Rac1 Modulates Stimulus-evoked Ca2 Release in Neuronal Growth Cones via Parallel Effects on Microtubule/Endoplasmic Reticulum Dynamics and Reactive Oxygen Species Production

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The small G protein Rac regulates cytoskeletal protein dynamics in neuronal growth cones and has been implicated in axon growth, guidance, and branching. Intracellular Ca²⁺ is another well known regulator of growth cone function; however, effects of Rac activity on intracellular Ca²⁺ metabolism have not been well characterized. Here, we investigate how Rac1 activity affects release of Ca²⁺ from intracellular endoplasmic reticulum (ER) stores stimulated by application **of serotonin (5-hydroxytriptamine). We also address how Rac1 effects on microtubule assembly dynamics affect distribution of Ca2 release sites. Multimode fluorescent microscopy was used to correlate microtubule and ER behavior, and** ratiometric imaging was used to assess intracellular Ca^{2+} dynamics. We report that Rac1 activity both promotes Ca^{2+} **release and affects its spatial distribution in neuronal growth cones. The underlying mechanism involves synergistic Rac1** effects on microtubule assembly and reactive oxygen species (ROS) production. Rac1 activity modulates Ca^{2+} by 1) enhancing microtubule assembly which in turn promotes spread of the ER-based Ca²⁺ release machinery into the growth cone periphery, and 2) by increasing ROS production which facilitated inositol $1,4,5$ -trisphosphate-dependent $Ca²⁺$ **release. These results cast Rac1 as a key modulator of intracellular Ca2 function in the neuronal growth cone.**

INTRODUCTION

The small G protein Rac plays a pivotal role in establishing cell polarity and controlling cell migration (Etienne-Manneville and Hall, 2002; Jaffe and Hall, 2005). Although Rac was initially characterized as a regulator of actin network assembly in lamellipodia of motile cells (Hall, 1998), later studies demonstrated that Rac has remarkable functional diversity. For example, Rac activation promotes microtubule assembly in epithelial cells (Wittmann *et al*., 2003, 2004) and the Rac activator DOCK7 inactivates the microtubule-destabilizing protein stathmin/Op18 in nascent axons (Watabe-Uchida *et al*., 2006). In addition, Rac is an essential component of NADPH oxidase complexes that produce reactive oxygen species (ROS) used for oxidative killing by phagocytes and for cellular signaling by nonmyeloid cells (Lambeth, 2004; Bedard and Krause, 2007). In neurons, genetic approaches suggest Rac is required for axon outgrowth, guidance, and branching (Luo, 2000; Guan and Rao, 2003); however, the underlying cell biological mechanisms are not yet well characterized.

Intracellular Ca^{2+} is another well known regulator of growth cone motility (Henley and Poo, 2004; Gomez and Zheng, 2006); however, Ca^{2+} effects on growth cone behavior seem to be pleiotropic. For example, in some cases Ca^{2+} increases induce growth cone collapse and inhibit neurite outgrowth (Cohan and Kater, 1986; Cohan *et al*., 1987; Mattson

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and Kater, 1987; Gomez *et al*., 1995, 2001; Gomez and Spitzer, 1999), but in others Ca^{2+} increases are correlated with filopodium protrusion and neurite elongation (Connor, 1986; Bedlack *et al*., 1992; Garyantes and Regehr, 1992; Lau *et al*., 1999; Cheng *et al*., 2002). Several lines of evidence indicate that Ca^{2+} regulates cytoskeletal function via Rac activation (Fleming *et al*., 1999; Price *et al*., 2003; Jin *et al*., 2005); in contrast, the inverse scenario, i.e., how Rac activity affects $Ca²⁺$ metabolism has not been well characterized.

Here, we address this question in the context of Ca^{2+} responses stimulated by application of the neurotransmitter serotonin (5-hydroxytryptamine, 5-HT) to neuronal growth cones. Specifically, we investigated how Rac effects on microtubule assembly and/or ROS production affect stimulus evoked Ca^{2+} release. We report that Rac1 activity promoted microtubule assembly-dependent endoplasmic reticulum (ER) advance which in turn resulted in spread of ER based $Ca²⁺$ release sites to the growth cone periphery. In parallel with these effects, Rac1 activity promoted ROS production in growth cones which facilitated $1,4,5$ -trisphosphate (IP_3) -dependent Ca^{2+} release in response to 5-HT application.

MATERIALS AND METHODS

Cell Culture and Chemicals

Primary culture of *Aplysia* bag cell neurons as described previously (Forscher *et al*., 1987). 5-HT, U73122, U73343, Xestospongin C (XeC), ryanodine (Rya), Taxol, H₂O₂, and *N*-acetyl-cysteine (NAC) were from Sigma-Aldrich (St. Louis, MO). Heparin sodium salt, vinblastine, apocynin, genistein, and ionomycin were from Calbiochem (San Diego, CA). L61 Rac1-glutathione-S-transferase (GST) and N17 Rac1-GST, referred to here as "CA Rac1" and "DN Rac1", respectively, were from Cytoskeleton (Denver, CO), and biological activity of these constructs was confirmed by the manufacturer. Calcium Green-1 dextran, potassium salt, 3000 molecular weight (MW) (CG-1), Texas

Red dextran, 3000 MW (Texas Red), BODIPY FL brefeldin A and CM-H2DCFDA (DCF) were purchased from Invitrogen (Carlsbad, CA). Anti-KDEL antibody MAC 256 was from Abcam (Cambridge, MA).

Solutions

Artificial seawater (Na-ASW) contained 400 mM NaCl, 10 mM KCl, 15 mM HEPES, 10 mM CaCl₂, and 55 mM MgCl₂ at pH 7.8. Na-AWS 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. Ca^{2+} free ASW was as for Na-ASW except that CaCl₂ is omitted and EGTA (0.5 mM) is included. Ca²⁺ injection buffer consisted of 100 mM potassium aspartate and 10 mM 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 Texas Red in Ca^{2+} injection buffer (needle concentration 11–15 and 0.3–0.4 mg/ml, respectively). For microtubule dynamics, neurons were injected with 1 mg/ml Alexa-Fluor 594-labeled tubulin. To generate CA and DN Rac1 backgrounds: L61 Rac1- GST (CA Rac1) or N17 Rac1-GST (DN Rac1) was included in the injection needle (concentration 0.7–1.0 mg/ml) as described previously (Zhang *et al*., 2003), along with tubulin probes or Texas Red volume tracer with or without $Ca²⁺$ dye. For ROS assays, neurons were injected with 0.4 mg/ml Texas Red as a volume tracer. Reagent or vehicle solution injections were typically \sim 10% of cell volume. After microinjection, cells were incubated in Na-ASW 1 h before imaging.

Ca2 Imaging and Analysis

Fluorescence images of growth cones loaded with Ca^{2+} dye CG-1 and Texas Red volume tracer were obtained using an Eclipse TE300 microscope (Nikon, Tokyo, Japan) equipped with a Quantix 57 back-illuminated frame transfer cooled charge-coupled device camera (Photometrics, Tucson, AZ) mounted
on the bottom port. Similar Ca²⁺ profiles and dynamics were observed in control experiments using 70,000 MW CG-1 ruling out the possibility that the 3,000 MW CG-1 used here could be portioning into intracellular stores. Two programmable fast filter wheels (100-ms filter changes) that include shutters were mounted with bandpass filters (Ludl,) for changing both excitation and emission wavelengths. Fluorescence filters used (denoted as center wavelength/bandwidth) were as follows: CG-1: 484/15 nm for excitation, and 523/15 nm for emission; Texas Red: 555/25 nm for excitation and 605/25 nm for emission (Chroma Technology, Brattleboro, VT). Paired images with comparable intensities of CG-1 and Texas Red were recorded every 10 or 20 s by using 400- to 600-ms integration times for fluorescein isothiocyanate (FITC) channel (Ca²⁺ signal) and 100- to 200-ms integration times for fluorescein isothiocyanate (TRITC) channel (volume signal). The ratio images (CG-1/ Texas Red) were created by dividing background corrected intensity values of CG-1 fluorescence by Texas Red fluorescence and then converted into timelapse montages or kymograph for data analysis. Average pixel intensity values were obtained from the entire C-domain and P-domain (see Figure 1D) of the growth cones of interest. Ca^{2+} changes over time were expressed as $\Delta F/F_0$, where $\Delta F = F_t - F_0$ and F_0 was the average Ca²⁺ level sampled during ~3- to 5-min baseline period (before 5-HT addition). $\Delta F/F_o$ (%) greater than 10% was considered significant. Ten- or 20-s sampling intervals were used for $Ca²⁺$ imaging. This sampling regime would not resolve $Ca²⁺$ transients with 1-s time constants such as observed in *Xenopus* growth cones (Gomez *et al*., 2001) and thus does not address potential contributions of these events.

To assess basal Ca^{2+} levels in growth cones, ratiometric Ca^{2+} images were performed first in Na-ASW to get basal fluorescence ratio. Subsequently, the bath solution was exchanged with Ca^{2+} -free AWS with addition of various concentrations of CaCl₂ (~0.1–0.4 mM) to yield free Ca²⁺ concentration ranging from ~45 to 720 nM (WEBMAXCLITE, http://www.stanford.edu/~cpatton/webmaxc/ webmaxclite115.htm). Ionomycin (10 µM) was included to equilibrate extra-
cellular and intracellular Ca²⁺ levels. The fluorescence ratio acquired in each corresponding bath solution was normalized to the basal fluorescence ratio. The graph of normalized fluorescence ratio (percentage) plotted against free Ca^{2+} concentration exhibited a linear regression with $R^2 > 0.96$.

Actin and Microtubule Immunocytochemistry

Bag cell neurons were fixed by rapid exchange of the medium with 3.7% formaldehyde in ASW supplemented with 400 mM sucrose. After fixation for 20 min, the cells were permeabilized for 5 min using 1% Triton X-100 in the fixation solution. Cells were then washed three times with phosphate-buffered saline (PBS) containing 0.1% Triton X-100 (wash solution). For actin filament labeling, Alexa 594-phalloidin (Invitrogen) was incubated at 1:50 in wash solution for 1 h. After three washes, the cells were blocked with 5% BSA in wash solution for 30 min and then incubated with mouse anti- α -tubulin (Sigma-Aldrich) at 1:100 in blocking solution for 30 min at room temperature (RT). After three washes, Alexa 488-goat anti-mouse immunoglobulin G (IgG) (Invitrogen) was added at 1:100 in blocking solution for 30 min at RT. The final wash solution was replaced with anti-fading solution Mowiol (Calbiochem, San Diego, CA). Microtubules and actin filaments were visualized using the same filters as CG-1 and Texas Red as described above.

ER Immunocytochemistry

An anti-KDEL antibody was used to visualize ER structure (Munro and Pelham, 1987; Saunders and Cohen, 1999). Cells were fixed with 4% paraformaldehyde for 15 min, permeabilized with 0.5% Triton X-100 for 5 min, and then incubated with 5% BSA and 10% normal goat serum for 1 h. After blocking, cells were stained with rat anti-KDEL antibody (MAC 256; diluted 1:100) for 1 h and visualized with Alexa 594-goat anti-rat IgG (Invitrogen) for 30 min. ER distribution and actin filaments (labeled with Alexa 488-phalloidin) were visualized in growth cones by spinning disk confocal microscopy (Andor, Belfast, Northern Ireland). Alexa 488 and Alexa 594 were excited simultaneously with 488 and 561 nm laser lines, and emission was monitored using 535/40 nm and 625/40 nm filters respectively (Chroma Technology).

ER Vital Labeling

BODIPY FL brefeldin A was used for ER staining (Deng *et al*., 1995; Cifuentes *et al*., 2001). Neurons were labeled in Na-ASW containing BODIPY FL brefeldin A (2 μ M) in total darkness for 30 min. At the end of the incubation period, cells were extensively washed and mounted in a laminar flow chamber for observation. In most cases, neurons were injected with CA Rac1 or DN Rac1. For visualizing microtubule and ER simultaneously, microtubule probe was also injected beforehand. The fluorescence filters used for the ER and microtubule imaging were the same as for CG-1 and Texas Red as described above.

ROS Detection

CM-H2DCFDA (DCF) was used for ROS detection (Xie *et al*., 1999). Neurons injected with Texas Red were loaded with DCF (40 μ M) for 1.5 h in the dark in Na-ASW and then washed thoroughly. Fifteen minutes later the coverslip was affixed to a chamber for visualization. In most cases, neurons were also injected with CA Rac1 or DN Rac1. The filter settings for the volume corrected ROS imaging were the same as for CG-1 and Texas Red as described above. Paired images of DCF and Texas Red were recorded using 200-to 500-ms integration times for FITC channel (ROS signal) and 100- to 200-ms integration times for TRITC channel (volume signal). The path length corrected image was generated by dividing background subtracted pixel values of DCF fluorescence by those of Texas Red fluorescence. To verify the sensitivity of this assay, a line scan in ROS kymograph derived from DCF/Texas Red images was done during H_2O_2 (10 μ M) exposure. To quantify ROS levels, the values of DCF/Texas Red in C-domain and P-domain were measured in different Rac1 backgrounds and plotted as box-whisker plots.

Quantitative Analysis

Microtubule and ER dynamics assessed by fluorescent speckle microscopy (FSM) a technique for direct quantitative assessment of cytoskeletal polymer dynamics in living cells (Waterman-Storer *et al*., 1998; Waterman-Storer and Salmon, 1998b). Multimode time lapse microscopy as described previously (Schaefer *et al*., 2002). In brief, three-channel recordings were made every 12 s by using 300- to 700-ms integration times for microtubules, 100 to 400 ms for ER visualization, and 50 ms for differential interference contrast (DIC) images. Microtubule dynamic parameters were assessed as reported previously (Schaefer *et al*., 2002). To quantify ER density and advance, the P-domain was divided into quartiles starting at the C–P-domain interface and ending at the leading edge (see Figure 1D). Microtubule/ER excursions were quantified by counting the number of microtubules or ER tubules spanning 25 or 75% of the P-domain (denoted 25%P or 75%P, respectively). ER–microtubule-coupling efficiency was estimated by dividing ER by microtubule excursions assessed at the same time. Data are expressed as mean \pm SEM, unless stated otherwise. Statistical analysis was done with two tailed paired or unpaired *t* test.

RESULTS

Rac1 Promotes Microtubule–ER Advance into P-Domain

Given that Rac1 activity can promote microtubule assembly in nonneuronal cells (Wittmann *et al*., 2003), we first characterized microtubule dynamics in different Rac1 activity backgrounds by using FSM as described previously (Schaefer *et al*., 2002) (also see Supplemental Figure S1). Growth cones are composed of two distinct cytoplasmic domains: 1) the central or C-domain, characterized by fast organelle transport and high microtubule density; and 2) the peripheral or P-domain, consisting of dense dynamic actin networks that tend to exclude organelles larger than ${\sim}50$ nm (Figure 1D; Forscher *et al*., 1987).

Figure 1. Rac1 promotes microtubule–ER advance into P-domain. (A) Immunocytochemistry of actin filament and ER distribution in a growth cone in CA Rac1 backgrounds (left) or DN Rac1 backgrounds (right). Top, actin. Bottom, ER. Red line, filopodial ends. Bar, 10 μ m. (B) Simultaneous labeling of microtubule and ER in CA Rac1 injected neuron. From left to right, DIC, microtubule and ER. Bar, 10 μ m. (C) As in B in DN Rac1-injected neurons. (D) DIC image indicating C- and Pcytoplasmic domains and reference lines used in calculating ER and microtubule excursion numbers (see *Materials and Methods*). C, Cdomain; P, P-domain. Red line, interface between C-domain and P-domain defined by organelle boundary; green line, 25%P; blue line, 75%P. Bar, 10 μ m. (E) Quantification of microtubule versus ER excursions past 25%P and 75%P boundaries in control, CA Rac1, or DN Rac1. Top, microtubule excursion number; middle, ER excursion number; and bottom, ER—microtubule-coupling efficiency. Data between growth cones normalized by boundary length. $N = 10$ (control), 12 (CA Rac1), and 8 (DN Rac1) for microtubule excursion assay; $N = 16$ (control), 14 (CA Rac1), and 12 (DN Rac1) for ER excursion assay; and $N = 9$ (control), 10 (CA Rac1), and 8 (DN Rac1) for ER microtubule-coupling efficiency assay. N denotes the number of growth cones tested. Values are expressed as mean \pm SEM. Statistical analysis by two-tailed unpaired *t* test. $*p < 0.01$ versus control and #p < 0.05 versus control.

We found that constitutively active (CA) Rac1 increased microtubule rescue frequency by 160% and decreased catastrophe frequency by $-\frac{28}{6}$ in the P-domain (Table 1). Dominant-negative (DN) Rac1 had the opposite effect: decreasing microtubule rescue frequency and increasing microtubule catastrophe frequency in the P-domain by -32 and 200%, respectively (Table 1). As a result, microtubules tended to spend a greater proportion of the time growing and relatively less time shrinking in CA Rac1 compared with DN Rac1 backgrounds. Absolute microtubule growth and shortening velocities did not seem to be significantly affected by Rac1 activity. It is important that more persistent microtubule growth in the presence of CA Rac1 promoted deep microtubule advances into the P-domain, consistent with previous observations in epithelial cells (Wittmann *et al*., 2003).

We next examined the effect of Rac1 activity on ER distribution in the growth cones by using an anti-KDEL antibody to visualize ER distribution and Alexa-phalloidin to locate characteristic cytoplasmic domains in fixed cells (Saunders

Plus-end periphery microtubules: control (n = 25 microtubules), CA Rac1 (n = 31 microtubules), and DN Rac1 (n = 25 microtubules). Mean \pm SEM elapsed recording time: $3-5$ min. $np < 0.001$ vs. control.

Given that ER distribution is dynamically regulated by microtubule-based transport in neuronal growth cones (Dailey and Bridgman, 1989, 1991) and other motile cells (Terasaki *et al*., 1986; Waterman-Storer and Salmon, 1998a), we next covisualized microtubule and ER dynamics in the growth cone by FSM and BODIPY brefeldin-A imaging (Deng *et al*., 1995; Cifuentes *et al*., 2001). In CA Rac1 backgrounds, growth cones tended to have well spread C-domains with microtubules and ER both extending deep into the P-domain (Figure 1B and Supplemental Movie 1). In contrast, in DN Rac1 backgrounds, growth cones tended to have more compact, focused C-domains (Figure 1C). Although microtubules still sometimes extended into the P-domain, the ER did not enter the P-domain nearly as efficiently (Figure 1C and Supplemental Movie 2). Note that individual images tend to underestimate the time-averaged density of the ER in the Pdomain because peripheral ER is associated with a highly dynamic population of microtubules (Schaefer *et al*., 2002; Suter *et al.*, 2004).

To quantify these Rac effects, we counted microtubule and ER excursions past fiducial marks placed at 25 and 75% of

Figure 2. Rac1 alters ER dynamic behavior on microtubules. (A) Two representative examples of time-lapse sequences of microtubule (top) and ER (bottom) in CA Rac1 backgrounds. Red and green arrowheads point to microtubule and ER tips, respectively. Note that the red arrowheads in ER panel indicate the corresponding position of microtubule tips. Elapsed time (seconds) is shown at the top. Arrows denote position transition zone between Cand P-domains. (B) DN Rac1 background. Regions are similar to white box in Figure 1D.

the width of the P-domain under all conditions (Figure 1D, green 25%P, blue 75%P lines, respectively; see *Materials and Methods*). The number of microtubule and ER excursions past the 25%P and 75%P marks increased in CA Rac1 and decreased in DN Rac1 backgrounds relative to controls (Figure 1E, top and middle, respectively). We also calculated the ratio of ER to microtubule excursions to estimate ER–microtubule-coupling efficiency, and found that CA Rac1 significantly increased this ratio in both the proximal (25%) and distal (75%P) regions of the P-domain relative to controls, whereas DN Rac1 had the opposite effect (Figure 1E, bottom).

A detailed analysis of microtubule versus ER dynamics in CA versus DN Rac1 backgrounds is shown in Figure 2. Distal microtubule and ER ends are denoted by red and green triangles, respectively. Under both conditions, microtubule ends could enter the P-domain without ER attached; however, ER was never found in the periphery without a microtubule present. CA Rac1 dramatically increased microtubule rescue frequency, resulting in microtubules spending more time growing (Table 1) which in turn promoted ER advance. Figure 2A shows an example of ER extending outward along what seems to be a previously assembled microtubule. However, it is difficult to distinguish between ER elongating along a microtubule versus ER tracking with a microtubule tip (Grigoriev *et al*., 2008) because microtubules can coalign, making unambiguous plus-end localization challenging. An example of two closely aligned microtubules separating is evident in the 48- to 72-s interval in Figure 2A, bottom. Colabeling microtubules with a microtubule tip*-*tracking protein such as EB3 will be needed to resolve this issue.

In contrast, increased microtubule catastrophe frequencies were observed in DN Rac1 (Table 1). This resulted in shorter

Figure 3. 5-HT evokes Ca^{2+} rise in CA Rac1 backgrounds. (A) Baseline Ca^{2+} level in growth cones in different Rac1 backgrounds. \overline{N} = 6 (control), 6 (CA Rac1), or 5 (DN Rac1). Values are expressed as mean \pm SEM. (B) Probability of 5-HT (10 μ M)-evoked Ca²⁺ responses in growth cones in control, CA Rac1-, or DN Rac1-injected neurons. $N = 15$ (control), 18 (CA Rac1), or 20 (DN Rac1). (C) Left, DIC image of growth cone. C, central domain (C-domain); P, peripheral domain (P-domain). Green dashed line, interface between Cdomain and P-domain defined by organelle boundary. Middle and right, Ca^{2+} ratio images of the entire growth cone before (ctl) and after 7 min in 5-HT (10 μ M throughout). Red line, leading edge. Bar, $10 \mu m$. (D and F) Time course of 5-HT $(10 \mu M)$ effects. DIC (top) and $Ca²⁺$ ratio images (bottom) in CA Rac1-(D) or DN Rac1 (F)-injected neuron obtained from area of interest as indicated in C (left, yellow rectangle). Elapsed time (minutes) is shown at the top. Red dotted line, peripheral boundary. (E and G) Summary of $\Delta F/F_0$ plots recorded in the entire C-domain (red line) or P-domain (blue line) of growth cones quantifying Ca^{2+} response to 5-HT over time in CA Rac1 (E) or DN Rac1 (G) backgrounds. $N = 5$ (CA Rac1) or 4 (DN Rac1). Values are expressed as mean \pm SEM. Black bar shows 5-HT presence. N denotes the number of growth cones tested. $Ca²⁺$ ratio image is coded by pseudocolors in linear scale (see bars).

average microtubule growth periods and consequent ER depletion from the P-domain (Figure 2B).

Rac1 Regulates the Probability of 5-HT– evoked Ca2 Elevation

5-HT has both neurotransmitter and neuromodulator roles in neuronal development (Gaspar *et al*., 2003). Because *Aplysia* 5-HT receptors have been extensively characterized previously (Li *et al.*, 1995; Barbas *et al.*, 2005), and Ca²⁺ has been reported to participate in 5-HT function in neurons (Dropic *et al*., 2005; Li *et al*., 2005), we investigated whether Rac1 activity might affect 5-HT-evoked Ca^{2+} responses. We first tested the effect of Rac1 activity on basal $Ca²⁺$ levels and no significant differences were observed in different Rac1 backgrounds (Figure 3A).

Next, Ca^{2+} responses to 5-HT were assessed. Under control conditions only 13% of growth cones tested exhibited $Ca²⁺$ increases after 5-HT application (Figure 3B), and these responses tended to be transient. It is interesting that when cells were injected with CA Rac1, nearly 80% of growth cones tested responded to 5-HT with Ca^{2+} increases (Figure 3B and Supplemental Movie 3). A representative response in CA Rac1 background is shown in Figure 3, C and D. 5-HT triggered a sustained increase in Ca^{2+} in the entire growth cone, including C- and P-domains. After 5-HT washout, Ca^{2+} returned to baseline levels in 30–40 min (data not shown). Ca^{2+} response curves over time in C-domain and P-domain were essentially indistinguishable, reaching plateau levels after 7–8 min in 5-HT with a maximal increase of 45% (Figure 3E).

In contrast, in DN Rac1 backgrounds 5-HT typically did not trigger significant Ca^{2+} changes, and the response probability was only 5% (Figure 3, B, F, and G). Brief Ca^{2+} transients $(<1$ min in duration) were sometimes observed in the axon shaft in DN Rac1 backgrounds; however, these events were not counted as positive responses because they did not affect Ca^{2+} levels in \bar{C} - or P-domains. Together, these results suggest that Rac1 activity increases the probability

CA Rac1+U73343

e. B CA Rac1+U73122 0.6 $Ca(4FF_o)$
 \sim 0.2 $\mathbf 0$ $5-HT$ **Figure 4.** 5-HT–evoked Ca^{2+} rise depends -2 Ω $\overline{2}$ $\overline{4}$ on $PLC \rightarrow IP_3$ signaling cascade. (A and C) Time (min) Time-lapse montage of DIC (top) and Ca^{2+} ratio imaging (bottom) during $5-HT$ (10 μ M) CA Rac1+XeC treatment in CA Rac1-injected neurons pre-Е treated with U73122 (2 μ M) (A) or U73343 (2 5-HT Ω μ M) (C) for 30 min. (E and G) Time-lapse sequence of DIC (top) and Ca^{2+} ratio imaging (bottom) during 5-HT (10 μ M) exposure in CA Rac1-injected neurons pretreated with XeC (20 μ M) for 30 min (E) or Rya (25 μ M) for 40 min α (G). Dotted line, peripheral boundary. Elapsed time is shown at the top of DIC in minutes. $Ca²⁺$ ratio image is coded by pseudocolors in linear scale (see bars). B, D, F, and H, summary of $\Delta F/F_0$ plots recorded in the entire C-domain (red line) or P-domain (blue line) of growth cones quantifying Ca^{2+} response to CA Rac1+XeC 5-HT over time in CA Rac1-injected neurons 0.6 pretreated with U73122 (2 μ M) (B), U73343 (2 (EF_0)
(4 0.2 μ M) (D), or XeC (20 μ M) (F) for 20–30 min or with Rya (25 μ M) (H) for 40–50 min. In H, the control curves for C-domain and P-domain යී from Figure 3E are shown for comparison. $\mathsf 0$ Black bar shows 5-HT presence. $N = 4$ (CA $5-HI$ $Rac1 + U73122$), 3 (CA $Rac1 + U73343$), 4 (CA -2 0 $\overline{2}$ $\overline{4}$ $Rac1 + XeC$, or 4 (CA $Rac1 + Rya$). N denotes Time (min) the number of growth cones tested. Values are expressed as mean \pm SEM.

A

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and amplitude of Ca^{2+} increases in response to 5-HT in the C- and P-domains of the growth cone.

5-HT-evoked Ca2 Responses Involve Phospholipase C $(PLC) \rightarrow IP_3$ *Signaling*

We next investigated the Rac1-dependent Ca^{2+} response mechanism. In many cell types, 5-HT activates PLC, leading to IP₃ formation and Ca²⁺ release from IP₃ receptors (Yoder *et al*., 1996; Chiu *et al*., 1999; Noda *et al*., 2003; Dropic *et al*., 2005; Li *et al*., 2005). To investigate a role for PLC in 5-HT– induced responses, CA Rac1-injected neurons were pretreated for 30 min with U73122, a well known PLC inhibitor (Jin *et al.*, 1994; Zhou *et al.*, 1999), and Ca²⁺ imaging was performed before and after 5-HT exposure. PLC inhibition completely suppressed Ca^{2+} elevation in response to 5-HT (Figure 4, A and B). In control experiments using U73343, an inactive U73122 analog (Smith *et al*., 1990; Jin *et al*., 1994), 5-HT elicited typical \check{Ca}^{2+} responses (Figure 4, C and D), ruling out nonspecific U73122 effects. These results indicate that $\bar{5}$ -HT-evoked Ca²⁺ responses depend on PLC activation. To investigate the role of IP_3 receptor activation, we assessed Ca^{2+} responses in the presence of the membranepermeable IP3 receptor blocker XeC (Gafni *et al*., 1997). XeC pretreatment completely blocked 5-HT dependent Ca^{2+} release (Figure 4, E and F). Similar inhibition of 5-HT effects on $Ca²⁺$ release was observed when neurons were injected with heparin (Takei *et al.*, 1998)—a chemically distinct IP₃ receptor antagonist (data not shown).

We also tested whether Ca^{2+} release from ryanodine receptor (RyaR)-gated Ca^{2+} stores could be involved in the observed 5-HT responses. Ca^{2+} release from RyaR stores is activated by low concentrations (10–100 nM) and blocked by high concentrations ($>$ 10 μ M) of ryanodine (Meissner, 1986; McPherson *et al*., 1991); therefore, CA Rac1-injected neurons were treated with an inhibitory dose of ryanodine (25 μ M) for 40 min and then exposed to 5-HT in the continued presence of the inhibitor. Under these conditions, 5-HT continued to evoke Ca^{2+} release (Figure 4G).

Although the early phase of the Ca^{2+} response was unaffected by ryanodine, Ca^{2+} responses plateaued at lower levels after ${\sim}5$ min (Figure 4H). These results are consistent with Ca²⁺-induced Ca²⁺ release from RyaR stores contributing mainly to the late phase of the 5-HT-elicited Ca^{2+} rise. Because the RyaR and the IP_3 receptors regulate a common ER-associated Ca²⁺ pool (Walton *et al.*, 1991; Solovyova and Verkhratsky, 2003), it is not surprising that IP_3 receptor activation could lead to further RyaR-dependent Ca^{2+} release. Together, these data strongly suggest CA Rac1 is

Figure 5. Rac1 effects on microtubule–ER distribution modulate the pattern of 5-HT–evoked Ca^{2+} release. (A) Immunocytochemistry of actin filament and microtubule distribution in a Taxol (0.1 μ M; 20 min) pretreated growth cone in CA Rac1 backgrounds. Left, Actin. Middle, microtubule. Right, merge of actin filaments (red) and microtubules (green). Red, green, and blue lines: boundaries as in Figure 1D. Bar, 10 μ m. (B) Taxol $(0.1 \mu M)$ effects on ER distribution in growth cones in CA Rac1 backgrounds. Top, actin staining. Bottom, ER staining. Left, control. Middle, Taxol treatment for 20 min. Right, Taxol treatment for 40 min. Red line, C–P-domain boundary. Bar, 10 μ m. (C) 5-HT–evoked Ca²⁺ release in a Taxol-pretreated $(0.1 \mu M; 20 \text{ min})$ growth cone in CA Rac1 background. (1) DIC and Ca^{2+} kymographs sampled from a region similar to the white bar in Figure 1D before and after 5-HT (10 μ M) addition. Note Taxol was present throughout. Data acquired at 10-s intervals. Blue line, 75%P; red line, C–P interface; yellow line, leading edge. P, P-domain; C, Cdomain. Ca^{2+} kymograph is coded by pseudocolors in linear scale (see bar). (2) Time course of 5-HT–evoked Ca^{2+} response sampled from blue line and red line. (3) \tilde{Ca}^{2+} profiles sampled at 0 and 7 min showing Ca^{2+} distribution across growth cone. Red line indicates C–P-domain interface. (4) $\Delta F/F_0$ plots of 5-HT response recorded from entire C-domain (red) or P-domain (blue) of growth cones pretreated with Taxol (0.1 μ M; 20–30 min). Control curves for C- and Pdomains from Figure 3E are shown for comparison (open symbols). $N = 5$ growth cones for each condition. Values are expressed as mean \pm SEM. Black bar shows 5-HT presence. (5) Comparison of 5-HT (10 μ M) evoked peak Ca²⁺ for control versus Taxol $(0.1 \mu M; 20-30 \text{ min})$ pretreated growth cones in CA Rac1 backgrounds. Data from entire C- or P-domains normalized to control conditions. $N = 5$ growth cones for each condition. Values are expressed as mean \pm SEM. $*$ p < 0.01 and #p < 0.05.

potentiating 5-HT–evoked Ca^{2+} release from IP₃- and ryanodine receptor-gated Ca^{2+} stores.

Ca2 Release Topography Depends on Microtubule Assembly

The observation that CA Rac1 promoted microtubule and ER advance into the growth cone periphery (Figure 1, A, B, and E) suggested the possibility that Rac1 activity might spatially regulate Ca^{2+} release. To test this hypothesis, we used low levels of Taxol (100 nM) or vinblastine (25 nM) that damp microtubule assembly and have been shown to result in rapid clearance of microtubules from P-domain due to persistent microtubule coupling to retrograde actin flow (Suter *et al*., 2004). Taxol treatment would be expected to result in rapid ER clearance from the P-domain given ER coupling to microtubule movements (Figure 2).

We first assessed actin filament and microtubule distributions in CA Rac1-injected neurons with or without Taxol (0.1 μ M; 20 min) treatment. In control growth cones, individual microtubules aligned with filopodia (Schaefer *et al*., 2002) and penetrated into the P-domain (Supplemental Figure S3); in contrast, microtubules rarely entered the P-domain at all in Taxol-treated growth cones (Figure 5A).

Next, we examined ER distribution before and after Taxol treatment in CA Rac1 backgrounds. Taxol treatment resulted in a dramatic relocalization of ER within the growth cone (Figure 5B and Supplemental Movie 4). Low levels of vinblastine that have similar effects on microtubule dynamics had similar effects on ER distribution (Supplemental Figure S4C). It is interesting that ER network density decreased in C-domain over time under these conditions, which could account for the decrease in C-domain Ca^{2+} responses observed. These results suggest microtubule presence alone (Figure 5A) is not sufficient to sustain Ca^{2+} release in the C-domain and point to a novel role for microtubule assembly considered in the *Discussion*.

To assess Taxol/vinblastine effects on ER distribution, we quantified ER excursion events in CA Rac1 backgrounds (Table 2). ER excursions past the 25%P domain boundary decreased by \sim 60% after Taxol or vinblastine addition and no ER extended past 75%P boundary (Table 2). These observations suggested that alteration in microtubule assembly could affect spatial patterns of Ca^{2+} release by causing redistribution of $ER Ca²⁺$ stores, especially in the growth cone P-domain.

To test this hypothesis, we assessed 5-HT-evoked Ca^{2+} release in CA Rac1 backgrounds after inhibition of microtubule dynamics with Taxol. Under these conditions, Ca^{2+} elevation was restricted to the C-domain and proximal re-

Table 2. Taxol/vinblastine changes ER distribution in growth cones in CA Rac1 background

Boundary	before Taxol	Taxol	before Vinblastine	Vinblastine
25%P	$5.42 + 0.17$	$2.32 + 0.16*$	5.36 ± 0.18	$2.12 + 0.25*$
75%P	$2.04 + 0.16$	$0 + 0*$	$1.97 + 0.17$	$0 + 0*$

CA Rac1-injected neurons were treated by Taxol (0.1 μ M) or vinblastine (25 nM) for 20–30 min. Data represents ER crossings/10- μ m boundary length. Thirteen and 10 growth cones were assessed for Taxol and vinblastine treatments, respectively. Values are expressed as mean \pm SEM. Statistical analysis by two-tailed paired *t* test. $\frac{k}{P}$ < 0.001 vs. before taxol or vinblastine addition.

gion of the P-domain (Figure 5C1 and Supplemental Movie $\bar{5}$). Ca²⁺ levels no longer changed at all beyond the 75%P peripheral domain boundary (Figure 5C, 2 and 3) as observed under control conditions (compare Figure 3C). Figure 5C, 4 and 5, compares pooled control versus Taxol-pretreated 5-HT-evoked Ca²⁺ responses. C- and P-domain responses are of similar magnitude under control conditions; in contrast, Ca^{2+} responses were reduced in both the C- and P-domains after Taxol treatment (Figure 5C, 4 and 5) in accord with the observed reduction of ER density. Similar results were obtained using vinblastine (Supplemental Figure S4, A and B). Reduction of Ca^{2+} release in the C-domain subsequent to Taxol treatment was somewhat of a surprise because Taxol did not markedly alter microtubule density there (cf. Figure 5A). This finding bears further investigation as it suggests the possibility that a dynamic pool of microtubules in the C-domain may play a role in maintaining ER distribution there.

ROS Derived from Rac1 Activation Facilitates 5-HT– evoked Ca2 Release

The above-mentioned results are consistent with Rac1 effects on microtubule assembly promoting ER transport into the P-domain, thereby facilitating 5-HT–dependent Ca^{2+} release there. However, Rac effects on microtubule dynamics alone are not sufficient to explain our observations. Specifically, we noted that DN Rac1 abolished *all* 5-HT-dependent Ca^{2+} release in the growth cone (Figure 3F), including release in the C-domain where both microtubules and ER are clearly still present (Figure 1C).

Given that Rac is also an essential regulator of a multicomponent NADPH oxidase complex that controls ROS production (Lambeth, 2004) and recent evidence that cellular redox state can modulate Ca²⁺ release (Gordeeva *et al.*, 2003; Davidson and Duchen, 2006), we decided to investigate the relationship between Rac1 activity, growth cone ROS levels, and 5-HT–evoked Ca^{2+} release.

To assess growth cone ROS activity, neurons were loaded with DCF, a live-cell fluorescent ROS detector (Xie *et al*., 1999). In control experiments, DCF levels corrected for volume were monitored during H_2O_2 exposure, and the response was obtained over time, confirming the sensitivity of DCF (Supplemental Figure S5). Next, ROS levels in different Rac1 backgrounds were quantified. CA Rac1 significantly increased baseline ROS levels in both C- and P-domains, whereas DN Rac1 had the opposite effect (Figure 6A). To test whether elevated baseline ROS levels in CA Rac1 backgrounds could be modulating Ca^{2+} mobilization, neurons were pretreated with an antioxidant NAC (Sundaresan *et al*., 1996; Suzukawa *et al*., 2000) and then exposed to 5-HT. Antioxidant pretreatment totally blocked 5-HT–evoked $Ca²⁺$ responses typical of CA Rac1-injected cells (Figure 6, B and F, and Supplemental Figure S6A; cf. Figure 3, C–E). Note that NAC $(1 \text{ mM}; 30 \text{ min})$ had no significant effect on ER distribution in growth cones (data not shown). Moreover, pretreatment of CA Rac1-injected neurons with apocynin, a specific inhibitor of NADPH oxidase complex assembly (Fellner and Arendshorst, 2005; Kimura *et al*., 2005), similarly suppressed 5-HT–evoked Ca^{2+} responses (Figure 6, C and F, and Supplemental Figure S6B). These data strongly suggest ROS derived from Rac1 dependent NADPH oxidase activation are necessary for Ca^{2+} responses to 5-HT in the entire growth cone, i.e., in either Cor P-domains.

If 5-HT–evoked Ca^{2+} release depends on the presence of ROS, exposure to exogenous ROS would be expected to rescue Ca^{2+} responses to 5-HT in DN Rac1 backgrounds. To

Figure 6. ROS derived from Rac1 activation promotes 5-HT–evoked Ca^{2+} response. (A) ROS levels measured in C- and P-domains in control, DN Rac1, or CA Rac1 backgrounds. $N = 35$ (control), 36 (DN Rac1), and 34 (CA) Rac1). N denotes the number of growth cones tested. Values are expressed as whisker plots. Means are represented by dots, medians are represented by middle lines, 75th and 25th quartiles are represented by top and bottom of the boxes, respectively, and 90th and 10th percentiles are represented by whiskers. Statistical analysis by two tailed unpaired *t* test. #p 0.05 and $p < 0.01$ compared with control for C-domain and P-domain, respectively. (B and C) Summary of $\Delta F/F_0$ plots recorded in the entire C-domain (red sign) or P-domain (blue sign) of growth cones preincubated with NAC $(1 \text{ mM}, 20-30 \text{ min}; \overline{B})$ or apocynin $(1 \text{ mM}, 1)$ 40–50 min; C) to quantify Ca^{2+} response to 5-HT over time in CA Rac1 backgrounds. The control curves for C-domain and P-domain from Figure 3E are shown for comparison. $N = 5$ (CA Rac1 + NAC), 7 (CA Rac1 + apocynin), or 5 (CA Rac1). (D and E) Summary of $\Delta F/F_0$ plots recorded in the entire Cdomain (red sign) or P-domain (blue sign) of growth cones pretreated with H_2O_2 (10 μ M, $\overline{20}$ –30 min; D) or H_2O_2 plus XeC (10 and 20 μ M, respectively, 20–30 min; E) to quantify $Ca²⁺$ response to 5-HT over time in DN Rac1 backgrounds. The control curves for Cdomain and P-domain from Figure 3G are shown in D , and the H_2O_2 -conditioned curves for C-domain and P-domain from D are shown in E for comparison. $N = 4$ for each condition. N denotes the number of growth cones tested. Values are expressed as mean SEM. Black bar shows 5-HT presence. (F) Summary of 5-HT (10 μ M)-evoked Ca²⁺ peak values presented in B–E. Data were obtained from the entire C-domain and P-domain and normalized to those in CA Rac1 conditions. Values are expressed as mean \pm SEM. *p < 0.01.

test this hypothesis, cells injected with DN Rac1 were pretreated with a low concentration of H_2O_2 , and Ca^{2+} responses were assessed. In sharp contrast to controls (Supplemental Figure S6C), addition of 5-HT to neurons conditioned with H₂O₂ (10 μ M) resulted in dramatic Ca²⁺ increases in the C-domain (Figure 6, D and F, and Supplemental Fig. S6D). Ca^{2+} levels plateaued with a 30% increase after \sim 7 min in 5-HT (Figure 6D). $\rm H_2O_2$ conditioning did not rescue Ca^{2+} responses in the P-domain (Figure 6, D and F, and Supplemental Figure S6D)—this is expected as a consequence of peripheral ER depletion in DN Rac1 backgrounds (Figure 1). Note that in DN Rac1 backgrounds, $H_2O_2(10 \mu M;$ 20 min) did not affect ER distribution in growth cones significantly (data not shown).

The H_2O_2 rescue responses mentioned above depended on IP₃ receptor activation, because Ca²⁺ release was absent in the presence of XeC (Figure 6, E and F, and Supplemental Figure S6E). In control or DN Rac1 injected neurons, H_2O_2 (10 μ M) alone had little effect on growth cone morphology, and baseline Ca²⁺ levels did not change significantly $(<5\%)$ in the continued presence of H_2O_2 (>45 min exposure; data not shown). These results are consistent with a report that histamine-stimulated Ca^{2+} oscillations could be rescued in human aortic endothelial cells expressing DN Rac1, if cells

were exposed to low concentration of H₂O₂ (Hu *et al.*, 2002). Together, these data suggest that Rac1 activity modulates 5-HT–evoked Ca²⁺ release via ROS-dependent sensitization of IP_3 receptors.

DISCUSSION

Given its signaling diversity, it is not surprising that Ca^{2+} dynamics are tightly regulated both spatially and temporally. Rac regulation of Ca^{2+} metabolism has been reported in other systems, for example, Rac/Cdc42-inactivating *clostridial cytotoxins* inhibited intracellular Ca²⁺ mobilization induced by FcRI-stimulation in rat basophilic leukemia cells (Djouder *et al.,* 2000); CA Rac enhanced Ca²⁺ release from ER in mast cells after antigen stimulation, whereas DN Rac dampened Ca^{2+} mobilization (Hong-Geller and Cerione, 2000; Hong-Geller *et al*., 2001). The current study extends the role of Rac in regulation of Ca^{2+} metabolism to neuronal growth cones, where $5-HT$ –evoked Ca^{2+} release topography, and sensitivity seems to be modulated by Rac1 activity via parallel effects on microtubule dynamics and $IP₃$ receptor sensitivity.

Microtubule-dependent Changes in the ER Ca²⁺ Store Distribution

Because intracellular Ca²⁺ stores reside in ER (Ross *et al.*, 1989; Walton *et al*., 1991), which is a cargo for microtubulebased transport, Rac effects on microtubule dynamics could affect ER distribution. Indeed, Rac1 promotes microtubule growth in epithelial cells via Pac kinase-dependent inhibition of the microtubule catastrophe factor Op18/stathmin (Wittmann *et al*., 2004). Moreover, Rac activator DOCK7 inactivates Op18/stathmin in nascent axons (Watabe-Uchida *et al*., 2006). In agreement, we found that CA Rac1 facilitated microtubule extension into the P-domain by increasing microtubule rescue frequency and decreasing catastrophe frequency (Table 1). These effects were correlated with increased presence of microtubules and ER in the Pdomain (Figures 1, A, B and E, and 2A). When microtubule assembly was disrupted by Taxol or vinblastine treatment (Figure 5A), ER was also cleared from the growth cone periphery (Table 2, Figure 5B, and Supplemental Figure S4C), resulting in attenuated 5-HT–evoked Ca^{2+} responses (Figure 5C and Supplemental Figure S4, A and B). Collectively, these data suggest that altering the spatial distribution of intracellular \check{Ca}^{2+} pools in the P-domain may be an important mechanism for modulating Ca^{2+} responses there.

We also found that paradoxically Taxol/vinblastine treatments attenuated Ca^{2+} responses to 5-HT in the C-domain (Figure 5C, 4 and 5, and Supplemental Figure S4B) and reduced ER density (Figure 5B and Supplemental Figure S4C) despite continued high microtubule density there (Figure 5A). These effects are interesting because they suggest the presence of a heretofore uncharacterized pool of dynamic microtubules in the C-domain that play a role in regulating ER distribution. Such a pool would be difficult to detect because the high density of microtubules in the Cdomain precludes unambiguous analysis by speckle microscopy. These observations are consistent with a report that Taxol decreased thapsigargin-mediated $Ca²⁺$ release in adenocarcinoma cells (Padar *et al*., 2004). Note that neither Taxol (100 nM) nor vinblastine (25 nM) significantly altered

basal Ca²⁺ levels in growth cones (data not shown), suggesting these pretreatments do not significantly perturb existing Ca^{2+} stores. It will be of interest to investigate these apparent microtubule assembly dependent effects on ER distribution in the C-domain in future studies.

In DN Rac1 backgrounds, microtubules and ER were still present in C-domain (Figure 1, A and C), but now 5-HT– dependent Ca^{2+} release was essentially absent (Figure 3, F and G). These data suggest that the Rac1 effects on microtubule assembly and ER transport discussed above are not sufficient to explain the potentiating effects of Rac activity on 5-HT–evoked Ca^{2+} responses. The role of Rac in ROS production seems to be the missing piece of this puzzle.

ROS-dependent Changes in Ca²⁺ Release

Rac plays an essential role in NADPH oxidase-mediated ROS production and related signaling (Lambeth, 2004; Bedard and Krause, 2007). Evidence suggests a close connection between Rac1 activation and ROS generation in a variety of cell types, including neurons (Suzukawa *et al*., 2000), liver cells (Cool *et al*., 1998), astroglioma cells (Lee *et al*., 2002), and fibroblasts (Sundaresan *et al*., 1996). Indeed, we found that CA Rac1 significantly increased ROS levels in the growth cone C- and P-domains, whereas DN Rac1 had the opposite effect (Figure 6A). Given that both IP_3 receptors and ryanodine receptors seem to be sensitized by ROS (Davidson and Duchen, 2006), elevated ROS levels we see in CA Rac1 backgrounds may facilitate $5-HT Ca²⁺$ responses. In agreement, depleting ROS using a reactive oxygen scavenger (NAC) or blocking NADPH oxidase function (with apocynin) abolished 5-HT–evoked Ca^{2+} responses in CA Rac1 backgrounds (Figure 6, B, C, and F, and Supplemental Figure S6, A and B). In contrast, addition of extracellular H_2O_2 restored 5-HT–evoked Ca²⁺ release from IP₃ gated stores in DN Rac1 backgrounds (Figure 6, D, E, and F, and Supplemental Figure S6, D and E). These results support the idea that Rac1 activation is driving production of ROS that in turn potentiates the observed $5-HT$ –evoked Ca^{2+} responses. Parenthetically, treatment with the ROS scavenger

Figure 7. Model showing that Rac1 promotes 5-HT–induced Ca^{2+} mobilization by promoting microtubule-based ER advance and ROS-sensitized Ca^{2+} release. Rac-dependent Ca^{2+} regulation module is indicated by gray box. CICR, Ca^{2+} induced Ca^{2+} release.

NAC (1 mM; 30 min) or H_2O_2 (10 μ M; 20 min) did not affect ER distribution (data not shown), consistent with segregation of Rac effects on microtubules and NADPH oxidase.

It is well known that ROS such as H_2O_2 inactivate protein tyrosine phosphatases by Cys oxidization and potentiate tyrosine kinase signaling (Tonks, 2005). It is unlikely that this plays a significant role here because pretreatment with genistein (100 μ M), a general protein tyrosine kinase inhibitor, had no effect on H_2O_2 rescue of 5-HT–evoked Ca²⁺ responses in DN Rac1 backgrounds (data not shown). Because both IP_3 receptors and RyaRs are known to be sensitive to their thiol redox state, with oxidation of critical thiols favoring Ca^{2+} release (Abramson and Salama, 1988; Missiaen *et al*., 1991; Bootman *et al*., 1992; Xia *et al*., 2000; Joseph *et al*., 2006), we speculate that ROS may oxidize IP_3 and Rya receptors and thereby reduce the threshold for Ca^{2+} channel opening. Rac1 has also been reported to directly activate PLC (Illenberger *et al*., 1998; Snyder *et al*., 2003); however, we do not think constitutive PLC activation is involved because 1) Ca^{2+} responses were rescued by H_2O_2 in DN Rac1 backgrounds (Figure 6, D and F, and Supplemental Figure S6D) and 2) Ca^{2+} responses still depended on IP₃ receptor activation (Figure 6, E and F, and Supplemental Fig. S6E).

Rac is emerging as a functional node for coordination of multiple biochemical processes (Figure 7); here, we add to this story by providing evidence that Rac "modulates" the topology and sensitivity of Ca^{2+} release by 1) regulating microtubule assembly, which in turn affects ER distribution; and 2) increasing ROS production, which leads to sensitization of evoked \tilde{Ca}^{2+} release from IP₃-dependent stores. Although Rac activation would create an environment that facilitates stimuli-evoked Ca^{2+} responses, it is difficult to predict the relative importance of Rac effects stemming from changes in microtubule dynamics versus ROS production because each of these processes could be influenced by other signaling pathways.

 $Ca²⁺$ diffusion is highly restricted by binding proteins and pumps (Augustine *et al*., 2003); thus, downstream effectors in growth cones need to be situated relatively close to sites of Ca^{2+} release. We speculate that localized regions of high Rac activity will tend to accumulate microtubules due to net effects on assembly, this in turn will bring ER stores into proximity with potential Ca^{2+} effector proteins. Positive external modulators of Rac activity could thus "prime" the P-domain for ligand-dependent \check{Ca}^{2+} release (Figure 7). In primed regions, ligand dependent responses involving the $PLC \rightarrow IP_3$ cascade would also be facilitated by increased ROS levels. The net result would be a spatio-temporal sensitization of ligand dependent Ca^{2+} responses. The dependence of 5-HT–evoked Ca^{2+} responses on Rac activity presented here are in line with this type of model. It will be of interest to see how these Rac dependent processes affect the cytoskeletal proteins dynamics involved in axon growth.

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