

# Growth Factors Mobilize Multiple Pools of $K_{Ca}$ Channels in Developing Parasympathetic Neurons: Role of ADP-Ribosylation Factors and Related Proteins

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**Chae, Kwon-Seok, Kwang-Seok Oh, and Stuart E. Dryer.** Growth factors mobilize multiple pools of  $K_{Ca}$  channels in developing parasympathetic neurons: role of ADP-ribosylation factors and related proteins. *J Neurophysiol* 94: 1597–1605, 2005. First published April 20, 2005; doi:10.1152/jn.00296.2005. In developing ciliary ganglion (CG) neurons, movement of functional large-conductance (BK type)  $Ca^{2+}$ -activated  $K^+$  ( $K_{Ca}$ ) channels to the cell surface is stimulated by the endogenous growth factors TGF $\beta$ 1 and  $\beta$ -neuregulin-1 (NRG1). Here we show that a brief NRG1 treatment (0.5–1.5 h) mobilizes  $K_{Ca}$  channels in a post-Golgi compartment, but longer treatments (>3.5 h) mobilize  $K_{Ca}$  channels located in the endoplasmic reticulum or Golgi apparatus. Specifically, the effects of 3.5 h NRG1 treatment were completely blocked by treatments that disrupt Golgi apparatus function. These include inhibition of microtubules, or inhibition of the ADP-ribosylation factor-1 (ARF1) system by brefeldin A, by over-expression of dominant-negative ARF1, or over-expression of an ARF1 GTPase-activating protein that blocks ARF1 cycling between GTP- and GDP-bound states. These treatments had no effect on stimulation of  $K_{Ca}$  evoked by 1.5 h treatment with NRG1, indicating that short-term responses to NRG1 do not require an intact Golgi apparatus. By contrast, both the acute and sustained effects of NRG1 were inhibited by treatments that block trafficking processes that occur close to the plasma membrane. Thus mobilization of  $K_{Ca}$  was blocked by treatments that inhibit ADP-ribosylation factor-6 (ARF6) signaling, including overexpression of dominant-negative ARF6, dominant-negative ARNO, or dominant-negative phospholipase D1. TGF $\beta$ 1, the effects of which on  $K_{Ca}$  are much slower in onset, is unable to selectively mobilize channels in the post-Golgi pool, and its effects on  $K_{Ca}$  are completely blocked by inhibition of microtubules, Golgi function and also by plasma membrane ARF6 and phospholipase D1 signaling.

## INTRODUCTION

The intrinsic membrane properties of excitable cells are a highly differentiated trait that emerges over an extended period of time during the development of postmitotic cells. For several years, we have studied the regulation of this process in identified populations of vertebrate neurons, especially in the parasympathetic neurons of the chick ciliary ganglion (Dryer et al. 2003). In chick ciliary neurons, significant functional expression of large-conductance (BK-type)  $Ca^{2+}$ -activated  $K^+$  ( $K_{Ca}$ ) channels occurs relatively late in embryonic development at stages that coincide with the formation of synapses with target tissues (E8-13) (Dourado and Dryer 1992).

The expression of ciliary neuron  $K_{Ca}$  channels requires inductive interactions with target tissues in the eye and with midbrain preganglionic neurons (Dourado et al. 1994) that are mediated by TGF $\beta$ 1 (Cameron et al. 1998) and  $\beta$ -neuregulin-1 (NRG1) (Cameron et al. 2001; Subramony and Dryer 1997), respectively. The effects of TGF $\beta$ 1 and NRG1 on ciliary neuron  $K_{Ca}$  channels persist when protein synthesis is blocked (Cameron et al. 1998; Chae et al. 2005; Lhuillier and Dryer 2000, 2002; Subramony and Dryer 1997; Subramony et al. 1996) and are associated with movement of preexisting  $K_{Ca}$  channels into the plasma membrane (Chae et al. 2004; Lhuillier and Dryer 2002). However, the mechanisms of growth factor-evoked  $K_{Ca}$  trafficking are poorly understood.

We have observed that NRG1 can evoke a robust increase in functional  $K_{Ca}$  in as little as 30 min (Chae et al. 2005), whereas a reliable response to TGF $\beta$ 1 is not usually observed in <6 h (Subramony et al. 1996). By contrast, both growth factors can activate the PI3K/Akt signaling cascades in a matter of minutes. One possible explanation for the different time courses of the physiological responses to the two factors is that NRG1 and TGF $\beta$ 1 cause mobilization of different pools of  $K_{Ca}$  channels. In the present study, we test that hypothesis.

The trafficking of membrane proteins is orchestrated by a host of small GTPases, including members of the ADP-ribosylation factor (ARF) subfamily (Donaldson 2003; Spang 2002; Takai et al. 2001). Two phylogenetically conserved members of this subfamily, ARF1 and ARF6, have received extensive attention in the context of membrane trafficking. ARF1 and its associated proteins regulate the budding and fusion of vesicles within the Golgi apparatus, in part by recruiting coatamer protein I to Golgi membrane surfaces (Presley et al. 2002; Stamnes 2002). By contrast, ARF6 regulates the movement of vesicles in and out the plasma membrane and mediates changes in cortical actin dynamics and membrane lipid composition that are associated with protein trafficking and cell motility (Donaldson 2003). As with other small GTPases, ARF1 and ARF6 exist in GTP-bound (active) and GDP-bound (inactive) states, and normal function requires that they cycle between these states. Moreover, ARF1 and ARF6 are regulated by a variety of guanine nucleotide exchange factors (GEFs), which mediate ARF activation, and GTPase-activating proteins (GAPs), which stimulate the intrinsic GTPase activity of ARFs and thereby terminate their action.

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In the present study, we show that  $K_{Ca}$  channels can be mobilized from multiple intracellular pools. These include a post-Golgi pool that can be rapidly mobilized in response to NRG1 by a process that requires activation of ARF6 but that does not require activation of ARF1, microtubules, or an intact Golgi apparatus. A different pool of  $K_{Ca}$  channels is mobilized more slowly and in a more sustained manner in response to TGF $\beta$ 1, as well as in response to higher concentrations of NRG1 (Chae et al. 2005). Mobilization of these  $K_{Ca}$  channels into the plasma membrane requires an intact Golgi apparatus based on the requirement for activation of ARF1 and related proteins and functional microtubular transport.

## METHODS

### *Cell isolation, culture, and transfection*

Chick ciliary ganglion neurons were isolated at embryonic day 9 (E9) and cultured in a medium containing 20 ng/ml recombinant rat ciliary neurotrophic factor as described previously (Cameron et al. 1998, 1999, 2001). Recombinant human TGF $\beta$ 1 and NRG1 were obtained from R&D Systems (Minneapolis, MN). For some experiments, colchicine, nocodazole, and brefeldin-A were obtained from Calbiochem (San Diego, CA). Cells were pretreated with these agents for 30 min before the addition of TGF $\beta$ 1 and NRG1, grown for various times as indicated in figure legends, and then used for electrophysiology. In other experiments, cells were transfected using biolistic methods (Lhuillier and Dryer 2003) and then used for electrophysiology or confocal microscopy. For transfection, E9 neurons were dissociated, cultured for 12 h, and plasmids were introduced into neurons using a biolistic particle-delivery system (PSD-1000; Bio-Rad, Hercules, CA). Plasmids were precipitated onto 1.0- $\mu$ m gold beads according to the manufacturer's protocol, and a helium shock wave with a pressure gradient of 650 psi was used to accelerate the coated beads onto cultured CG neurons. In these experiments, plasmids encoding dominant-negative or constitutively active mutants of signaling proteins were co-transfected with *Renilla* green fluorescent protein (GFP) (Stratagene, La Jolla, CA) at a ratio of 1:1. Plasmids encoding ARF6 (T27N), ARF1 (T31N), and ARF6 (Q67L) were provided by Dr. Julie G. Donaldson (NHLBI, National Institutes of Health, Bethesda, MD). A dominant-negative form of ARNO (E156K) was obtained from Dr. James E. Casanova (University of Virginia Health Sciences Center, Charlottesville, VA). Plasmids encoding phospholipase D1 (PLD1 K898R) and wild-type PLD1 were obtained from Dr. Michael Frohman (University Medical Center at Stony Brook, Stony Brook, NY). GFP-ARF-GAP273, in which sequences encoding 273 carboxyl-terminal noncatalytic amino acids of ARF-GAP1 were joined to the carboxyl-terminus of GFP, were obtained from Dr. Michael G. Roth (University of Texas Southwestern Medical Center, Dallas, TX).

### *Electrophysiology*

Whole cell recordings of  $K_{Ca}$  and voltage-activated  $Ca^{2+}$  currents were made using methods that we have standardized in many previous studies on ciliary neurons (Cameron et al. 1998, 1999, 2001; Chae and Dryer 2005; Chae et al. 2005; Dourado and Dryer 1992; Dryer et al. 1991; Lhuillier and Dryer 2000, 2002, 2003; Subramony and Dryer 1997; Subramony et al. 1996). Briefly, 25-ms depolarizing steps to 0 mV were applied from a holding potential of  $-40$  mV in normal and nominally  $Ca^{2+}$ -free salines containing 250 nM tetrodotoxin, and the net  $Ca^{2+}$ -dependent currents were obtained by digital subtraction using Pclamp software (Axon Instruments). Ciliary neurons are nearly spherical, and surface areas were calculated from cell diameters measured in two orthogonal axes and used to estimate current density. Similar results were obtained when currents were normalized to cell

capacitance estimated as described in Martin-Caraballo and Dryer (2002). Recording electrodes were made from thin-wall borosilicate glass (3–4 M $\Omega$ ) and filled with a solution consisting of (in mM) 120 KCl, 2 MgCl $_2$ , 10 HEPES-KOH, and 10 EGTA, pH 7.2. This solution buffers the bulk cytosolic levels of  $Ca^{2+}$  to low levels and prevents slow changes in intracellular  $Ca^{2+}$  associated with longer distance diffusion, such as diffusion that would occur between endosomal stores and plasma membranes but does not effectively buffer rapid  $Ca^{2+}$  changes in the immediate vicinity of  $Ca^{2+}$  channels. Normal external salines for measurements of  $K_{Ca}$  contained (in mM) 145 NaCl, 5.4 KCl, 0.8 MgCl $_2$ , 5.4 CaCl $_2$ , 5 glucose, and 13 HEPES-NaOH, pH 7.4. Voltage-activated  $Ca^{2+}$  currents were analyzed the same way except that KCl in the recording pipettes was replaced with CsCl as described previously (Dourado and Dryer 1992; Dourado et al. 1994; Lhuillier and Dryer 2000, 2002). Throughout, error bars represent SE. Data were analyzed by one-way ANOVA followed by post hoc analysis using Tukey's honest significant difference test for unequal  $n$  using Statistica software, with  $P < 0.05$  regarded as significant. In every experiment, data were collected from a minimum of two platings of ciliary ganglion neurons (i.e., from multiple cultures).

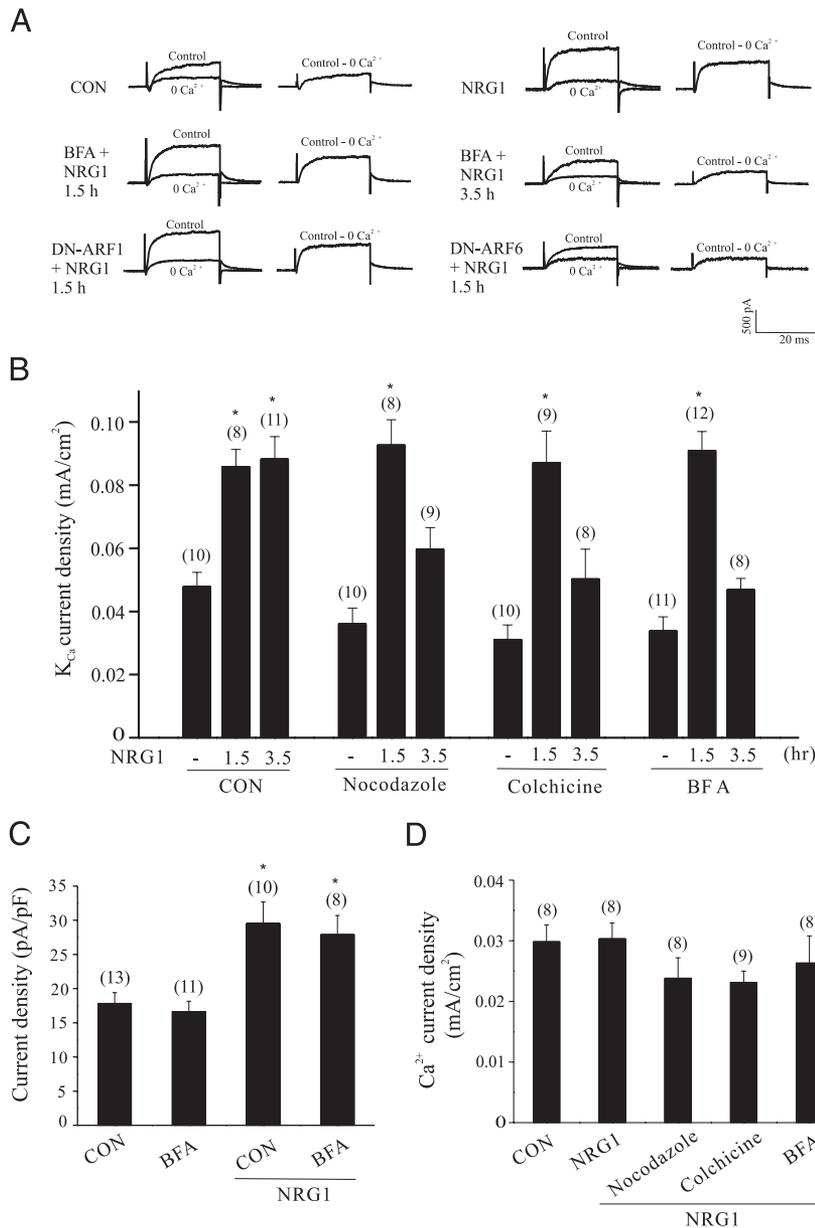
### *Golgi staining*

The Golgi apparatus was visualized using the fluorescent marker *N*-([4-[4, 4-difluoro-5-(2-thienyl)-4-bora-3a,4a-diaza-*s*-indacene-3-yl]phenoxy]acetyl) sphingosine (BODIPY TR C $_5$  ceramide) as described by Gangalum et al. (2004). This probe is commercially available as a complex with bovine serum albumin (Molecular Probes, Eugene, OR). Cells were treated with neuregulin in the presence or absence of brefeldin A for 1.5 or 3.5 h, washed in PBS, fixed in 4% paraformaldehyde for 6 min, blocked in PBS containing 1% BSA and 0.1% Triton X-100 for 30 min, and then stained with 15  $\mu$ M BODIPY TR C $_5$  ceramide for 30 min in the dark. Cells were then rinsed, mounted, and the integrity of the Golgi determined by fluorescence microscopy. The excitation wavelength was 580 nm and emission was monitored at 620 nm using a  $\times 60$  objective and an Olympus confocal microscope.

## RESULTS

### *Role of the Golgi apparatus, microtubules and ARF1 in NRG1 stimulation of $K_{Ca}$*

Previous studies from our laboratory (Lhuillier and Dryer 2002) showed that the stimulatory effects of TGF $\beta$ 1 on  $K_{Ca}$  expression require an intact Golgi apparatus because the effects of this growth factor were blocked by the Golgi inhibitor brefeldin-A as well as by inhibitors of microtubules (Lhuillier and Dryer 2002; see also Hamm-Alvarez and Sheetz 1998). TGF $\beta$ 1-evoked mobilization of  $K_{Ca}$  is a relatively slow process and is best observed  $\geq 5$  h after the onset of treatment. Functional mobilization of ciliary  $K_{Ca}$  channels by NRG1 is much more rapid and can occur in as little as 30–90 min (Chae et al. 2005). Therefore in the initial experiments of the present study, we examined the role of microtubules and the Golgi apparatus in NRG1-evoked trafficking of  $K_{Ca}$  channels. We quantified increases in macroscopic  $K_{Ca}$  by means of standard whole cell recordings (Fig. 1A) immediately after 1.5- or 3.5-h exposures to 10 nM NRG1. We observed that pretreatment with the microtubule inhibitors nocodazole (10  $\mu$ M) or colchicine (5  $\mu$ M) for 30 min prior to the onset of NRG1 exposure blocked the responses observed after 3.5 h but had no significant effect on responses observed at 1.5 h. A similar pattern was observed



**FIG. 1.** Inhibition of microtubules or the Golgi apparatus blocks sustained but not acute mobilization of  $K_{Ca}$  by NRG1. Embryonic day 9 (E9) ciliary ganglion (CG) neurons were incubated for 1.5 or 3.5 h in the presence or absence of 10 nM  $\beta$ -neuregulin-1 (NRG1), 20  $\mu$ M nocodazole, 5  $\mu$ M colchicine, or 1  $\mu$ g/ml brefeldin A, as indicated. Treatment with the inhibitors agents commenced 30 min prior to the onset of growth factor treatment.  $K_{Ca}$  and voltage-activated  $Ca^{2+}$  currents were quantified at the end of the treatment by whole cell recordings. **A:** examples of typical whole cell recordings in ciliary neurons treated with growth factors, brefeldin A (BFA), or overexpressing dominant-negative forms of signaling proteins described in subsequent figures. **B:** treatment with nocodazole or colchicines (which block microtubules) or brefeldin-A (which disrupts the Golgi apparatus) had no effect on  $K_{Ca}$  density measured 1.5 h after the start of NRG1 but blocked the effect of NRG1 measured 3 h after the start of treatment. **C:** in a different group of cells, brefeldin A produced the same quantitative pattern when currents were normalized for whole cell capacitance. **D:** voltage-activated  $Ca^{2+}$  currents measured at 3 h were not affected by these drugs.

in ciliary neurons pretreated with 1  $\mu$ g/ml brefeldin-A, an agent that disrupts the Golgi apparatus (Lippincott-Schwartz et al. 1989) (Fig. 1B). The same quantitative conclusion was obtained when currents were normalized for cell size by means of whole cell capacitance (Fig. 1C). Consistent with our previous observations (Lhuillier and Dryer 2002), none of these treatments affected the density of voltage-activated  $Ca^{2+}$  currents (Fig. 1D). These treatments, and the others used in subsequent experiments in this study, had no effect on kinetics of  $Ca^{2+}$  current activation or deactivation (data not shown). The ability of brefeldin A treatment to disrupt the Golgi apparatus in ciliary neurons, as it does in other cell types, was addressed directly by means of the fluorescent Golgi marker BODIPY TR C<sub>5</sub> ceramide (Fig. 2). The Golgi apparatus in ciliary neurons was completely disrupted by brefeldin A treatment. Therefore these data are consistent with a model in which NRG1 initially causes mobilization of  $K_{Ca}$  channels

located in post-Golgi pools but subsequently recruits channels from the Golgi apparatus.

To further test this model, we examined responses to NRG1 in E9 ciliary neurons over-expressing a form of ARF1 (T31N) that preferentially binds GDP and thereby functions as a dominant-negative (Dascher and Balch 1994). Cells were concurrently transfected with GFP to allow for identification of transfected cells during electrophysiology (Chae and Dryer 2005; Chae et al. 2005; Lhuillier and Dryer 2003). Control neurons were transfected with GFP alone. Transfections were carried out using biolistic methods, and growth factor treatments and whole cell recordings were carried out 24 h after transfection. Our earlier studies have shown that expression of constructs from CMV promoters in chick ciliary neurons is robust 6 h after transfection (Lhuillier and Dryer 2003). We observed that 1.5-h exposure to 10 nM NRG1 continued to cause mobilization of  $K_{Ca}$  in ciliary neurons co-expressing

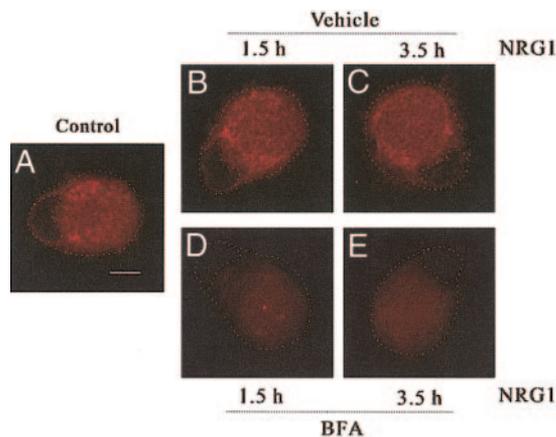


FIG. 2. Brefeldin A treatment causes complete disruption of the Golgi apparatus. Confocal images of Golgi were obtained using the fluorophore marker *N*-{[4-(4, 4-difluoro-5-(2-thienyl)-4-bora-3a,4a-diaza-*s*-indacene-3-yl)phenoxy]acetyl}sphingosine (BODIPY TR C<sub>5</sub> ceramide). ···, outline of cell surface and nucleus. Control shows Golgi apparatus in an untreated ciliary neuron. Other images are from different cells treated with 10 nM NRG1 for 1.5 h (left) or 3.5 h (right) in the absence (top) or presence (bottom) of 1 μg/ml brefeldin A.

GFP and dominant-negative ARF1, and the responses were indistinguishable from those obtained in ciliary neurons expressing GFP alone. By contrast, overexpression of dominant-negative ARF1 completely blocked responses to NRG1 monitored 3.5 h after the onset of growth factor treatment (Fig. 3A). Dominant-negative ARF1 and GFP overexpression had no effect on voltage-activated Ca<sup>2+</sup> currents monitored after 3.5 h of NRG1 (Fig. 3B). A similar conclusion was obtained using a different method to disrupt ARF1 function. ARF-GAP1 is the founding member of a class of GTPase-activating proteins known to terminate the actions of ARF1 (Cukierman et al. 1995). Overexpression of wild-type ARF-GAP1 causes disruption of the Golgi apparatus, most likely owing to increased GTP hydrolysis on ARF1 (Aoe et al. 1997). In the present experiments, we used an ARF1-GFP fusion protein (GFP-ARF-GAP273) that blocks protein translocation through the Golgi when overexpressed (Yu and Roth 2002). As with dominant-negative ARF1, overexpression of GFP-ARF-GAP273 for 24 h had no effect on the response of NRG1 monitored after a 1.5-h exposure but blocked responses observed after a 3.5-h exposure (Fig. 4A). Overexpression of this construct had no significant effect on the expression or properties of Ca<sup>2+</sup> currents (Fig. 4B). Collectively these data point to multiple pools of K<sub>Ca</sub> channels in different cellular compartments that can be mobilized by NRG1 on different time scales.

#### ARF6 signaling is required for both the acute and sustained stimulation of K<sub>Ca</sub>

ARF6 regulates a number of protein trafficking events that occur in and near the plasma membrane (Donaldson 2003). To determine whether ARF6 is involved in mobilization of K<sub>Ca</sub> channels, E9 ciliary ganglion neurons were transfected with a mutant form of ARF6 (T27N) defective in GTP binding that thereby functions as a dominant-negative (D'Souza-Schorey et al. 1998; Macia et al. 2004). As with ARF1 constructs, transfection was carried out 24 h before application of growth factors. We observed that overexpression of ARF6 (T27N)

blocked responses to 10 nM NRG1 monitored both 1.5 and 3.5 h after the onset of treatment (Fig. 5A). However, expression of this construct had no effect on voltage-activated Ca<sup>2+</sup> currents (Fig. 5B). The activity of ARF6 is regulated by a guanine nucleotide exchange factor known as ARF nucleotide binding site opener (ARNO), one of a conserved family of Sec7-domain containing proteins that regulate various ARF proteins (Jackson and Casanova 2000). The Sec7 domain is the catalytic center of these proteins, and a conserved glutamate residue within this domain is required for catalytic activity in all members of this family (Beraud-Dufour et al. 1999). Therefore we examined NRG1 mobilization of K<sub>Ca</sub> in E9 ciliary neurons over-expressing an ARNO mutant (E156K) that lacks this conserved residue and that functions as a dominant nega-

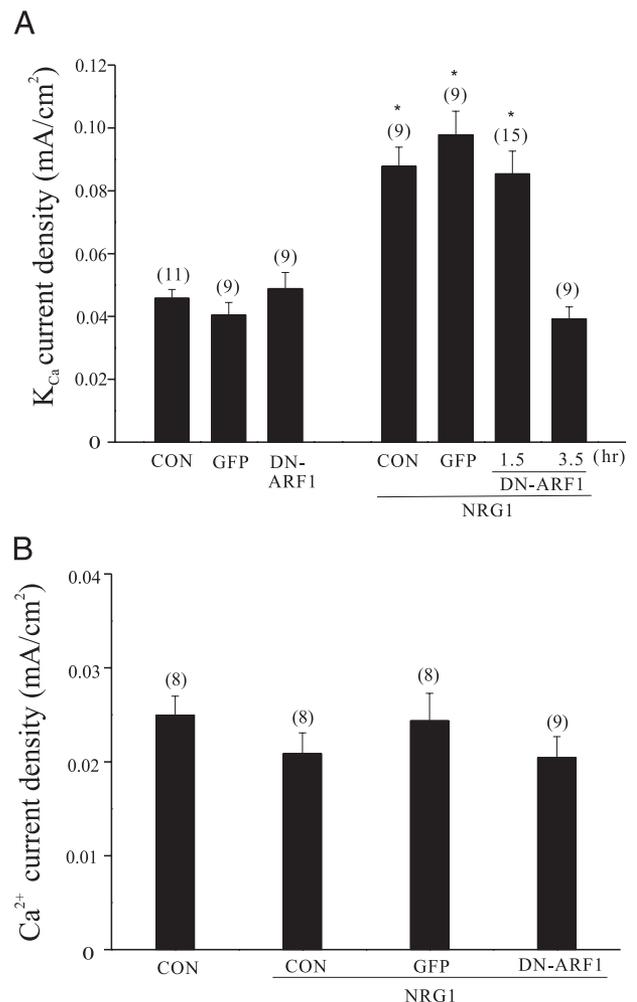


FIG. 3. ADP-ribosylation factor-1 (ARF1) is necessary for the sustained but not for the acute mobilization of K<sub>Ca</sub> evoked by NRG1. These experiments utilized biolistic overexpression of ARF1 (T31N), a dominant-negative form of ARF1. *A*: macroscopic K<sub>Ca</sub> density in ciliary neurons transfected with *Renilla* green fluorescent protein (GFP), a combination of GFP and ARF1 (T31N), or from nontransfected cells in the same dish (CON). Note robust stimulation by NRG1 in nontransfected cells, in cells expressing only GFP, and in cells incubated for 1.5 h with NRG1 after overexpressing of GFP and ARF1 (T31N). However, over-expression of ARF1 (T31N) with GFP completely blocked the responses to NRG1 measured 3.5 after the start of treatment. *B*: overexpression of ARF1 (T31N) or GFP had no effect on mean Ca<sup>2+</sup> current density in any treatment group. *C*: E9 ciliary neurons overexpressing a GFP-tagged form of ARF1 were incubated for 1.5 or 3.5 h in the presence or absence of 10 nM NRG1. NRG1 treatment promoted movement of ARF1 into the Golgi apparatus, especially after 3.5 h.

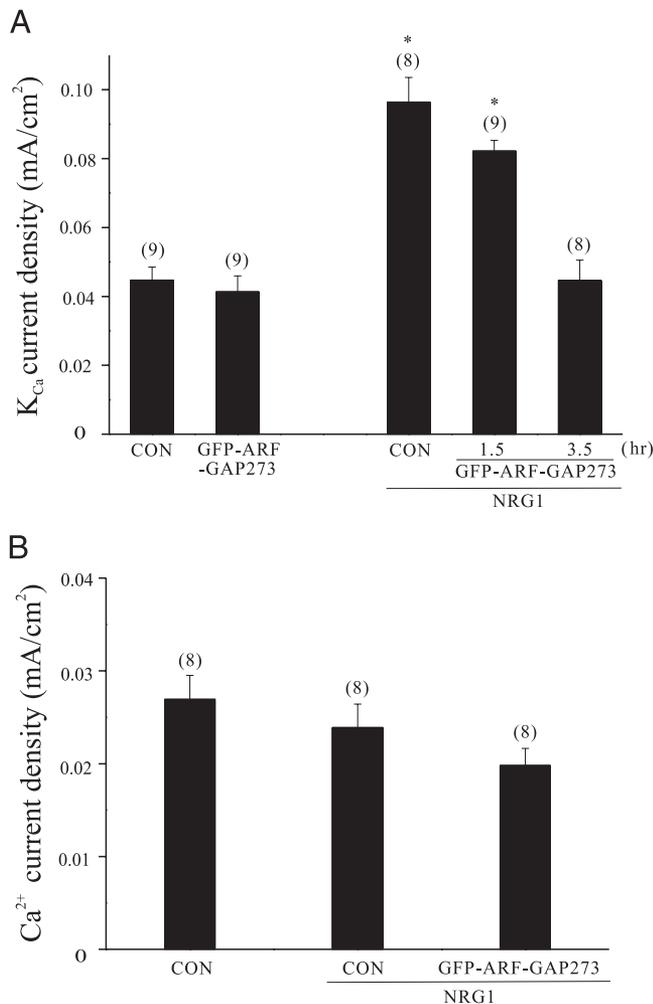


FIG. 4. ARF1 is necessary for the sustained but not for the acute mobilization of  $K_{Ca}$  evoked by NRG1. These experiments used overexpression of ARF-GAP273, an agent that stimulates the intrinsic GTPase activity of ARF1, and thereby inhibits its function. *A*: overexpression of ARF-GAP273 has no effect on response to NRG1 measured at 1.5 h but completely blocked responses to NRG1 measured at 3 h. These results are similar to those obtained with dominant-negative ARF1. *B*: overexpression of ARF-GAP273 did not affect voltage-activated  $Ca^{2+}$  currents.

tive (Mukherjee et al. 2001). We observed a pattern similar to that obtained with the ARF6 dominant negative; specifically, both the acute (1.5 h) and sustained (3.5 h) responses to 10 nM NRG1 were abolished, but  $Ca^{2+}$  currents were fully normal (Fig. 6).

ARF6 affects membrane and cytoskeleton dynamics by regulating multiple effector molecules (Donaldson 2003). One of these is PLD1, which is thought to be necessary to induce transient changes in membrane lipid composition required for vesicle trafficking to the plasma membrane of neurons and neuroendocrine cells (Caumont et al. 1998; Vitale et al. 2001). To explore the role of PLD1 in  $K_{Ca}$  trafficking in ciliary neurons, we overexpressed a catalytically dead form of PLD1 (K898R) that functions as dominant-negative (Denmat-Ouisse et al. 2001; Rizzo et al. 1999; Vitale et al. 2002). We observed that over-expression of PLD1 (K898R) in E9 ciliary neurons blocked the increase in macroscopic  $K_{Ca}$  evoked by a 1.5-h treatment with NRG1 (Fig. 7A) or a 3-h treatment with NRG1 (data not shown). Overexpression of this construct had no

effect on expression of voltage-activated  $Ca^{2+}$  channels in E9 neurons (Fig. 7B). We also examined the role of PLD1 using a different experimental design that entailed use of a constitutively active form of ARF6 (Q67L). This mutant cannot hydrolyze bound GTP and is therefore unable to cycle between GTP- and GDP-bound states. As with other constitutively-active small GTPases, this mutant is able to recapitulate some but not all of the actions of active wild-type ARF6 (Santy 2002). We observed that overexpression of ARF6 (Q67L) had no effect on basal  $K_{Ca}$  expression in E9 ciliary neurons (Fig. 7C). Similarly, overexpression of wild-type PLD1 had no effect on basal  $K_{Ca}$  density. However, overexpression of ARF6 (Q67L) together with wild-type PLD1 caused a modest but statistically significant increase in  $K_{Ca}$  (Fig. 7C). As an aside, it bears noting that over-expression of ARF6 (Q67L) not only

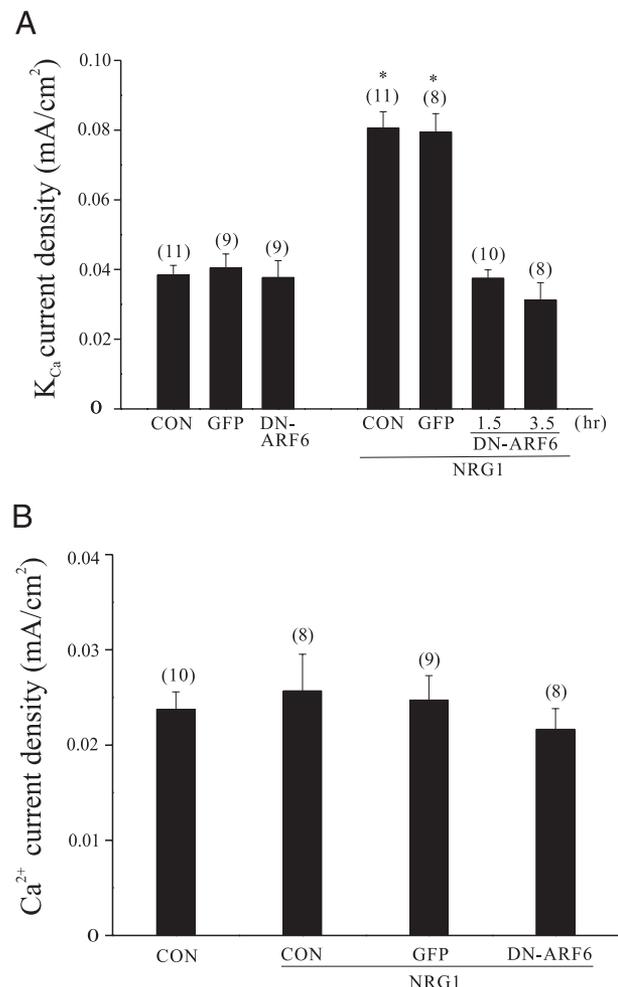


FIG. 5. ARF6 is necessary for both the acute and sustained mobilization of  $K_{Ca}$  evoked by NRG1. Macroscopic  $K_{Ca}$  and  $Ca^{2+}$  currents were recorded from ciliary neurons overexpressing GFP or a combination of GFP and ARF6 (T27N), a dominant-negative form of ARF6. Recordings were also made from nontransfected cells in the same culture dish (CON). *A*: overexpression of ARF6 (T27N) completely blocked the effect of NRG1 on functional expression of  $K_{Ca}$  measured at all times after treatment. *B*: voltage-activated  $Ca^{2+}$  currents were not affected by overexpression of ARF6 (T27N). *C*: E9 ciliary neurons overexpressing a GFP-tagged form of ARF6 were incubated for 1.5 or 3.5 h in the presence (NRG1) or absence (control) of 10 nM NRG1. NRG1 treatment evoked some movement of tagged ARF6 toward the periphery of the cell at 1.5 h, with some movement back into intracellular compartments apparent at 3 h. The latter phenomenon may reflect accumulation of GDP-bound ARF6 that has completed its cycle.

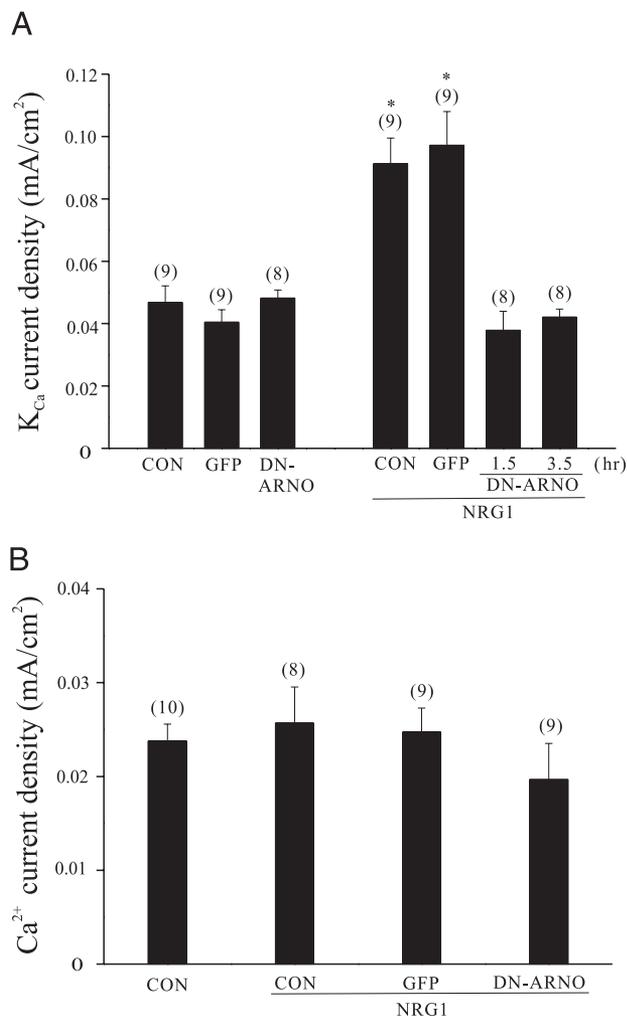


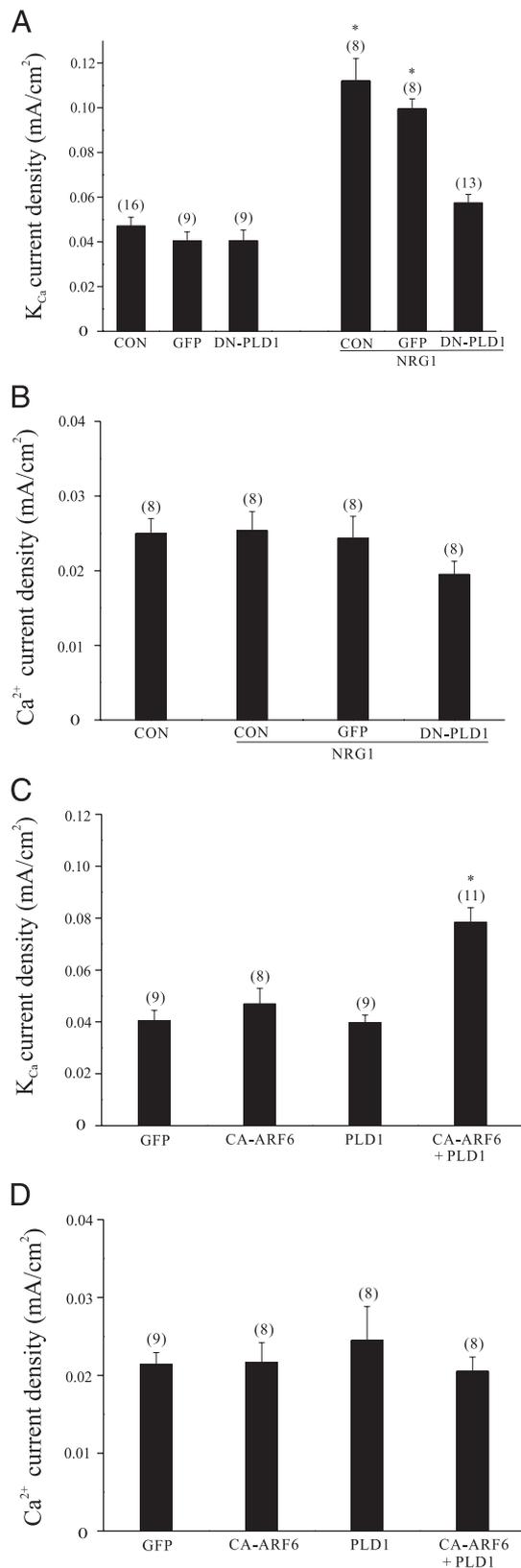
FIG. 6. ARF nucleotide-binding site opener (ARNO) is required for both the acute and sustained mobilization of  $K_{Ca}$  channels evoked by NRG1. These experiments employed overexpression of the dominant-negative mutant ARNO (E156K). *A*: overexpression of ARNO (E156K) completely blocked mobilization of  $K_{Ca}$  measured 1.5 and 3.5 h after the onset of NRG1 treatment. *B*: voltage-activated  $Ca^{2+}$  currents were not affected by overexpression of ARNO (E156K).

failed to increase trafficking of  $K_{Ca}$  channels by itself, it also blocked the stimulation of  $K_{Ca}$  normally observed following 1.5- or 3.5-h treatments with NRG1 (data not shown). In other words, with respect to mobilization of  $K_{Ca}$ , the GTPase-deficient ARF6 mutant also behaved as a dominant-negative, as reported in other cell types (Zhang et al. 1998). Collectively, these data indicate that an ARF6-PLD1 mechanism is necessary for  $K_{Ca}$  trafficking to the plasma membrane.

Finally, it bears noting that ARF1, ARF6, ARNO, and PLD1 are also necessary for sustained posttranslational responses to

FIG. 7. PLD1 is required for both the acute and sustained mobilization of  $K_{Ca}$  evoked by NRG1. *A*: overexpression of dominant-negative phospholipase D1 (PLD1) (K898R) completely blocked the effect of NRG1 on functional expression of  $K_{Ca}$  measured 1.5 h after the onset of growth factor treatment. *B*: voltage-activated  $Ca^{2+}$  currents were not affected by overexpression of PLD1 (K898R). *C*: overexpression of the constitutively-active mutant ARF6 (Q67L) together with wild-type PLD1 caused a modest but statistically significant increase in  $K_{Ca}$ , even in the absence of growth factor treatment. Overexpression of these constructs by themselves was not sufficient to cause mobilization of  $K_{Ca}$ . *D*: voltage-activated  $Ca^{2+}$  currents were not affected by overexpression of constitutively active ARF6 (Q67L) and/or PLD1.

TGF $\beta$ 1, another growth factor required for normal  $K_{Ca}$  expression in developing ciliary neurons (Fig. 8). In those experiments, we examined  $K_{Ca}$  expression in transfected E9 ciliary neurons after a 6.5-h exposure to TGF $\beta$ 1 using the constructs already described. The results suggest that two growth factors



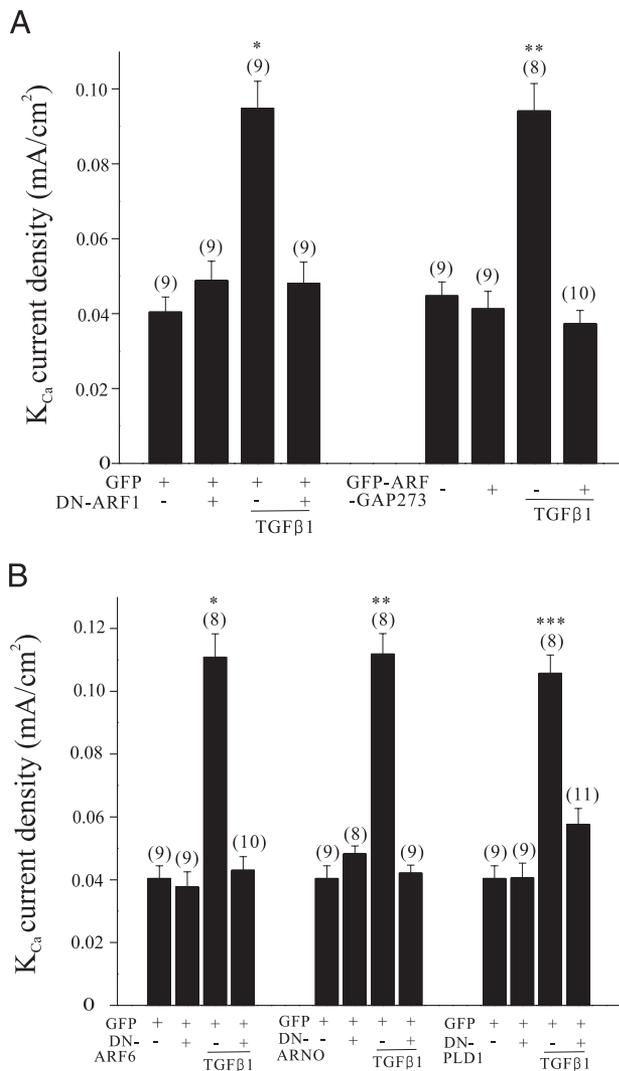


FIG. 8. ARF1, ARF6, ARNO, and PLD1 are required for sustained mobilization of  $K_{Ca}$  evoked by TGFβ1. These experiments used over-expression of the constructs described in previous figures. Mobilization of  $K_{Ca}$  was quantified in E9 ciliary neurons by whole cell recording after a 6.5-h exposure to 1 nM TGFβ1. Shorter treatments (e.g., 1.5–3 h) do not cause statistically significant stimulation of  $K_{Ca}$ . *A*: overexpression of ARF1 (T31N) or ARF-GAP273 completely blocked the effect of TGFβ1 on functional expression of  $K_{Ca}$ . *B*: overexpression of ARF6 (T27N), ARNO (E156K) or PLD1 (K898R) completely blocked the effect of TGFβ1 on functional expression of  $K_{Ca}$  measured at 6.5 h. Voltage-activated  $Ca^{2+}$  currents were not affected by overexpression of these mutants as noted in previous figures.

share at least some of the pathways required for mobilization of  $K_{Ca}$  channels in the endoplasmic reticulum and/or Golgi apparatus of ciliary neurons. Responses to TGFβ1 with shorter treatment times are weak and highly variable and are not detectable at all at 1.5 h. Thus TGFβ1 appears to be much less effective than NRG1 at mobilizing the pool of  $K_{Ca}$  channels associated with post-Golgi compartments.

## DISCUSSION

The data in this study are consistent with a model in which functional BK-type  $K_{Ca}$  channels of developing ciliary neurons are mobilized into the plasma membrane from two separate compartments in response to physiologically relevant growth factors. One pool of channels in the ER and/or Golgi is

mobilized slowly (>3 h) in response to TGFβ1 or high concentrations of NRG1 (Chae et al. 2005; Lhuillier and Dryer 2002). A second  $K_{Ca}$  pool in a post-Golgi compartment moves into the plasma membrane more rapidly after the onset of NRG1 exposure (0.5–1.5 h) but cannot be effectively mobilized by TGFβ1. We have previously shown that stimulation of  $K_{Ca}$  evoked by lower concentrations (1 nM) of NRG1 is transient; responses are significant 1–1.5 after the onset of treatment but return to baseline after 3 h, even in the continuous presence of NRG1 (Chae et al. 2005). By contrast, treatment with 10 nM NRG1, the concentration used in the present study, evokes responses that last essentially as long as NRG1 is present. This suggests that the post-Golgi pool appears to be preferentially mobilized by low concentrations of NRG1 (Chae et al. 2005). For reasons as yet unknown, TGFβ1 is unable to rapidly mobilize the  $K_{Ca}$  channels in the post-Golgi pool.

The two-pool model of  $K_{Ca}$  regulation is supported by several lines of evidence. Thus the sustained physiological responses to NRG1 and TGFβ1 are completely inhibited by treatments that cause disruption of Golgi function, including microtubule inhibitors (Hamm-Alvarez and Sheetz 1998), overexpression of dominant-negative forms of ARF1 (Dascher and Balch 1994), brefeldin-A (Lippincott-Schwartz et al. 1989), and expression of a mutant form of ARF-GAP-1 that blocks ARF1 cycling (Yu and Roth 2002). None of these treatments affected the acute responses to NRG1, defined in this study as increases in macroscopic  $K_{Ca}$  observed ≤1.5 h after the onset of treatment. Further, none of these treatments affected the expression or properties of voltage-activated  $Ca^{2+}$  currents, consistent with our previous studies (Lhuillier and Dryer 2002).

Both the acute and sustained responses to growth factors ultimately entail insertion of  $K_{Ca}$  channels into the plasma membrane. Thus NRG1 and TGFβ1 increase the number of immunohistochemically detectable cell-surface SLO1 proteins (Chae et al. 2005). Moreover, we observed here that inhibition of ARF6 signaling cascades completely blocked the increases in macroscopic  $K_{Ca}$  evoked by NRG1 or TGFβ1. This was observed with both short (1.5 h) and long (>3.5 h) growth factor treatments that are differentially sensitive to inhibition of ARF1 cascades, and equivalent blockade was produced by overexpression of dominant-negative forms of ARF6 (D'Souza-Schorey et al. 1998), ARNO (Mukherjee et al. 2001), and PLD1 (Rizzo et al. 1999). Clearly, ARF6 activation is necessary for growth factor mobilization of  $K_{Ca}$ . However, it does not appear to be sufficient, as overexpression of a constitutively active form of ARF6 (Dascher and Balch 1994) did not lead to an increase in macroscopic  $K_{Ca}$ , except for a modest effect in cells that were concurrently overexpressing wild-type PLD1. Indeed, the constitutively-active form of ARF6, which is limited in its ability to cycle between GTP- and GDP-bound states, not only failed to cause mobilization of  $K_{Ca}$ , it actually inhibited the physiological responses to NRG1. A similar pattern has been observed with constitutively-active forms of ARF6 in other systems (Brown et al. 2001; Santy 2002) and with some other small GTPases (e.g., Lin et al. 1997).

It is likely that ARF6 acts on multiple effectors to regulate insertion of functional  $K_{Ca}$  complexes into the plasma membrane. In other systems, ARF6 causes activation of PLD1, leading to focal changes in the composition of the plasma

membrane (Donaldson 2003). Consistent with this, our current data indicate that an ARF6-PLD1 pathway is an essential component of the cascades leading to  $K_{Ca}$  mobilization in ciliary neurons. This is true when mobilization is evoked by either NRG1 or TGF $\beta$ 1. However, ARF6 activation also causes changes in the dynamics of a dense layer of filamentous actin adjacent to the plasma membrane of most cell types (Schafer et al. 2000). This so-called cortical actin layer can play a variety of roles in the late stages of membrane protein trafficking (Eitzen 2003). In a recent study, we observed that cortical actin acts as a barrier to prevent the constitutive insertion of  $K_{Ca}$  channels into the plasma membrane of ciliary neurons. Specifically, we observed that agents that cause depolymerization of cortical F-actin, such as cytochalasin D, rapidly stimulate insertion of  $K_{Ca}$  into the plasma membrane. Conversely, treatments that stabilize F-actin, such as phalloidin, tend to suppress ciliary neuron  $K_{Ca}$  mobilization (Chae and Dryer 2005). These data raise the possibility that actin rearrangements mediate some of the effects of ARF6 on  $K_{Ca}$  mobilization.

The mechanisms whereby NRG1 and TGF $\beta$ 1 lead to activation of ARFs in ciliary neurons are not known. However, we have previously shown that PI3 kinase and Akt activation are required for NRG1- or TGF $\beta$ 1-evoked mobilization of  $K_{Ca}$  channels (Chae et al. 2005; Lhuillier and Dryer 2000). Indeed, over-expression of a membrane-targeted (myristoylated) form of Akt is sufficient in itself to cause mobilization of  $K_{Ca}$  (Chae et al. 2005). Several ARF-associated GEFs of the Sec7 family, such as ARNO and GRP1, contain tandem pleckstrin homology (PH) domains that allow for binding of the lipid products of PI3 kinase, and a consequent association with cell membranes (Klarlund et al. 1998; Venkateswarlu et al. 1998). Akt also contains a PH domain that is necessary for its activation, and it is therefore possible that ARF-related proteins and Akt are brought into close proximity on endomembranes and plasma membranes of ciliary neurons in response to NRG1 or TGF $\beta$ 1. To date there have been no published reports of direct interactions between ARFs or ARF-associated proteins and Akt. However, a number of Akt substrates, such as p70 S6 kinase-1 (Qian et al. 2004) and p21-activated kinase (Bokoch 2003) are able to modulate cortical F-actin dynamics. Thus there are several potentially interacting mechanisms whereby Akt and ARF6 cascades could converge to coordinately regulate trafficking of  $K_{Ca}$  in ciliary neurons. Indeed, we cannot exclude that overexpression of a membrane-targeted form of Akt can induce dynamic changes in trafficking that, under more physiological circumstances, are mediated by ARFs.

It is surprising that two growth factors regulate different pools of  $K_{Ca}$  channels in ciliary neurons, and the question arises as to the functional significance of the rapid regulation of macroscopic  $K_{Ca}$  evoked by NRG1. TGF $\beta$ 1 and NRG1 are both required for developmental regulation of  $K_{Ca}$  in ciliary neurons developing in vivo (Cameron et al. 1998, 2001). It is possible that NRG1 regulation persists past development and plays a role in fully formed ciliary ganglion, as occurs with growth factor regulation of potassium channels in other systems (Fadool et al. 2000). By analogy, previous work suggests that NRG1 plays a role in the maintenance as well as the formation of nicotinic acetylcholine receptors in the developing mammalian neuromuscular junction (Sandrock et al. 1997).

In summary, we have shown that developing ciliary neurons maintain at least two distinct pools of  $K_{Ca}$  channels. These include a rapidly accessible pool located in a post-Golgi compartment and a major reserve pool located in the ER-Golgi. ARF1 and associated proteins are required for movement of  $K_{Ca}$  channels through the Golgi apparatus, whereas an ARF6 cascade that also requires PLD1 regulates the late stages of translocation of functional  $K_{Ca}$  channels into the plasma membrane. These data also provide plausible mechanisms for growth-factor-evoked regulation of the neuronal cytoskeleton in the context of protein trafficking and the developmental regulation of excitability.

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