Development/Plasticity/Repair

# Transient Receptor Potential Canonical 5 Channels Activate $Ca^{2+}$ /Calmodulin Kinase I $\gamma$ to Promote Axon Formation in Hippocampal Neurons

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Functionality of neurons is dependent on their compartmentalized polarization of dendrites and an axon. The rapid and selective outgrowth of one neurite, relative to the others, to form the axon is critical in initiating neuronal polarity. Axonogenesis is regulated in part by an optimal intracellular calcium concentration. Our investigation of  $Ca^{2+}$ -signaling pathways involved in axon formation using cultured hippocampal neurons demonstrates a role for  $Ca^{2+}$ /calmodulin kinase (CaMKK) and its downstream target  $Ca^{2+}$ /calmodulin kinase I (CaMKI). Expression of constitutively active CaMKI induced formation of multiple axons, whereas blocking CaMKK or CaMKI activity with pharmacological, dominant-negative, or short hairpin RNA (shRNA) methods significantly inhibited axon formation. CaMKK signals via the  $\gamma$ -isoform of CaMKI as shRNA to CaMKI $\gamma$ , but not the other CaMKI isoforms, inhibited axon formation. Furthermore, overexpression of wild-type CaMKI $\gamma$ , but not a mutant incapable of membrane association, accelerated the rate of axon formation. Pharmacological or small interfering RNA inhibition of transient receptor potential canonical 5 (TRPC5) channels, which are present in developing axonal growth cones, suppressed CaMKK-mediated activation of CaMKI $\gamma$  as well as axon formation. We demonstrate using biochemical fractionation and immunocytochemistry that CaMKI $\gamma$  and TRPC5 colocalize to lipid rafts. These results are consistent with a model in which highly localized calcium influx through the TRPC5 channels activates CaMKK and CaMKI $\gamma$ , which subsequently promote axon formation.

## Introduction

Establishment of morphological and functional polarity of neurons is seminal to formation of neuronal circuits in the CNS. Cultured rat embryonic hippocampal neurons are a well established model system to study the development of neuronal polarity in vitro (Craig and Banker, 1994). In this system, neuronal polarity is initiated by axon formation that occurs within the first 24-48 h. On plating, neurons initially form lamellapodia (stage 1) that develop into multiple short neurites that undergo repeated stochastic episodes of extension and retraction (stage 2) until one neurite extends rapidly to become the axon (stage 3). Amplification of a positive signaling feedback loop in one neurite presumably accelerates its outgrowth, which, when combined with negative-feedback signals from the other neurites, ensures formation of a single axon (Andersen and Bi, 2000). Here, we examined the role of calcium influx-activated signaling via CaM kinases (CaMKs) in the development of neuronal polarity.

Calcium regulates axon outgrowth and growth cone motility in complex ways (Gomez and Spitzer, 2000; Gomez and Zheng, 2006). Calcium transients occur spontaneously in growth cones and are linked to preaxonogenesis events (Bentley et al., 1991; Davenport and Kater, 1992). Transient receptor potential canonical (TRPC) channels are implicated in Ca<sup>2+</sup>-dependent growth cone dynamics (Li et al., 2005; Shim et al., 2005; Wang and Poo, 2005). TRPC members (TRPC1-7) form homo- or heterotetramers that function as nonselective cation channels and are abundantly expressed in embryonic brain (Strübing et al., 2001; Ambudkar and Ong, 2007). Notably, TRPC4- and 5-containing TRPC channels are the most prevalent TRPC channels in rodent brain (Fowler et al., 2007), and TRPC5-containing channels are present in growth cones of stage 2 hippocampal neurons (Greka et al., 2003). Given the increased calcium permeability of TRPC5 homomeric channels (Ambudkar et al., 2006; Beech, 2007), they are an attractive candidate for mediating Ca<sup>2+</sup> entry into developing neurons.

Transduction of Ca<sup>2+</sup> gradients in axonal growth cones is thought to be dictated by calmodulin-dependent protein kinases (CaMKs) or the CaM-dependent phosphatase, calcineurin (Wen et al., 2004). Previous studies concluded that CaMKII mediates neurite outgrowth since pharmacological inhibitors of CaMKII [KN-62 (1-[*N*, *O*-bis(5-isoquinolinesulfonyl)-*N*-methyl-L-tyrosyl]-4-phenylpiperazine) and KN-93 (2-[*N*-(2-hydroxyethyl)]-*N*-

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(4-methoxybenzenesulfonyl)amino-N-(4-chlorocinnamyl)-N-methylbenzylamine)] (Tokumitsu et al., 1990) generally suppress Ca<sup>2+</sup>-dependent neurite outgrowth (Zheng et al., 1994; Kuhn et al., 1998). However, we now know these inhibitors are not specific for CaMKII but also inhibit CaMKIV and CaMKI (Mochizuki et al., 1993; Enslen et al., 1994) as well as some Ca<sup>2+</sup>-activated potassium channels (Ledoux et al., 1999) and voltage-gated Ca<sup>2+</sup> channels (Anderson et al., 1998). We previously demonstrated that the Ca<sup>2+</sup>/calmodulin kinase kinase (CaMKK)/CaMKI cascade (Soderling, 2000), which is present in neurons throughout development (Kamata et al., 2007a,b), regulates various stages of neuronal development including stage 3 axonal growth cone motility and outgrowth (Wayman et al., 2004), stage 4 dendritic arborization (Wayman et al., 2006), and stage 5 spine/synapse formation (Saneyoshi et al., 2008). Using multiple approaches, we demonstrate here that CaMKK/ CaMKIy, activated by Ca<sup>2+</sup>-permeable TRPC5 channels, promote axon formation in cultured hippocampal neurons.

## **Materials and Methods**

*Pharmacological inhibitors and antibodies.* 1,8-Naphthoylene benzimidazole-3-carboxylic acid (STO-609) was purchased from Tocris. 1-[ $\beta$ -[3-(4-Methoxyphenyl)propoxy]-4-methoxyphenethyl]-1*H*-imidazole, HCl (SKF96365), anthra[1,9-*cd*]pyrazol-6(2*H*)-one (SP600125), and nifedipine were from Calbiochem. Monoclonal antibody for Tau-1 was from Millipore Bioscience Research Reagents. Monoclonal anti-Flotillin-1 antibody was from BD Biosciences Transduction Laboratories. Monoclonal anti-Transferrin receptor (TfR) antibody was from Zymed. Rat polyclonal anti-TRPC5 and anti-TRPC6 antibodies were from Alomone Labs. E7 monoclonal ascites to detect  $\beta$ -tubulin by Western blotting and immunocytochemistry was purchased from Developmental Studies Hybridoma Bank. Phospho-CaMKI and CaMKI $\gamma$  antibodies were gifts from our collaborator, Dr. Naohito Nozaki (Kanagawa Dental College, Kanagawa, Japan).

Cell culture and transfection. Low-density hippocampal neuron cultures were prepared from embryonic day 18 (E18) rat hippocampi and cultured using the glial sandwich method as previously described in detail (Kaech and Banker, 2006; Oliva et al., 2006). For cDNA and plasmid based short hairpin RNA (shRNA) expression,  $0.5 \times 10^6$  E18 hippocampal neurons were electroporated with 1–5  $\mu$ g of cDNA or with 2.5  $\mu$ g of Stealth small interfering RNA (siRNA) using AMAXA nucleofection system before plating [details described by Kaech and Banker (2006)]. Approximately 30–50% of the electroporated neurons ( $\sim 0.15-0.25 \times 10^6$ ) were plated on four poly-L-lysine-coated 18 mm glass coverslips that were placed in a 60 mm culture dish with plating media. For pharmacological inhibition of signaling pathways, the inhibitors were added to the plates containing glia (and glial conditioned media) just before introducing the glass coverslips containing the hippocampal neurons after the 2-4 h attachment period. When using plasmid based shRNA knockdown approach, to extend the time window for effective shRNA knockdown of CaMKK/CaMKI pathway, we incubated the neurons with 20 µM SP600125 for the first 48 h in culture. We then released the suppressed stage 2 neurons from the SP600125 block by gently moving the coverslips to new glial plates (with glial conditioned media) without inhibitors. This constituted a washout of SP600125. This technique has been previously described by Oliva et al. (2006). Once transferred to new glial plates, the neurons were kept in culture for an additional 48 h to allow axon formation.

shRNA, siRNA, and plasmids. The construction, validation, and specificity of the plasmid-based shRNA constructs for knockdown of CaMKK/CaMKI cascade have been described previously (Wayman et al., 2006; Saneyoshi et al., 2008). For selective knockdown of TRPC5 and TRPC6 subunits of TRPC channels we used Block-iT Stealth RNA interference (RNAi) (Invitrogen). siTRPC5 was custom synthesized targeting nucleotide starting position 1900 (CATGCTGATATTGAGTGGAAGTTTG; accession number NM\_080898; rat TRPC5), and siTRPC6 was generated by custom synthesis of RNAi against nucleotide starting position

1506 (AGGAATGATATGGGCTGAATGTAAA; accession number NM\_053559; rat TRPC6). siTRPC5 and siTRPC6 were validated for knockdown and specificity by cotransfection of Myc-mTRPC5 and HA-mTRPC6 with 20 and 40 nm of corresponding siRNA in COS7 cells.

Constitutively active CaMKI (caCaMKI), dominant-negative CaMKI (dnCaMKI), dnCaMKK, CaMKKins, and enhanced green fluorescent protein (EGFP)-CaMKIIN have been described previously (Wayman et al., 2004, 2006; Saneyoshi et al., 2008). Truncated Kinesin-1 construct used to identify axon(s) has also been described previously (Jacobson et al., 2006). Human CaMKIγ (hCaMKIγ) was cloned from human hippocampal cDNA library from BD Biosciences (primers based on accession number NM\_020439) and subsequently subcloned into pCAGGS vector with N-terminal Flag epitope tag using ClaI and XhoI sites. EGFPhCaMKIy was constructed by first subcloning hCaMKIy into pEGFP-C1 followed by subcloning of the EGFP-hCaMKIγ fusion into pCAGGS. To generate truncation mutant CaMKI $\gamma_{\rm DPP}$ , Met407 was mutated to a TAG stop codon using QuickChange Site-Directed Mutagenesis kit (Stratagene). All constructs were verified by sequencing and their expression was tested in transiently transfected COS7 cells and cultured hippocampal neurons before use in experiments.

Immunocytochemistry. For Tau-1 immunostaining, neurons were fixed at indicated times, in prewarmed 4% paraformaldehyde (PFA) in PHEMS buffer (60 mm PIPES, 25 mm HEPES, 2 mm MgC1<sub>2</sub>, 1 mm EGTA, pH 6.9) for 20-30 min at room temperature (RT). After fixation, neurons were permeabilized with 0.1% Triton X-100 in PBS for 5–10 min and blocked for 1 h at RT with PBS containing 0.5% Fish Gelatin (Sigma-Aldrich). Axons were labeled by overnight incubation (at 4°C) with anti-Tau-1 monoclonal antibody diluted at 1:3000 in blocking buffer followed by brief wash (three times; 5 min each) and incubation with donkey anti-mouse Cy3 secondary antibody (1:3000) for 1 h at RT. After secondary antibody incubation, neurons were washed with PBS (four times; 10 min) and coverslips were mounted using elvanol mounting medium. Endogenous TRPC5 was immunostained using the primary rabbit anti-TRPC5 antibody (Alomone Labs; 1:1000) and donkey anti-rabbit Cy3 (1:5000) secondary antibody. β-Tubulin was immunostained using E7 ascites (1:3000) and Alexa Fluor 350-conjugated anti-mouse IgG (1:1000). Flotillin-1 immunocytochemistry was performed with monoclonal antibody diluted 1:1000 in blocking buffer, followed by incubation with Alexa Fluor 488-conjugated goat anti-mouse (1:1000). Endogenous monoganglioside 1 (GM1) in 1 d in vitro (DIV 1) E18 hippocampal neurons plated at low density was stained live with Alexa Fluor 546conjugated cholera toxin B subunit (CTxB) (Invitrogen) diluted to a final concentration of 2 µg/ml in prewarmed artificial CSF (ACSF) for 6-8 min at 37°C. Immediately after staining, the coverslips were briefly rinsed (2-4 s) in prewarmed ACSF and fixed with 4% PFA diluted in PHEMS buffer for 20 min at room temperature. For colocalization with EGFP-CaMKIy, the coverslips were mounted and imaged immediately. Flotillin-1 costaining with CTxB::GM1 was done as described above.

To measure the level of endogenous TRPC5 knockdown by siTRPC5 with immunocytochemistry, neurons were electroporated with control siRNA or siTRPC5 along with soluble EGFP as a transfection marker and were fixed and immunostained with anti-TRPC5 after 48 h as described in Materials and Methods. To quantify the level of TRPC5 knockdown by immunocytochemistry, multiple random fields were captured from both control and siTRPC5 conditions. We measured the pixel intensity (P.I.) (in arbitrary units) of immunofluorescence from siTRPC5-transfected (i.e., EGFP-positive) neuron relative to neighboring nontransfected neuron(s) within the same field that served as internal positive controls.

Microscopy, imaging, and quantification. Images were acquired by using a cooled CCD camera (Hamamatsu Photonics) attached to a Zeiss Axioplan 2 (Carl Zeiss) inverted microscope with a 10 or  $20 \times$  lens for quantification of neuronal polarity and with a  $63 \times$  oil-immersion lens for imaging immunostaining of endogenous TRPC5, Flotillin-1, β-tubulin, GM1, and for colocalization of EGFP-CaMKIγ and TRPC5. Images were processed using ImageJ and Adobe Photoshop 10. For analyzing numbers of neurons that formed axons under various conditions, 15–30 random fields from two to four individual coverslips (usually representing  $\sim$ 100–200 neurons) were captured as described above using the filter sets for EGFP (soluble fill as marker for transfection) and

Cy3 (Tau-1). Total neurons, neurons expressing Tau-1-positive process (termed axon), and where indicated, number of Tau-1-positive axons per neurons were manually counted using ImageJ (W. S. Rasband, ImageJ, National Institutes of Health, Bethesda, MD; http://rsb.info.nih.gov/ij/; 1997–2008). The percentage of neurons under basal conditions that formed an axon ranged from 60 to 85% because of natural variability in cultures. To facilitate statistical analysis between experiments, the control average from each experiment was set to 100%, and the experimental data (average  $\pm$  SEM) were normalized relative to the control. Statistical analyses and graphing were performed using unpaired Student's t test using Microsoft Excel or GraphPad Prism 3.0. To measure axonal length, the NeuronJ plugin was used in conjunction with ImageJ. Raw axonal length data from NeuronJ were adjusted to micrometers based on calibration ruler measurements. Data in the graphs are represented as averages plus SEMs from two to five independent experiments.

Lipid raft fractionation and Western blotting. Lipid raft fractionation was done from postnatal day 0 (P0) rat forebrains with a discontinuous sucrose gradient using protocols adapted from previously described methods (Maekawa et al., 2001; Suzuki et al., 2004). Briefly, we made homogenates from 1.3 g of P0 forebrain material per 10 ml of buffer A (50 mm HEPES, pH 7.4, 1 mm MgCl<sub>2</sub>, 1 mm NaF plus protease and phosphatase inhibitors) with a Dounce homogenizer. This crude homogenate was centrifuged at 200  $\times$  g for 5 min in a SS-34 rotor to pellet larger particulate material. The supernatant from this was centrifuged at 50,000 rpm (179,000 g force) in a 70Ti rotor for 30 min. The resulting supernatant was designated as the crude cytosol and was discarded. The pellet fraction represented the crude membranes. To concentrate the crude membrane fraction before loading on the gradient, the membranes were resuspended in 5 ml of buffer B (buffer A plus 1% Triton X-100) by homogenization with a Dounce homogenizer. After homogenization, this membrane suspension was incubated on ice for 30 min. This membrane fraction was then loaded on a discontinuous sucrose gradient as follows: 0.5 ml of the crude membrane suspension was mixed with 0.5 ml of 90% sucrose (in buffer A). This 1 ml fraction composed of a 50:50 mix of membranes in 45% sucrose was loaded on the bottom of SW41Ti rotor tubes; this was gently overlaid with 8 ml of 35% sucrose in buffer A and finally 3.5 ml of 5% sucrose in buffer A. The gradient was centrifuged at  $140,000 \times g$  for 18 h in a SW41Ti rotor at 4°C (low brake setting). One milliliter fractions were collected by puncturing the tubes from the bottom and collecting fraction from bottom to top (bottom-most fraction is designated as fraction 12). Protein concentration was determined using Bradford reagent (Bio-Rad), and fractions were loaded on SDS-PAGE and transferred with polyvinylidene difluoride. Western blotting was with indicated antibodies at the following dilutions: anti-TRPC5, 1:1000; anti-TRPC6, 1:1000; anti-CaMKI, 1:100; anti-Flotillin-1, 1:2000; and anti-TfR, 1:2000. Western blot detection (IR700- and IR800-conjugated secondary antibodies) and quantification were done using the Odyssey Infrared System (LI-COR Biosciences).

#### Results

## Constitutively active CaMKI promotes multiple axons

We have previously demonstrated a role for CaMKK and CaMKI in regulation of axonal extension and growth cone motility in neonatal hippocampal neurons (Wayman et al., 2004). These experiments were conducted using stage 3 (DIV 2-4) cultured neurons that are already polarized with a single extended axon and several short neurites that subsequently become dendrites. Under these conditions, expression of dnCaMKK or dnCaMKI, but not nuclear localized dnCaMKIV or a protein inhibitor of CaMKII, caused growth cone collapse and significantly reduced total axonal length and branching. Conversely, transfection of a caCaMKI stimulated axonal extension in stage 3 neurons. In the current study, we investigated the role of the CaMKK/CaMKI signaling pathway in the early steps involved in the establishment of neuronal polarity. We used a low-density embryonic (E18) hippocampal culture system that is an optimal and well defined model system for studying neuronal polarity (Craig and Banker,

1994; Kaech and Banker, 2006). In these cultures, most neurons do not display an axon at 24 h (stage 2; all neurites are similar length), whereas at 48 h a majority (>70%) have one neurite that is more than twice the length of the others, and is specified as the axon (stage 3 neuron) (Deitch and Banker, 1993). To modulate axon formation with pharmacological inhibitors and transfection protocols, we electroporated neurons before plating and initiated all pharmacological treatments at 2 h after plating.

To investigate the role of the CaMKK/CaMKI cascade in axon formation, we transfected hippocampal neurons with constitutively active CaMKI $\alpha$  that has been previously tested (caCaMKI) (see below). Overexpression of caCaMKI should act as a generic constitutively active CaMKI (Wayman et al., 2004, 2006; Saneyoshi et al., 2008). Approximately 70% of neurons transfected with caCaMKI were polarized by 48 h, and although transfection with caCaMKI did not alter the length of the primary projection (control,  $263.9 \pm 13.4 \mu m$ ; caCaMKI,  $251.5 \pm 12.6$ ; n = 50 neurons), neurons expressing caCaMKI had significantly longer minor processes (Fig. 1A). Most of these secondary projections were in fact axons, as defined by staining for the well established axonal marker, Tau-1 (Mandell and Banker, 1996) (Fig. 1B). In Figure 1 *B*, the left vertical panels show representative images of soluble EGFP cotransfected with empty vector (control, top) or ca-CaMKI (bottom); the middle vertical panels show Tau-1 staining in one distinct elongated process in control neurons but in multiple elongated processes in neurons expressing caCaMKI. Analyses from five independent experiments revealed that control neurons have an average of 1.12  $\pm$  0.16 axons per neuron (n =400 neurons) versus  $2.7 \pm 0.25$  (n = 276 neurons) axons in neurons expressing caCaMKI (Fig. 1D).

To further validate that these Tau-1-positive projections induced by caCaMKI were axons, we used another confirmed marker of axons, Kinesin-1. It was recently shown that truncated Kinesin-1 selectively accumulates in the emerging axon and is excluded from all other neurites (Jacobson et al., 2006). We cotransfected neurons with an EGFP-tagged truncated Kinesin-1 (Kif5c<sup>560</sup>) along with empty vector (control) or caCaMKI. Neurons were stained with  $\beta$ -tubulin antibody to identify all projections (pseudocolored green). As shown in Figure 1C, EGFP-Kif5C<sup>560</sup> (pseudocolored red) was selectively accumulated in the only in the tips of axonal branches of control cells and was absent from other neurites. However, in caCaMKI-expressing neurons, Kif5C<sup>560</sup> was present in tips of multiple projections, demonstrating that expression of caCaMKI shifted multiple minor neurites toward biochemically defined axons. The caCaMKI induced a  $\sim$ 2.4-fold increase in the number of Kif5C <sup>560</sup>-containing axons per neuron compared with control neurons (caCaMKI, 2.5 ± 0.09 axons/neuron, vs control, 1.12  $\pm$  0.1 axons/neuron; n = 30neurons for both conditions) (Fig. 1*E*).

We also performed live imaging of hippocampal neurons expressing control vector or caCaMKI to visualize axon formation (supplemental movies, available at www.jneurosci.org as supplemental material). As seen in the movies, neurons expressing caCaMKI show enhanced outgrowth of multiple axons compared with a single axon in control transfection. The above data suggest that the CaMKI pathway may play a role in regulating not just axonal outgrowth of stage 3 neurons as reported previously (Wayman et al., 2004) but may regulate axon formation itself. Therefore, we designed additional experiments to specifically test whether modulation of CaMKI pathway alters axon formation and axonal length in low-density embryonic hippocampal cultures.

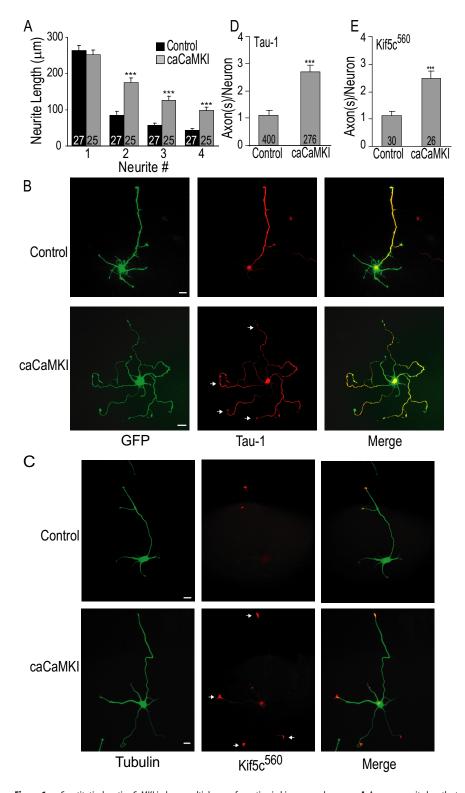


Figure 1. Constitutively active CaMKl induces multiple axon formation in hippocampal neurons. *A,* Average neurite length at 48 h from neurons transfected either with control plasmid (empty vector) or caCaMKl. *B,* Low-density E18 hippocampal cultures were electroporated before plating with either pCAGGS (control; top panel) or pCAGGS-caCaMKl (bottom panel) and soluble EGFP (marker for transfection) for 48 h followed by fixation and staining as described in Materials and Methods. Representative immunofluorescent images demonstrate formation of single Tau-1-positive neurite defined as the axon in control cells. In contrast, expression of caCaMKl strongly increases the number of Tau-1-positive axons. *C,* GFP-tagged Kif5C <sup>560</sup> (pseudocolored red) was cotransfected with control or caCaMKl before plating, and neurons were fixed and stained with anti-β-tubulin (pseudocolored green) at 48 h. Representative images are shown. *D, E,* Quantification (means ± SEM) of number of Tau-1-positive axons per neuron from three independent experiments (*D*) and GFP-tagged Kif5C <sup>560</sup>-containing neurite tips from control and caCaMKl (*E*). Scale bars: 20 μm (all panels). Here and in all subsequent figures, all graphs show means ± SEM (*n* values are shown as numerical insets within bars of graphs; where indicated,  $p \le 0.05$ , \*\* $p \le 0.005$ , \*\* $p \le 0.005$  by Student's *t* test).

## The CaMKK inhibitor, STO-609, blocks axon formation and elongation

In the preceding experiments, we used a mutant of CaMKI that is constitutively active, but the physiological upstream activator of CaMKI is CaMKK. To assess the cellular role of CaMKK, we incubated E18 hippocampal neurons with the cellpermeable CaMKK inhibitor STO-609 (Tokumitsu et al., 2002, 2003). In vitro, STO-609 selectively blocks CaMKK with little direct effect on CaMKI, CaMKII, CaMKIV, PKA (protein kinase A), PKC (protein kinase C), MLCK (myosin light chain kinase), or p42/44 MAPK (mitogenactivated protein kinase) (Tokumitsu et al., 2002), and we have shown that STO-609 can selectively block activation of CaMKI in cultured hippocampal neurons without suppressing CaMKII activation (Wayman et al., 2004). Treatment with 5.2 µm STO-609, beginning at 2 h after plating, resulted in only 45.6 ± 3% of neurons developing an axon at 48 h compared with 75% of control neurons (Fig. 2A, B). We performed a time course experiment to determine whether incubation of neurons with STO-609 only delays axon formation or results in a chronic block of axonogenesis. Hippocampal neurons were transfected with soluble EGFP, incubated with 5.2  $\mu$ M STO-609 for up to 96 h in culture, fixed, and stained with Tau-1 antibody to mark axons. As shown in Figure 2B, incubation with STO-609 for up to 96 h resulted in a significant number of neurons remaining chronically suppressed at stage 2 (average decrease of  $51.1 \pm 2.8\%$  at 24-96 h). Inhibition of CaMKK by STO-609 selectively suppresses axonal development, since the length of the minor neurites, which were Tau-1 negative, was not significantly altered in the presence of this inhibitor (supplemental Fig. 1*A*, available at www. jneurosci.org as supplemental material). This is consistent with our previous observation that inhibition of CaMKK does not suppress basal (i.e., activityindependent) stage 4 development of dendrites from neurites (Wayman et al., 2006). The suppression of axon formation in hippocampal neurons with STO-609 was reversible (data not shown). These data suggest a regulatory role for CaMKK in axon formation during stage 2 to stage 3 progression.

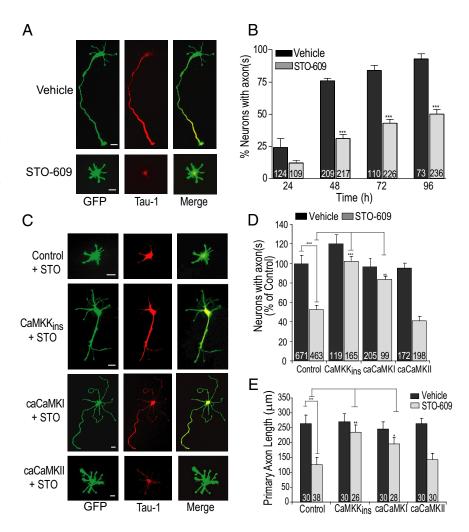
Although STO-609 has strong selectivity for inhibition of CaMKK relative to other protein kinases (Tokumitsu et al., 2002), we wanted to confirm that its inhibitory effect was in fact attributable to suppression of CaMKK. To this end, we

used a STO-insensitive CaMKK mutant (designated CaMKK<sub>ins</sub>) that has a single amino acid mutation of CaMKKα (L233F), rendering it 100-fold less sensitive to STO-609 (Tokumitsu et al., 2003). We tested whether expression of this STO-insensitive CaMKK could rescue axon formation in the presence of STO-609. In the presence of STO-609, 52.2% of neurons remained blocked at stage 2 relative to vehicle-treated control neurons (set to 100%) at 48 h. Expression of CaMKK<sub>ins</sub> in the presence of STO-609 showed complete rescue of axon formation and axonal length (Fig. 2C-E). The effects of STO-609 were attributable to inhibition of CaMKK and downstream signaling in neurons, and not the supporting glial cells, because in glial conditioned medium alone STO-609 suppressed axon formation to the same degree as with the neuron-glial sandwich cultures (supplemental Fig. 1B, available at www.jneurosci. org as supplemental material).

By suppressing activity of CaMKK, STO-609 inhibits activation of its downstream targets including CaMKI. Thus, expression of caCaMKI, which bypasses the requirement for activation of CaMKI by CaMKK, should also restore axon formation in STO-609-treated cells. Transfection with caCaMKI significantly reversed the STO-609 phenotype by increasing Tau-1-positive neurons to 83.8% in the presence of STO-609 (compare with  $52.8 \pm 4\%$  with empty vector) (Fig. 2C, D). Even in the presence of STO-609, a significant number of caCaMKI-expressing neurons still exhibited more than one axon (as in Fig. 1). Expression of a constitutively active kinase poses the risk of promiscuous substrate activation (Wayman et al., 2008), since various CaMKs have overlapping substrate specificity determinants (e.g., an Arg in the P-3 position). CaMKII is the other major neuronal cal-

cium/calmodulin kinase with functional implications in aspects of neuronal development (Wen et al., 2004), so it was important to test whether expression of caCaMKII [H282R mutant (Brickey et al., 1994)] might also reverse the STO-609 phenotype. However, expression of caCaMKII was unable to rescue the suppression of axon formation or length of axon by STO-609 (Fig. 2C–E). Therefore, it is unlikely that the effects of CaMKK $_{\rm ins}$  and caCaMKI are attributable to nonspecific substrate phosphorylation.

Since STO-609 produces only a partial block ( $\sim$ 50%) (Fig. 2D) of axon formation, we examined axonal length in those neurons that form a morphologically distinct axon in the presence of STO-609. Axon elongation is a critical component of establishing neuronal polarity in the sense that continuous growth of one neurite permits the establishment of axonal characteristics. The neurons that still formed an axon in the presence of STO-609 had significantly shorter axons (Fig. 2E) (control average length of the primary axon, 263.9  $\pm$  13.4  $\mu$ m; STO-609-treated cells,



**Figure 2.** Inhibition of CaMKK by STO-609 chronically suppresses axon formation. **A**, Low-density hippocampal cultures electroporated before plating with soluble EGFP were incubated with vehicle (control) or  $5~\mu$ m STO-609 from 2 h until fixation at 48 h. Representative images with axonal marker Tau-1 shows normal formation of a single axon in control neurons with suppressed axon formation in neurons incubated in STO-609. **B**, STO-609 time course with neurons fixed at 24, 48, 72, and 96 h reveals chronic partial block of axon formation. **C**–**E**, Expression of STO-609-insensitive CaMKK (CaMKK<sub>ins</sub>) or caCaMKI, but not caCaMKII, rescue STO-609 inhibition of axon formation and outgrowth. All inhibitor and inhibitor rescue data are shown normalized to control conditions set to 100% [60 –90% of neurons polarized under basal (control) conditions] in this and subsequent figures. Representative images from STO-609-insensitive CaMKK, caCaMKI, and caCaMKII are shown in **C**. Quantification of three independent experiments are shown in **D**. Average axonal length as measured by length of primary Tau-1-positive axon is shown in graph in **E**. Scale bars: 20  $\mu$ m (all panels).

 $128.6\pm7.5~\mu m$ ). Transfection with CaMKK<sub>ins</sub> or caCaMKI, but not caCaMKII, rescued inhibition of axonal outgrowth by STO-609 (Fig. 2E). These data together indicate a specific role of the CaMKK/CaMKI signaling cascade in axon formation.

In studies of axon specification, neuronal polarization is typically defined by the appearance of a projection that is longer than the other neurites and displays hallmark axonal markers. One neuronal polarity model suggests that axon fate is conferred on one neurite because of accumulation of "axon-specifying" components in this neurite, followed by its rapid outgrowth to form the axon. Another polarity model conceptualizes that accelerated growth of one neurite is the critical initial step in axonogenesis, leading to preferential enrichment of key axonal components and consequential acquisition of axonal fate by this neurite (Jiang and Rao, 2005; Winckler, 2007). Because of experimental dependence on appearance of a morphologically discernable axon in our cultured neuron model system, it is difficult to mechanistically dif-

ferentiate between "specification" and "outgrowth" as both are requisite steps. The fact that transfection with caCaMKI promoted formation of multiple axons may indicate it has a role in axon "specification." However, our other experiments cannot differentiate between specification and outgrowth, so we have used a more general term "axon formation" that is commonly used in this field.

## Dominant-negative CaMKK and CaMKI block axon formation

To establish the physiological role of a signaling pathway, it is essential to use multiple, independent protocols to modulate the functionality of that pathway (for review, see Wayman et al., 2008). In addition, to further investigate the potential role of endogenous CaMKII in axon development, we expressed CaMKIIN, a highly selective endogenous protein inhibitor of CaMKII (Chang et al., 1998). Transfection of CaMKIIN completely blocks activation of CaMKII, even on depolarization with KCl, without effect on axonal outgrowth of stage 3 neurons (Wayman et al., 2004). Expression of dnCaMKK mimicked the effects of STO-609 such that only 56.3  $\pm$  3.5% of neurons had Tau-1-positive axons (Fig. 3A,B). Of the dnCaMKK-expressing neurons that did develop axons, their axonal length was significantly reduced (control, 210.3  $\pm$  19.1  $\mu$ m, vs dnCaMKK, 124  $\pm$  12  $\mu$ m) (Fig. 3C). The dnCaMKI had a similar effect with 69.7  $\pm$  5.8% of neurons developing axons compared with control. We previously showed that nuclear-restricted dnCaMKIV (dnCaMKIV<sub>NIS</sub>) had no effect on stage 3 axonal outgrowth (Wayman et al., 2004). Here, in our experiments with the stage 1 to stage 2 neurons in the low-density culture system, expression of dnCaMKIV<sub>NLS</sub> resulted in significant toxicity, thereby making the results uninterpretable. Overexpression of CaMKIIN did not significantly suppress axon formation or affect axonal length. These data corroborate the results from the preceding experiments indicating that CaMKK and CaMKI, but not CaMKII, play a modulatory role in axon formation of E18 hippocampal neurons in our system.

# shRNA knockdown of CaMKK and CaMKI decreases axonogenesis

Since ectopic expression of dnCaMKK and dnCaMKI may result in unknown regulation of other signaling pathways, we verified these results by acutely knocking down the expression of endogenous CaMKK and CaMKI using plasmid-based shRNA approach. Both CaMKK isoforms ( $\alpha$  and  $\beta$ ) and CaMKI subfamily members that include four isoforms ( $\alpha$ ,  $\beta$ ,  $\gamma$ , and  $\delta$ ) are present in the hippocampus at varying levels during early development (Nishimura et al., 2003; Sakagami et al., 2005; Kamata et al., 2007a,b). To determine which CaMKI isoform(s) regulates axon formation in neurons, we used CaMKI isoform-specific shRNAs that have been previously validated for knockdown efficacy and specificity in HEK293 cells (Wayman et al., 2006) as well as shRNAs for  $\alpha$  and  $\beta$  CaMKK (Saneyoshi et al., 2008). As shown in Figure 2B, axon formation in these cultures occurs precipitously between 24 h (24  $\pm$  7% stage 3 neurons) and 48 h (76  $\pm$  2% stage 3 neurons), but with the plasmid-based shRNA approach it is difficult to achieve substantial reductions of proteins within 24 h. To increase the duration of shRNA expression to allow sufficient knockdown of CaMKK and CaMKI before axon formation initiates, we arrested neuronal development at stage 2 with 20  $\mu$ M SP600124 from 2 to 48 h in culture, followed by release from block to initiate axon formation for another 48 h. SP600125 is a

Jun kinase inhibitor that reversibly blocks axon formation without affecting neuronal viability (Oliva et al., 2006). To release hippocampal neurons from SP600125 block, the coverslips were transferred to new plates containing glia and their glial-conditioned media. Incubating neurons with SP600125 for 48 h suppressed axon formation in 98.5  $\pm$  1% of neurons—this block was almost completely reversible, with 90  $\pm$  2% of neurons developing axons after the 48 h washout (supplemental Fig. 1*C*, available at www.jneurosci.org as supplemental material).

Transfection of E18 hippocampal neurons with a mixture of shRNAs for both CaMKK isoforms ( $\alpha$  and  $\beta$ ) decreased the number of neurons exhibiting Tau-1-positive axons to only 63  $\pm$  4.3% of control values (control, 100%). Of the four distinct isoforms of CaMKI ( $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\delta$ ), shCaMKI $\gamma$  significantly blocked axon formation (only 58.4  $\pm$  6.9% neurons formed Tau-1-positive axons) to the same degree as shCaMKK (Fig. 4*A*,*B*). shCaMKI $\alpha$  also had a modest but statistically significant effect on axon formation (83.1  $\pm$  7.3% formed an axon). As shown in Figure 4*C*, the neurons that did develop an axon in the presence of shCaMKK and shCaMKI $\gamma$  had significantly shorter axonal lengths (52.2% decrease with shCaMKK and 40.1% decrease with shCaMKI $\gamma$ ) relative to control. Thus, the shRNAs phenocopied the inhibitory effects of STO-609, dnCaMKK, and dnCaMKI.

To ensure that the effects of shCaMKIγ are specific, we rescued the effect by cotransfection with human CaMKIy. Experiments in HEK293 cells showed that the shCaMKIy, designed against the rat CaMKIy nucleotide sequence, did not suppress expression of hCaMKIγ (data not shown). In Figure 5A, the representative images demonstrate that coexpression of hCaMKIy with shCaMKIγ significantly rescued the inhibition of both axon formation and length. Quantification of these data (Fig. 5B,C) demonstrate that only 63 ± 5% of shCaMKIγ-expressing neurons formed axons with average axonal length of 130.8  $\pm$  15  $\mu$ m versus 86 ± 7% of shCaMKIy plus hCaMKIy-expressing neurons that formed an axon with average axonal length of 200.6  $\pm$ 26.5 μm. Furthermore, if CaMKIγ regulates axon formation, transient overexpression of CaMKIy might increase percentage of neurons that exhibit an axon. To test this hypothesis, we transfected hippocampal neurons with wild-type EGFP-CaMKIγ, and fixed neurons at 24 h when most control neurons do not exhibit an axon (Fig. 2B). Hippocampal neurons expressing EGFP-CaMKIy showed a dramatic acceleration of axon formation at 24 h in culture. At 24 h, only 24.3  $\pm$  2.5% of control neurons had a defined axon compared with 56.4  $\pm$  2.2% of neurons expressing EGFP-CaMKI $\gamma$  (Fig. 5D,E). These axons were also significantly longer (171.8  $\pm$  8.6  $\mu$ m) than control neurons (85.3  $\pm$  5.5  $\mu$ m) (Fig. 5F). These data strongly support a role for CaMKI $\gamma$  in axon formation.

## TRPC5 channel plays a role in axon formation

Calcium plays multiple and complex roles in determining neuronal development both *in vitro* and *in vivo* (Mattson, 1992, 1999; Dotti and Poo, 2003; Jin et al., 2005; Zheng and Poo, 2007). The spatiotemporal frequency and amplitude of calcium oscillations trigger varying responses, mediated in part by CaM kinases and CaM-dependent protein phosphatases, which are critical for axon formation, outgrowth, and growth cone guidance as well dendritic arborization and synaptogenesis (Mattson and Kater, 1987; Gomez and Spitzer, 2000; Spitzer et al., 2000; Gomez and Zheng, 2006; Zheng and Poo, 2007). Our pharmacological, dominant-negative, and shRNA data establish a role for CaMKK and CaMKIγ in axon formation of cultured hippocampal neurons. Since the only known mechanism for activation of both of

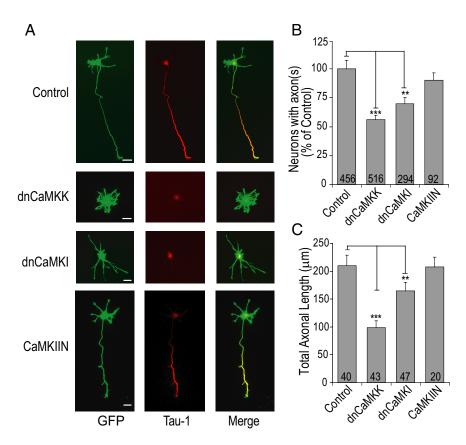
these enzymes is an increase in intracellular calcium, we investigated the source of this calcium upstream of CaMKK and CaMKIy in axon development. L- and N-type voltage-gated calcium channels are present in hippocampal neurons during early developmental stages (Pravettoni et al., 2000). Multiple recent studies also implicate the TRPC cation channel in regulation of growth cone morphology (Greka et al., 2003), neurite outgrowth (Wu et al., 2007), spinogenesis (Zhou et al., 2008), and as the source of calcium influx during chemotropic turning of growth cones in response to neurotrophins, netrin-1, and BDNF (Gomez, 2005; Li et al., 2005; Shim et al., 2005; Wang and Poo, 2005).

To determine the source of intracellular calcium elevation in neurites of stage 2 hippocampal neurons, we used pharmacological inhibitors. As shown in Figure 6A, the L-type channel blocker nifedipine (10  $\mu$ M) had no effect on the rate of axon formation in control neurons or those expressing caCaMKI. However, inhibition of TRPC channels with 3  $\mu$ M SKF96365 markedly blocked axon formation in control cells (34  $\pm$  4% polarized compared with 100% of control), and caCaMKI partially rescued axon formation (71  $\pm$  3% polarized) in the presence of SKF96365.

We next ascertained whether inhibition of TRPC channels also suppressed activation of CaMKI $\gamma$ . Because of its low abundance in young cultures, endogenously phosphorylated CaMKI $\gamma$  was not

detected by our phosphospecific CaMKI antibody. However, in neurons as well as heterologous systems, we are able to detect phosphorylation of transiently transfected CaMKI $\gamma$ . To monitor the phosphorylation of CaMKI $\gamma$ , we electroporated neurons with Flag-tagged CaMKI $\gamma$  and subsequently treated them 50  $\mu$ M SKF96365 or vehicle for 1 h. The activation state of Flag-CaMKI $\gamma$  by CaMKK was determined by Western blotting with antiphospho-CaMKI antibody. As shown in Figure 6B (top, representative blot; bottom, quantification), acute inhibition of TRPC channels strongly suppressed phospho-CaMKI $\gamma$ , but total Flag-CaMKI $\gamma$  remained constant. Thus, CaMKI $\gamma$  activation in stage 2 hippocampal neurons was dependent on TRPC channel activity.

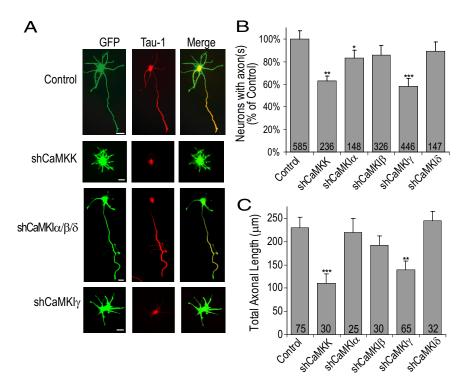
The TRPC channel family is composed of seven members (TRPC1-7) that assemble into homomeric or heteromeric channels with diverse properties including mode of activation, calcium conductance, and signaling partners (Clapham et al., 2001; Ambudkar, 2006; Ambudkar and Ong, 2007; Beech, 2007). Based on subunit assembly, TRPC channels are divided into distinct groups—TRPC3/6/7 heteromers, TRPC1/4/5 heteromers, and TRPC subunit homomers. It has been recently established that, by acting at extracellular sites close to the pore, TRPC4/5 heteromers and TRPC5 homomers are distinctively potentiated by the lanthanum (La<sup>3+</sup>) in the micromolar range (10–100  $\mu$ M) (Jung et al., 2003; Plant and Schaefer, 2003; Beech, 2007), a concentration that typically is used to suppress other TRPC and other calciumpermeable channels (Beedle et al., 2002). Because of this unique property of TRPC5 channels to be potentiated by La<sup>3+</sup>, we used it



**Figure 3.** dnCaMKK and dnCaMKI inhibit axon formation and decrease axonal length. **A**, E18 hippocampal neurons were coelectroporated with control (empty vector), dnCaMKK, dnCaMKI, or CaMKIIN containing plasmid and soluble EGFP before plating. **A** shows representative images of neurons fixed at 48 h and stained with the axonal marker Tau-1. **B**, Quantification of axon formation as measured by number of neurons with Tau-1-positive projections from three to four independent experiments. **C**, Quantification of average axonal length as measured by the length of the primary Tau-1-positive neurite. Scale bars: 20 μm (all panels).

as an experimental tool to test the role of calcium influx by TRPC5-containing TRPC channels in axon formation while inhibiting other calcium channels. When E18 neurons were incubated in media containing 100  $\mu$ M La  $^{3+}$ , there was an approximately twofold increase in neurons exhibiting axon(s) at 20–24 h in culture (Fig. 6C) as well as an approximately threefold increase in neurons with more than one axon (Fig. 6D). Both the La  $^{3+}$ -stimulated axon formation and multiple axon effects were strongly blocked by coincubation with the TRPC inhibitor, SKF96365. These data are similar to the effect of transfection with caCaMKI (Fig. 1). This observation with La  $^{3+}$  is supportive of a hypothesis that TPRC4 or -5 channels are a relevant source of Ca  $^{2+}$  influx during axon formation.

To further investigate the role of TRPC5 channels, we used siRNA techniques. TRPC5 channels are of particular interest in neuronal development as these channels are present in growth cones of stage 2 hippocampal neurons grown in culture and in hippocampus of intact brain (Okada et al., 1998; Strübing et al., 2001; Greka et al., 2003; Fowler et al., 2007) and are the most abundant TRPC channel in the hippocampus (Chung et al., 2006; Fowler et al., 2007). We specifically tested the involvement of TRPC5-containing TRPC channels in activation of CaMKIγ and subsequently axon formation with selective knockdown of this subunit with RNAi. As initially validated in COS7 cells, siTRPC5 specifically knocked down expression of Myc-TRPC5 without nonspecific targeting of related channel subunit TRPC6 (expressed as HA-TRPC6) (supplemental Fig. 1 E, available at www.



**Figure 4.** Knockdown of CaMKK and CaMKI $\gamma$  protein expression inhibits axonogenesis and decreases axonal length. **A**, E18 hippocampal neurons were electroporated before plating with pMU6pro alone (control) or pMU6pro containing shCaMKK, shCaMKI $\alpha$ , shCaMKI $\beta$ , shCaMKI $\beta$ , or shCaMKI $\gamma$ . Between 2 and 48 h in culture, low-density cultures of electroporated neurons were incubated with 20 μmSP600125 to suppress intrinsic neuronal polarization to achieve effective knockdown with the plasmid based shRNA (see Results). At 48–72 h in culture, electroporated neurons were released from SP600125 block by transfer of coverslips to new glial culture plates for an additional 48 h. Representative images of neurons fixed and stained with Tau-1 at 96 h are shown. The third from top panel shows staining from CaMKI $\alpha$  shRNA. This image is also representative of neurons transfected with shCaMKI $\beta$  or shCaMKI $\delta$ . **B**, Summary of three to five independent experiments. **C**, Average axonal length as measured by length of primary process containing Tau-1 staining from three independent experiments. Scale bars: 20 μm (all panels).

ineurosci.org as supplemental material). To determine the extent of endogenous TRPC5 knockdown, we nucleofected hippocampal neurons with siTRPC5 and harvested neurons at 24, 48, and 96 h after nucleofection. Immunoblotting with anti-TRPC5 antibody and anti- $\beta$ -tubulin (loading control) revealed a timedependent reduction of TRPC5 (percentage reduction: 24 h, 28%; 48 h, 46%; 96 h, 65%) (Fig. 6E). To specifically test the role of TRPC5 in activation of CaMKIy, control siRNA or siTRPC5 and Flag-CaMKIy were coelectroporated into cultured hippocampal neurons before plating and harvested on DIV 3. In Figure 6F, expression of siTRPC5 markedly reduced the level of pCaMKIγ by  $39.5 \pm 5.5\%$  compared with vector control. The extent of inhibition of CaMKI activation (40%) closely matched with the extent of TRPC5 suppression at 48 h (46%). These data support the results attained with the pharmacological inhibition of TRPC channels with SKF96365.

TRPC5 channels form functional channels by homo- or heterotetrameric assembly of individual subunits. It has been recently reported that the first ankyrin domain within the N terminus of TRPC5 is critical for the assembly of functional tetrameric channels (Schindl et al., 2008). Furthermore, Schindl et al. demonstrated that N-terminal fragments containing the first ankyrin repeat functioned as a dominant-negative species on TRPC4/TRPC5 heteromeric and homomeric channels by disrupting channel assembly resulting in downregulation of channel currents. We generated a dominant-negative TRPC5 construct based on these findings that included the N-terminal 330 aa containing the ankyrin repeats of mTRPC5, designated as TRPC5

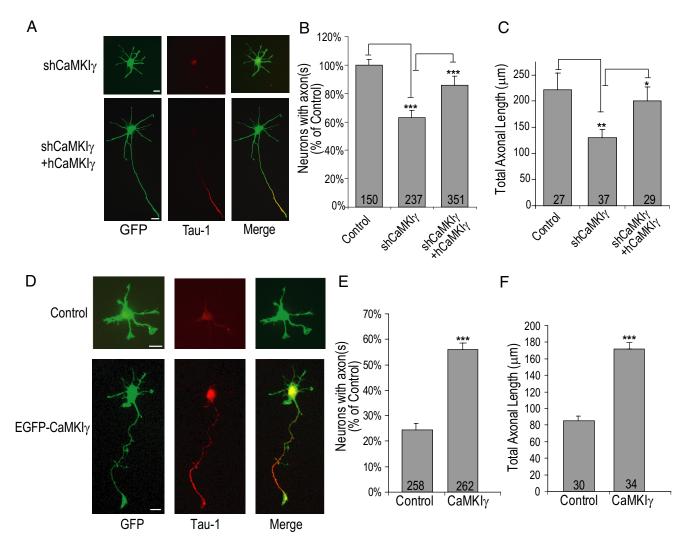
NT330. Transfection of this dominantnegative TRPC5 modestly but significantly reduced axon formation (72.5  $\pm$ 8% formed an axon compared with 100% for control) in hippocampal neurons at 48 h (supplemental Fig. 1D, available at www.jneurosci.org as supplemental material). We further evaluated the role of TRPC5 in the developmental progression from stage 2 to stage 3 hippocampal neurons by expressing RNAi to the TRPC5 subunit. We validated the knockdown of TRPC5 by immunocytochemistry with TRPC5-specific antibodies (as shown by Western blotting in Fig. 6E). As shown in Figure 7A, left panels, siTRPC5, but not control siRNA, reduced the anti-TRPC5 staining intensity throughout the neuron compared with neighboring nontransfected cell. The level of knockdown was variable between individual neurons, ranging from reductions of 30-70%, with an average reduction to  $56.5 \pm 5.8\%$  of the control level. These data confirmed the specificity of the anti-TRPC5 antibody used for immunocytochemistry.

Next, we evaluated the effect of siTRPC5 knockdown on axon formation. As shown in Figure 7*B*, transfection with siTRPC5 before plating revealed a significant reduction in the percentage of polarized neurons (siTRPC5, 57.2  $\pm$  4%, compared with control set at 100%). siTRPC5 also produced a small but signif-

icant decrease in axonal length of the remaining neurons that polarized in the presence of siTRPC5 (Fig. 7*B*, right bottom graph). A recent report indicates that TRPC6 channels are involved in synapse formation in mature (DIV 21) hippocampal neurons (Zhou et al., 2008). TRPC6 channel immunoreactivity peaks around P14 in hippocampus and is not detectable in E17 brain (Zhou et al., 2008) or DIV 1 hippocampal neurons (data not shown), so it is not likely to play a major role in axonogenesis. Indeed, siRNA to TRPC6, which was effective against TRPC6 but not TRPC5 in transfected COS7 cells (supplemental Fig. 1*E*, available at www.jneurosci.org as supplemental material), had no significant effect on axon formation under our conditions (Fig. 7*B*).

### Endogenous CaMKIγ and TRPC5 reside in lipid rafts

CaMKI $\gamma$  has a unique C-terminal extension containing multiple cysteines, not present in other CaMKI isoforms, that undergo prenylation and palmitoylation to target the  $\gamma$ -isoform to plasma membranes, Golgi and sarcoplasmic reticulum compartments (Takemoto-Kimura et al., 2003, 2007). Of the various TRPC channel members, TRPC1, TRPC3, and TRPC4 have recently been reported to localize within detergent-insoluble, lipid raft compartment in various non-neuronal cell lines (Remillard and Yuan, 2006). This compartmentalization modulates the functional properties of TRPC1. TRPC1 is a store-operated channel when in lipid rafts and a receptor-operated channel in nonraft plasma membranes (Alfonso et al., 2008; Alicia et al., 2008). These functional differences are likely attributable to selective protein–protein interactions that are compartment specific.



**Figure 5.** CaMKl γ expression promotes axon formation and increases axonal length. **A–C**, Coexpression of shRNA-insensitive human CaMKl γ rescues the effect of shCaMKl γ. Representative images are shown in **A**. Quantification of percentage of neurons exhibiting an axon and total axonal length from shCaMKl γ and shCaMKl γ plus rescue plasmid containing hCaMKl γ are shown in **B** and **C**, respectively. **D**, Representative images of EGFP-wtCaMKl γ induced accelerated axon formation at 24 h in culture. Controls contained soluble EGFP. **E**, **F**, Quantification of percentage of neurons exhibiting an axon at 24 h and total axonal length of neurons polarized at 24 h, respectively. Scale bars: 20 μm (all panels).

However, to date, there are very little data on the submembrane distribution of endogenous TRPC5 in developing brain.

We postulated that endogenous CaMKIy and TRPC5 channels may colocalize in a membrane microdomain to generate efficient coupling between calcium influx, activation of kinase, and highly localized downstream signaling to promote axonogenesis. Fractionation of P0 rat forebrains and Western blotting revealed endogenous CaMKK and CaMKI y were identified both in the lipid raft fraction 4 (marked by Flotillin-1) and nonraft membrane fraction 12 (marked by TfR) (Fig. 8A). The other isoforms, which are soluble, were not detectable in either fraction 4 or 12. Moreover, we found strong enrichment of TRPC5 in the lipid raft fraction (fraction 4), compared with the nonraft membrane fraction (fraction 12). In striking contrast with this result, TRPC6 is located almost completely in fraction 12. These are the first data to reveal distinct compartmentalization of TRPC5- and TRPC6-containing TRPC channels, which may allow distinct signaling of the different channel subtypes within different microdomains of the neurons. We performed immunocytochemistry to determine whether TRPC5 colocalized with the neuronal lipid raft marker, Flotillin-1. In cultured hippocampal neurons, TRPC5 (pseudocolored red) partially colocalizes with Flotillin-1

(pseudocolored green) particularly in the central region of the growth cone, with lesser colocalization along the length of the neurite. We also evaluated colocalization of EGFP-CaMKIy with the well established lipid raft marker GM1 as detected with CTxB. CTxB selectively binds the GM1 and subsequently clusters GM1, thus permitting live labeling of the lipid raft marker (Harder et al., 1998; Nichols et al., 2001; Hullin-Matsuda and Kobayashi, 2007). We observed significant colocalization between EGFP-CaMKIy and the raft marker, GM1, particularly in the axonal growth cone (Fig. 8C, arrows). We also tested whether the GM1containing lipid rafts colocalized with the raft marker, Flotillin-1. We found overlap of Flotillin-1 and CTxB throughout the cell and particularly in and the axonal growth cone (arrow) of the nascent axon in the stage 2+ and stage 3 hippocampal neurons (supplemental Fig. 1 F, available at www.jneurosci.org as supplemental material). Since CaMKI $\gamma$  and TRPC5 biochemically cofractionate in the low-density raft fraction, we also confirmed their colocalization in cultured hippocampal neurons. Since we do not have an antibody specific for detecting endogenous CaMKIy by immunocytochemistry, we electroporated hippocampal neurons with EGFP-CaMKIy, before plating, and fixed at 24 h, followed by immunostaining with anti-TRPC5 an-

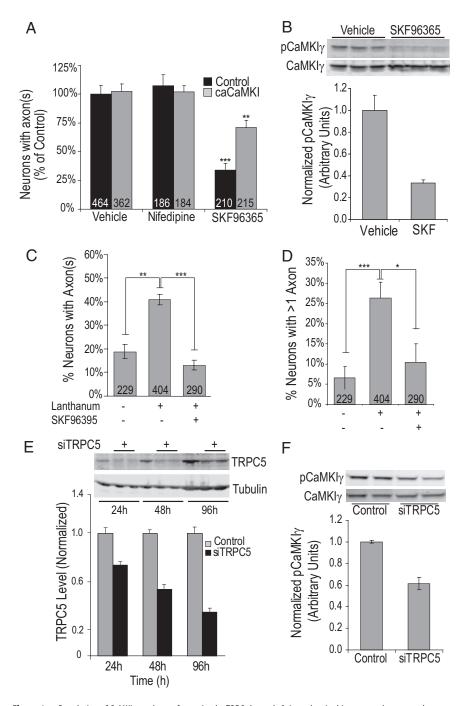


Figure 6. Regulation of CaMKI  $\gamma$  and axon formation by TRPC channel. A, Low-density hippocampal neurons electroporated before plating were incubated for 46 h (2 h after plating until fixation at 48 h) in culture with vehicle, the L-type channel blocker nifedipine (10 µm), or the TRPC blocker SKF96395 (3 µm). Neurons fixed at 48 h were stained with the axonal marker Tau-1. Quantification of results from three independent experiments is shown. B, Representative blot showing regulation of CaMKly by TRPC antagonist, SKF96365. P0 hippocampal neurons were electroporated with Flag-CaMKI $\gamma$  before plating and 24 h later exposed to 50  $\mu$ m SKF96365 for 60 min. Lysates were immunoblotted with antiphospho-CaMKI antibody (top blot) or anti-Flag antibody to detect total CaMKI γ (bottom blot). The bottom graph shows average normalized phospho-CaMKl $\gamma$ . C, D, Graphs depict quantification of axon formation (C) and induction of multiple axons ( $\boldsymbol{p}$ ) in E18 neurons that were incubated for 20 h (from 2 to 22 h) with growth medium containing 100  $\mu$ M lanthanum chloride, a selective activator of TRPC4 and -5 channels. E, High-density hippocampal neurons were electroporated with 2.5  $\mu g$  of siTRPC5 and harvested at the indicated times with SDS sample buffer. Western blots show the time course of reduction of endogenous TRPC5 (top blot) and  $\beta$ -tubulin (bottom blot) with siTRPC5. The bottom graph shows quantification of TRPC5 levels relative to tubulin at 24, 48, 72 h (with control at 24 h set to arbitrary unit of 1).  $\bf F$ , Flag-CaMKI $\gamma$  and control siRNA or siTRPC5 were coelectroporated in P0 hippocampal neurons and harvested after 48 – 72 h in SDS sample buffer. Phospho-CaMKI  $\gamma$  and total CaMKI  $\gamma$  (Flag) were detected by multiplexed Western blotting and the bottom graph represents summary of ratio of pCaMKl $\gamma$ /total CaMKl $\gamma$  from four independent experiments.

tibody and anti- $\beta$ -tubulin antibodies. We observed prominent colocalization of EGFP-CaMKI $\gamma$  and endogenous TRPC5 in growth cones of stage 2 neurons as well as in the axonal growth cone of stage 3 neurons (Fig. 8D; insets show growth cone). These data confirm the biochemical fractionation results, suggesting CaMKI $\gamma$  and TRPC5 may reside in close proximity in the submembrane, lipid raft domain.

Localization of transfected EGFP-CaMKIy in lipid raft fractions in cultured cortical neurons was dependent on the prenylation/palmitoylation of its unique C terminus—mutational deletion of these lipid modification sites ablated the incorporation of transfected CaMKI vinto lipid rafts and its ability to enhance dendritic outgrowth (Takemoto-Kimura et al., 2007). If the lipid modification-mediated targeting of CaMKIy to plasma membrane and submembrane compartments such as lipid rafts, is important for its role in axon formation, deletion of these lipid modification sites should likewise obviate the ability of transfected CaMKIy to accelerate axonogenesis (Fig. 5E,F). Therefore, we truncated CaMKIγ at residue M407 to remove its unique C terminus containing the prenylation/palmitoylation sites (CaMKI $\gamma_{\Delta PP}$ ) (supplemental Fig. 1G, available at www.jneurosci.org as supplemental material). When hippocampal neurons were transfected with CaMKI $\gamma_{\Lambda PP}$ , there was no increase in the formation of axons in contrast to wild-type CaMKIγ (Fig. 8 E). This result leads to the conclusion that membrane localization of CaMKIy, possibly in the lipid raft fraction, is important for axon formation.

Wild-type CaMKIγ requires binding of Ca<sup>2+</sup>/CaM for full activation. Therefore, we also tested the ability of wild-type CaMKIy to accelerate axon formation with reduced calcium contribution of TRPC5-containing channels in activation of CaMKIy, by cotransfecting siTRPC5 with CaMKI $\gamma$  and fixing neurons at 24 h. As shown in Figure 8F, reduction of TRPC5 levels by siRNA significantly reduced the ability of CaMKIγ to accelerate axon formation, suggesting that the calcium influx from TRPC5 is important for activation of CaMKI v during this process. In contrast, a caCaMKIγ construct with a mutant CaM binding domain that renders its catalytic activity independent of Ca<sup>2+</sup>/CaM binding was able to rescue the suppression of axon formation by siTRPC5 (Fig. 8G). The simplest interpretation of these data is that the calcium influx mediated by TRPC5 channels in developing

hippocampal neurons is required for full activation of CaMKI and its subsequent acceleration of axon formation. Since CaMKI and TRPC5 seem to biochemically codistribute and colocalize to detergent-resistant submembrane (lipid raft) compartment, together they may efficiently control signaling dynamics in a highly restricted way to facilitate axon formation.

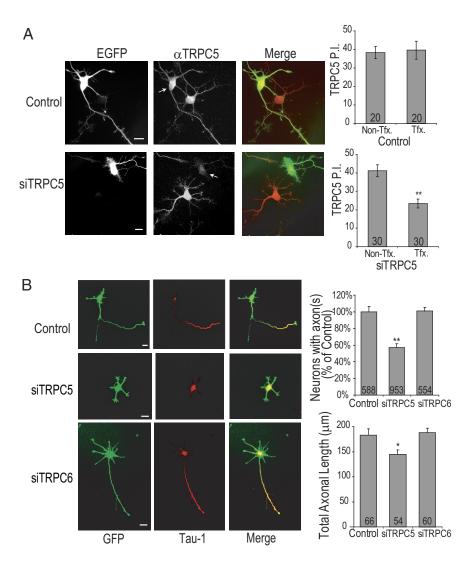
## Discussion

Despite extensive evidence for a the role of calcium signaling in axonal outgrowth and guidance in cultured neurons, relatively little is known about signaling downstream of calcium during the initial steps of neuronal polarization defined by formation of the axon. In the current study, we demonstrate that activation of CaMKK and its downstream target CaMKIy promote axon formation in cultured hippocampal neurons. Treatment with the CaMKK pharmacological blocker, STO-609, and suppression of CaMKK or CaMKI by expression of dominant-negative constructs and shRNA resulted in 40-50% reduction in the number of embryonic hippocampal neurons that elaborated a morphologically and biochemically defined axon. Moreover, modest overexpression of CaMKIy resulted in pronounced acceleration of axon formation.

CaMKIγ appears to be involved in several aspects of neuronal development as we previously reported a role for CaMKIγ in activity-dependent dendritic arborization of hippocampal neurons (Wayman et al., 2006). This conclusion was extended to demonstrate a role CaMKIγ in BDNF-stimulated dendritic growth of cortical neurons (Takemoto-Kimura et al., 2007). Takemoto-Kimura et al. did not observe any effect of CaMKIγ knockdown by shRNA on axonal length in high-density cortical neurons and they did not measure

the percentage of polarized stage 2/3 neurons. Measurement of neuronal polarity and axonal length of individual neurons would likely be challenging in these high-density cultures because of the extent of their axonal arborization. We used the low-density E18 hippocampal ("glial sandwich") culture system that is optimal for investigation of neuronal polarity (Craig and Banker, 1994; Bradke and Dotti, 2000; Kaech and Banker, 2006). This allowed us to measure the percentage of polarized neurons as well as measure their axonal length with a high degree of confidence because of the clear separation of neurons. Furthermore, our conclusion that CaMKI $\gamma$  is involved in axon formation is based on multiple, independent techniques.

We also document a novel role for TRPC channels during axon formation. In our experiments, incubation with the TRPC inhibitor SKF96365 blocked activation of CaMKI $\gamma$  and strongly suppressed axon formation—this latter effect was rescued by caCaMKI. As we (Fig. 8D) and others (Greka et al., 2003) demon-



**Figure 7.** Regulation of axon formation by TRPC5 channels. **A**, E18 Hippocampal neurons were electroporated before plating with control siRNA or siTRPC5 plus soluble EGFP. Neurons were fixed at 48 h after electroporation and stained with anti-TRPC5 polyclonal antibody and imaged as described in Materials and Methods. Representative images from control siRNA (top) and siTRPC5 (bottom) are shown in left panel and the graphs summarizing quantification ( *y*-axis shows TRPC5 P.I. in arbitrary units) are shown on right. **B**, E18 neurons were electroporated before plating with control siRNA, siTRPC5, or siTRPC6. Neurons were fixed at 48 h and evaluated for axon formation by staining with anti-Tau-1 antibody (see Materials and Methods). Representative images are shown in left panel(s) and quantification of percentage of polarized neurons (top graph) and length of (bottom graph) in the neurons that did polarize is shown on right. Scale bars: 20 μm (all panels).

strate, TRPC5 channels are expressed in the cell body and axonal growth cones of stage 2 and stage 3 cultured hippocampal neurons. These results suggest that under basal conditions TRPC5 channels, which can exhibit significant spontaneous activity (Yamada et al., 2000), may mediate calcium influx that activates CaMKK and subsequently CaMKIy. Furthermore, micromolar La<sup>3+</sup>, known to inhibit most Ca<sup>2+</sup>-permeable channels including TRPC channels except for TRPC4 and 5, strongly stimulated axon formation (Fig. 6B,C). It has been previously shown that short-term neurite outgrowth (1-3 h duration) was stimulated by 10  $\mu$ M La<sup>3+</sup>, and this effect required extracellular calcium (Mattson and Kater, 1987). Last, expression of dominantnegative N-terminal fragment of TRPC5, which disrupts functional channel assembly at plasma membrane, as well siRNA suppression of TRPC5, but not TRPC6, before plating caused a decrease ( $\sim$ 42%) in the number of neurons that formed an axon and reduced axonal length in the remaining neurons. Recently,

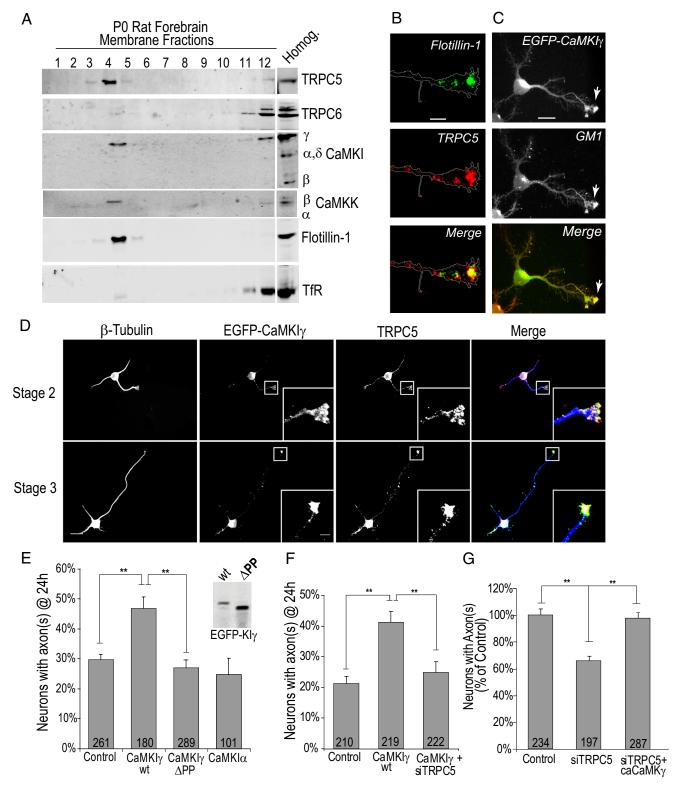


Figure 8. CaMKI $\gamma$  and TRPCS colocalize with lipid raft markers in rat brain and cultured hippocampal neurons. A, Endogenous CaMKI $\chi$ , and TRPCS are present in the lipid raft fraction (fractions 4) as confirmed by enrichment of the raft marker Hotillin-1 in fraction 4 of a discontinuous sucrose density gradient (fractions 1– 4,5% sucrose; fractions 5–11,35% sucrose; fraction 12,50% sucrose) from PO rat forebrains. The input is the crude homogenate before any centrifugation steps separating cytosol and membranes. B, Immunofluorescent staining of endogenous TRPCS (pseudocolored red) and endogenous Flotillin-1 (pseudocolored green) in axonal growth cone of E18 hippocampal neuron. Scale bars:  $5 \mu m$ . C, Colocalization of Alexa Fluor 546-CTXB labeled raft marker GM1 (as described in Materials and Methods) and EGFP-CaMKI $\gamma$  in DIV 1 hippocampal neurons. Scale bars: D, 20  $\mu$ m. Immunofluorescent staining of endogenous  $\beta$ -tubulin, TRPC5 with electroporated EGFP-CaMKI $\gamma$  in stage 2 (top row) and stage 3 (bottom row) neurons shows extensive colocalization of TRPC5 and EGFP-CaMKI $\gamma$ . In final column, the images are a merge of all three channels, with  $\beta$ -tubulin pseudocolored blue, EGFP-CaMKI $\gamma$  pseudocolored green, and endogenous TRPC5 pseudocolored red in stage 2 and stage 3 hippocampal neurons. The insets show additional magnification of largest growth cones. Scale bars:  $\beta$ -tubulin (bottom) panel, 10  $\mu$ m; EGFP-CaMKI $\gamma$  (bottom inset showing growth cone), 5  $\mu$ m. E, Quantification of neurons with Tau-1-positive axons expressing wild-type CaMKI $\gamma$ , CaMKI $\gamma$ , alone or wild-type CaMKI $\gamma$  plus siTRPC5 at 24 h. Note, in D and D, neurons are fixed at 24 h to test mutant CaMKI $\gamma$  and siTRPC5 in acceleration of axon formation assay. D, Quantification of rescue experiment from neurons expressing control, siTRPC5 plus rescue with caCaMKI $\gamma$ .

Greka et al. (2003) showed that transfection of a dominant-negative TRPC5 (dnTRPC5-EGFP) construct at DIV 1 resulted in increased neurite and filopodial length (Greka et al., 2003). It is unclear whether Greka et al. evaluated neuronal polarity or measured axonal (vs neurite) length in stage 2/3 hippocampal neurons. There may be differences in the signaling responses to TRPC5-mediated localized calcium influx on the outgrowth of neurites versus the axon in stage 2 or stage 3 neurons. For instance, in NG108 neuroblastoma cells and primary cerebellar granule neurons, early stages of neurite initiation and differentiation require calcium influx via TRPC5 channels; however, after differentiation, both cell types exhibited a sharp decline in TRPC5 levels (Wu et al., 2007). Our data using multiple approaches implicate a role for TRPC5 channels in the development of axons in low-density cultured hippocampal neurons.

We also show, by biochemical fractionation, the colocalization of CaMKK, CaMKIy, and TRPC5 in the low-density lipid raft fraction, identified the specific marker Flotillin-1. These biochemical data were supported by immunostaining for endogenous TRPC5 and Flotillin-1 in DIV 1 hippocampal neurons. Another lipid raft marker, GM1, also colocalized with EGFP-CaMKIy in these neurons. Targeting of signaling molecules to lipid rafts results in microdomains that serve to enrich and spatially restrict the signaling from membrane receptors/ion channels to the downstream effectors, thereby creating localized signal amplification. Various studies indicate that the segregation of signaling molecules to rafts may be an important mechanism during axon guidance and growth (Kamiguchi, 2006). TRPC5 channels in lipid rafts may mediate spatially restricted calcium entry into the growth cone to locally activate CaMKK/CaMKIγ during axon formation. Consistent with this model, transfection with the CaMKIy truncation mutant that lacked the lipid modification sites necessary for its plasma membrane and lipid raft targeting, did not accelerate axon formation similar to wild-type CaMKIy.

Recently, Bezzerides et al. (2004) demonstrated that treatment of DIV 1 hippocampal neurons with NGF or BDNF significantly increased the number of functionally available surface TRPC5 channels by membrane insertion from an intracellular vesicular pool. This increase in TRPC5 surface expression was dependent on PI3 kinase and Rac1 activity. In addition, Wu et al. (2007) showed that integrin receptor-mediated calcium influx via activation of TRPC5 channels was required for differentiation of NG108 cells and cerebellar granule neurons. In this case, TRPC5 channel activation and subsequent neurite outgrowth were also dependent on phospholipase Cy (PLCy) and PI3 kinase signaling. Neurotrophins such as BDNF and NGF, and guidance factors such as Netrin-1 that are coupled with PLCy, PI3 kinase, and Rac1 signaling, are well established modulators of neuronal polarization in vivo and in vitro (Estrach et al., 2002; Da Silva et al., 2005; Barnes et al., 2007; Shelly et al., 2007). This suggests a model in which asymmetric presentation of axonal guidance or growth cues, such as BDNF or Netrin-1, may increase the activity of existing surface TRPC5 channels via PLCy/DAG (diacylglycerol) (Beech, 2007) and/or also enhance the number of surface TRPC5 channels by membrane insertion (Bezzerides et al., 2004). In cultured neuron model systems, because of the presumed uniform extracellular milieu, the symmetry breaking event that is required for neuronal polarization is thought to occur stochastically (Andersen and Bi, 2000; Fivaz et al., 2008). Therefore, in cultured hippocampal neurons, TRPC5 channel activation in the growth cone of one neurite may be stochastically enhanced by increased surface expression or by higher spontaneous channel activity.

The subsequent increase in spatially restricted intracellular calcium and activation of CaMKK/CaMKI $\gamma$  may alter cytoskeletal or other signaling dynamics favoring accelerated growth of this neurite and eventual axon formation.

Because of the lack of a phosphospecific antibody that only detects the CaMKK-phosphorylated  $\gamma$ -isoform of CaMKI, we were unable to demonstrate enrichment of phospho-CaMKIγ in a single neurite of a fixed stage 2 neuron with the assumption that this neurite is the presumptive future axon. This conclusion has been made for several other signaling proteins that are putative "specificity determinants." However, interpretation of such observations in fixed stage 2 cells is questionable. There are only two examples of live-cell imaging of such proteins in hippocampal neurons during stage 2 and their subsequent progression into stage 3—Kif5c and shootin1. For both, Kif5C (Jacobson et al., 2006) and shootin1 (Toriyama et al., 2006), it is clear that these "specificity proteins" do not stably localize in one neurite but show fluctuating accumulation in multiple stage 2 neurites before eventually stably associating with the axon in stage 3 (see movie S6, www.cell.com/neuron/supplemental/S0896-6273(06)00095-X, and video 2 in http://jcb.rupress.org/cgi/content/full/jcb.200604160/ DC1). Thus, based on these findings, we conclude that that demonstrating selective enrichment of a signaling protein in one neurite of a fixed stage 2 neuron is not a definitive or predictive measurement of the future axon. Only repetitive imaging beginning at stage 2 and transitioning through stage 3 can identify which neurite will become the axon. However, the development of the necessary tools and technology to image activation of the CaMKK/KI cascade goes beyond the scope of this paper.

There are multiple potential downstream targets of CaMKI that might regulate axon formation. One possibility is the small G-protein Rac1. Activation of Rac1 may be critical for the localized suppression of RhoA and for activation of Pak1, both signaling pathways previously shown to be important for determination of neuronal polarity (Nishimura et al., 2005; Mertens et al., 2006; Watabe-Uchida et al., 2006; de Curtis, 2008). It was recently proposed that CaMKIy may regulate Rac1 as shRNA knockdown of CaMKIy reduced GTP loading of Rac1 in 3-d-old cortical neurons (Takemoto-Kimura et al., 2007). The Rac1-guanine exchange factor (GEF) βPIX can be activated by CaMKK/CaMKI signaling (Saneyoshi et al., 2008), and Tiam1 and Stef, other Rac1 GEFs, are also potential mediators (Takemoto-Kimura et al., 2007). Therefore, it is possible that increased localized activation of CaMKIy because of calcium influx via the TRPC5 channel will allow for RacGEF-mediated modulation of Rac1 signaling in the growth cone of one neurite that is essential for elongation of this neurite. Microtubules and local microtubule stabilization in one neurite are suggested to be a physiological signal that specifies initial neuronal polarity (Witte et al., 2008). Microtubule polymerization is weakened by sequestration of tubulin because of binding of depolymerizing factors such as stathmin family members. Interaction of stathmin with tubulin is suppressed by phosphorylation of Ser16 by CaMKIV, thereby favoring microtubule polymerization (Melander Gradin et al., 1997). In cotransfected HEK293 cells or *in vitro* with purified proteins, CaMKI can also phosphorylate Ser16 (M. A. Davare and G. A. Wayman, unpublished observation), so CaMKIy may regulate microtubule assembly by this mechanism.

Over the past few years, significant effort has led to the identification of multiple signaling molecules, including PI3 kinase, the Par3/Par6/aPKC complex, CRMP2 and LKB1 kinases, which regulate axon specification (Yoshimura et al., 2006; Barnes et al., 2007; Shelly et al., 2007). Our study is one of the first to demon-

strate a calcium-dependent signaling pathway that directly affects axon formation in cultured embryonic hippocampal neurons. It will be of interest to understand whether and how the CaMKK cascade impinges on or has positive cross talk with any of the previously described pathways regulating axon formation. Our future effort will be directed at determining the downstream targets and signaling pathways modulated by CaMK $\gamma$  that are responsible for the acceleration of axon formation.

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