ratio image is coded by pseudocolors in the linear scale (see scale bar). (B, C) Line scan analysis of the ratio of phosphorylated vs. total

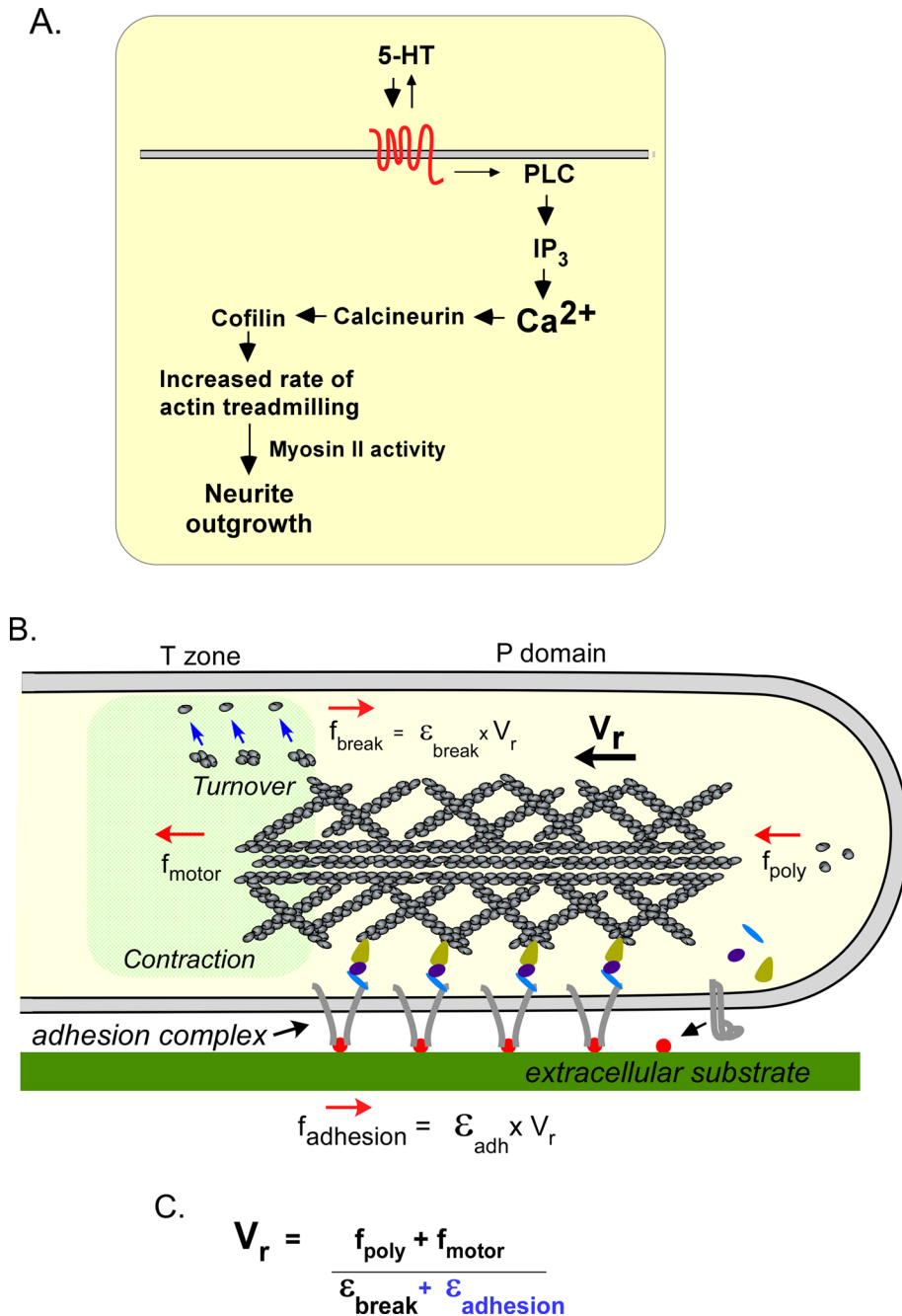


FIGURE 9: Model. (A) Growth on a laminin substrate elevates basal Rac activity levels and supports 5-HT-evoked Ca^{2+} release from IP_3 -gated internal stores. Ca^{2+} release leads to calcineurin-dependent cofilin activation, with increased actin filament turnover promoting faster network treadmilling. Accelerated network treadmilling can occur independent of myosin II activity; however, 5-HT-dependent increases in neurite outgrowth need myosin II activity. (B, C) Schematic and formula showing components that can define retrograde actin flow rate.

10–15 mg/ml for CG-1 and 0.8–0.9 mg/ml for volume tracer). For actin dynamics, neurons were injected with Alex 568–G-actin (needle concentration, 0.4 mg/ml) or Alexa Fluor 594–phalloidin

outside the cell. The ratio images (CG-1/volume) were then created by dividing background-corrected intensity values of CG-1 fluorescence by volume fluorescence and converted into time-lapse

(needle concentration, 20 μ M). For simultaneous Ca^{2+} imaging and actin dynamics, neurons were injected with CG-1, Alexa 647, and Alexa Fluor 594–phalloidin (needle concentration, 11.5 mg/ml, 0.9 mg/ml, and 38 μ M, respectively). Reagent or vehicle solution injections were typically ~10% of cell volume. After microinjection, cells were incubated in Na-ASW 1 h before imaging.

Confocal microscopy

Images were acquired using an Andor Revolution XD spinning disk confocal system (Andor, Belfast, United Kingdom) with a CSU-X1 confocal head (Yokogawa, Tokyo, Japan) and mounted on a Nikon TE 2000E inverted microscope with Perfect Focus (Nikon, Melville, NY). Confocal images were acquired using an Andor iXon^{EM+} 888 electron-multiplying charge-coupled device (CCD) camera. Transillumination was provided by a halogen lamp and controlled by a SmartShutter (Sutter Instrument, Novato, CA). Confocal excitation was provided by an Andor Laser Combiner with three laser lines at 488, 561, and 647 nm. Emission wavelength was controlled using a Sutter LB10W-2800 filter wheel outfitted with bandpass filters from Chroma Technology (Bellows Falls, VT). Image acquisition and all other peripherals were controlled by iQ software (Andor). A Nikon CFI Plan Apo 100 \times /1.4 numerical aperture (NA) objective was used.

Ca^{2+} imaging and analysis

Fluorescence images of growth cones loaded with the Ca^{2+} dye CG-1 and volume tracer were obtained using the described Andor confocal microscope. Paired images with comparable intensities of CG-1 and Alexa 568 or Alexa 647 were recorded every 10 s using 300- to 500-ms integration times for Ca^{2+} signal (488-nm laser line) and 200- to 300-ms integration times for the volume signal (561- or 647-nm laser line). The emission fluorescence filters used (denoted as center wavelength/bandwidth) were 535/40, 605/40, and 700/40 nm, respectively (Chroma Technology). For each paired image, Gaussian convolution was used to reduce noise levels, and a binary mask was also used to eliminate noise amplification

apCofilin1 fluorescence in growth cones. Scattered dots represent data set from individual growth cones. Solid lines represent the population average. *N*, growth cones measured. (B) Comparison between cells treated with vehicle (control, red) and cells treated with 5-HT (10 μ M, 30 min, blue). (C) In FK-506 backgrounds (2.5 μ M FK-506 pretreatment for 30 min), comparison between cells treated with vehicle (control, purple) and cells treated with 5-HT (10 μ M, 30 min, green) in the continuous presence of FK-506. (D) Quantification of average ratio of phosphorylated vs. total cofilin in sectors 1 and 3 (see inset for schematic) for each condition in B and C. *N*, growth cones measured. **p* < 0.001 with two-tailed unpaired *t* test.

montages for data analysis as reported previously (Zhang and Forscher, 2009). Average pixel intensity values were obtained from the entire area of the growth cones of interest. The Ca^{2+} changes over time were expressed as $\Delta F/F_0$, where $\Delta F = F_t - F_0$ and F_0 is the average Ca^{2+} level sampled during the 3- to 5-min baseline period (before 5-HT addition). $\Delta F/F_0$ (%) levels of >10% are considered significant.

Quantification of actin dynamics by fluorescent speckle microscopy

Two methods were used to visualize F-actin for FSM: 1) fluorescently labeled G-actin incorporated into actin filaments and 2) low levels of fluorescent phalloidin, which specifically binds F-actin but not G-actin. Briefly, two-channel images were acquired using 500- to 900-ms integration times for actin fluorescent probe and 120 ms for DIC with 5- or 10-s intervals as previously described (Burnette *et al.*, 2007). Kymography and automated speckle tracking were used to determine rates of F-actin movement. For kymographs, analysis was done as reported (Zhang *et al.*, 2003). For automated flow tracking, an adaptive multiframe correlation algorithm was performed, as described (Ji and Danuser, 2005). For image presentation only, the contrast of F-actin FSM images was enhanced by processing with an unsharp mask, followed by low-pass spatial filters.

Tandem Ca^{2+} ratio and F-actin FSM imaging

Ratiometric Ca^{2+} imaging was used to measure Ca^{2+} levels in the growth cone at the same time as F-actin dynamics was assessed using the methods described. Cells were injected with CG-1, Alexa 647-dextran, and Alex Fluor-594 phalloidin. DIC, Ca^{2+} level, volume signal, and F-actin were recorded in tandem with 5- or 10-s sampling intervals. CG-1, Alex Fluor 594-phalloidin, and Alexa 647-dextran were excited simultaneously with 488-, 561-, and 647-nm laser lines and emission monitored using 535/40-, 605/40-, and 700/40-nm filters, respectively (Chroma Technology). The F-actin dynamics was quantitatively compared and contrasted with corresponding Ca^{2+} levels in the growth cone.

Neurite outgrowth analysis

For long-term time-lapse experiments a Zeiss Axiovert 10 microscope with phase contrast optics (10× AchroStigmat/NA 0.25) and a CoolSnapHQ (Photometrics, Tucson, AZ) cooled CCD camera were used. Hardware and image acquisition were controlled with the open-source μ -Manager device adapter library (www.micromanager.org) through a custom Java user interface. To quantify growth cone advance 1 h before and after 5-HT addition in culture, a Nikon Eclipse TE300 microscope equipped with a Photometrics Quantix 57 backilluminated cooled CCD camera and MetaMorph instrumentation control software were used (Molecular Devices, Sunnyvale, CA). The displacement of growth cone's leading edge along the presumed growth axis in 1 h was used to depict neurite outgrowth.

Cloning *A. californica* cofilin1 (apCofilin1) and *A. californica* cofilin2 (apCofilin2)

Primers (apCofilin1_{for} and apCofilin1_{rev}; apCofilin2_{for} and apCofilin2_{rev}) were designed to amplify apCofilin1 and apCofilin2 from cDNA while introducing *Nco*I sites at the initiation codon and *Bam*HI restriction sites after the stop codon. Extra nucleotides were included outside the restriction sites to allow efficient digestion of PCR products. PCR was performed using Herculase polymerase (Stratagene, Santa Clara, CA). A cDNA library was constructed using mRNA extracted from bag cell neurons. PCR products were

separated on 1% agarose gels, and bands of the correct size (~450 base pairs) were cut out and purified with the QIAquick Gel Extraction Kit (Qiagen, Valencia, CA). Eluted DNA was digested with *Bam*HI and *Nco*I restriction enzymes (New England BioLabs, Ipswich, MA), ligated into pET15b (Novagen, Gibbstown, NJ) vector digested with the same enzymes, and transformed into DH5 α bacteria (Invitrogen) for amplification. Plasmids were confirmed by the Keck DNA Sequencing Facility (Yale University, New Haven, CT).

Antibody generation

Bacterially expressed, recombinant apCofilin1 and apCofilin2 were purified and sent to Proteintech Group (Chicago, IL) for generation of antibodies. apCofilin1 and apCofilin2 antibodies were generated in a rabbit or a guinea pig host against full-length apCofilin1 and apCofilin2, respectively, and were used without affinity purification. Generation of phospho-specific anti-apCofilin1 was performed by 21st Century Biochemicals (Marlboro, MA) from phosphorylated peptide corresponding to the first 11 amino acids of apCofilin1 (PP1). The antibody was generated in a sheep host, affinity depleted against the nonphosphorylated peptide (NP1), and affinity purified with the PP1 peptide. An attempt to generate a phospho-specific anti-apCofilin2 antibody against the first 10 amino acids of apCofilin2 was unsuccessful.

Western blots

Western blots were performed using standard methodology. Protein samples were resolved by SDS-PAGE, transferred to nitrocellulose membranes (Scheicher & Schuell BioScience, Dassel, Germany) by semidry transfer (TransBlot SD; Bio-Rad, Hercules, CA), probed with the indicated primary and secondary antibodies, developed with SuperSignal West Pico Chemiluminescent Substrate (Pierce, Rockford, IL), and digitally exposed using the Epi Chemi II Darkroom (UVP Laboratory Products, Upland, CA). For antigen competition assays, the primary antibody was preincubated with excess antigen for 30 min at 4°C before use; the control was preincubated with buffer. For alkaline phosphatase treatment, the nitrocellulose membranes were incubated in CIP buffer (50 mM Tris-HCl, pH 7.9, 100 mM NaCl, 10 mM MgCl₂, 1 mM dithiothreitol) at 37°C for 60 min, with or without 25 U/ml calf intestinal phosphatase (CIP; New England BioLabs). The membranes were washed extensively in TBS-T (50 mM Tris, pH 7.5, 150 mM NaCl, 0.1% Tween-20) and then processed normally.

Immunocytochemistry

In a flow chamber, cells were incubated in Fix (4% Formalin, 400 mM NaCl, 10 mM KCl, 15 mM HEPES, pH 7.8, 10 mM CaCl₂, 55 mM MgCl₂, and 400 mM sucrose) for 20–30 min and 1% Triton X-100 in Fix for 30 min before three washes with PBS-T (0.1% Triton X-100, 137 mM NaCl, 2.7 mM KCl, 10 mM phosphate, pH 7.5). For antibody labeling, cells were blocked for 30 min in 5% BSA/PBS-T, incubated with primary antibody for 20–30 min in 5% BSA/PBS-T, washed three times in 5% BSA/PBS-T, and incubated for 30 min to 1 h in secondary antibody diluted in 5% BSA/PBS-T. For antigen competition assays, the diluted primary antibody was incubated with 100- to 500-fold excess antigen for 30 min at 4°C with rotation before use. For actin visualization, Alexa 594 or Alexa 488-phalloidin was included in the secondary antibody solution at 0.66 μ M. Cells were washed three times in PBS-T and mounted in Mowiol media.

Cofilin line scan analysis

Line scans of ratio images of background-corrected intensity values of phosphorylated cofilin divided by total cofilin were used to

analyze the spatial intensity distribution of phosphorylated (inactive) relative to total cofilin. A 50-pixel-wide line was drawn from the leading edge to 1.5× peripheral domain (P domain) width along the presumed growth axis. Average intensity was measured with the plot profile function in ImageJ (National Institutes of Health, Bethesda, MD) and the data exported to Excel (Microsoft, Redmond, WA). Intensity was plotted versus distance normalized by growth cone size for population analysis of line scans. Distance was normalized by setting the beginning (left end) of the lines scan at the leading edge and letting the two-thirds position be the peripheral-central domain interface. Alternatively, to compare phosphorylated cofilin normalized to total cofilin in P domain under different conditions, the peripheral domain was divided into three equal annular sectors parallel to the leading edge, and average intensities in sectors 1 and 3 were calculated (Figure 8D, inset) and compared between different conditions.

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