Development/Plasticity/Repair

# PLD1 Negatively Regulates Dendritic Branching

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Neurons have characteristic dendritic arborization patterns that contribute to information processing. One essential component of dendritic arborization is the formation of a specific number of branches. Although intracellular pathways promoting dendritic growth and branching are being elucidated, the mechanisms that negatively regulate the branching of dendrites remain enigmatic. In this study, using gain-of-function and loss-of-function studies, we show that phospholipase D1 (PLD1) acts as a negative regulator of dendritic branching in cultured hippocampal neurons from embryonic day 18 rat embryos. Overexpression of wild-type PLD1 (WT-PLD1) decreases the complexity of dendrites, whereas knockdown or inhibition of PLD1 increases dendritic branching. We further demonstrated that PLD1 acts downstream of RhoA, one of the small Rho GTPases, to suppress dendritic branching. The restriction of dendritic branching by constitutively active RhoA (V14-RhoA) can be partially rescued by knockdown of PLD1. Moreover, the inhibition of dendritic branching by V14-RhoA and WT-PLD1 can be partially ameliorated by reducing the level of phosphatidic acid (PA), which is the enzymatic product of PLD1. Together, these results suggest that RhoA-PLD1-PA may represent a novel signaling pathway in the restriction of dendritic branching and may thus provide insight into the mechanisms of dendritic morphogenesis.

### Introduction

Dendritic morphogenesis is an important step for the establishment of neural circuitry. The numbers of primary dendrites arising from the cell body, higher order dendrites emerging from primary dendrites, and dendritic branching patterns appear to be critical for neuronal function (Jan and Jan, 2010). Although intracellular pathways promoting dendritic growth and branching are being elucidated, the mechanisms that restrict the overgrowth of dendrites remain enigmatic.

The small GTPases of the Rho subfamily are critical regulators of dendritic morphogenesis (Luo, 2000). The best studied Rho GTPases are RhoA, Rac1, and Cdc42. Among these Rho GTPases, RhoA is unique in its ability to inhibit dendritic branching both in cultured neurons and mosaic *Drosophila* brains (Lee et al., 2000; Li et al., 2000). Exactly how RhoA negatively regulates dendritic branching is not comprehensively defined. Several kinases downstream of RhoA have been identified including Rho kinase

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DOI:10.1523/JNEUROSCI.5378-11.2012 Copyright © 2012 the authors 0270-6474/12/327960-10\$15.00/0 system represents one of the major mechanisms of cellular signaling. Phospholipase D (PLD), which catalyzes the hydrolysis of phosphatidylcholine (PC) to generate phosphatidic acid (PA) and choline, has emerged as a signal transduction phospholipase in recent studies (Cockcroft, 1996; Exton, 1999; Rudge and Wakelam, 2009). PLD isozymes, including PLD1 and PLD2, are present in many types of mammalian cells, including neurons (Meier et al., 1999). Recently, both PLD1 and PLD2 have been included in the production of positive parts in present in the production.

(Matsui at el., 1996) and LIM kinase 1 (Maekawa et al., 1999; Ohashi et al., 2000). However, the roles of several proteins down-

stream of RhoA in dendritic branching have not been confirmed

by genetic studies. For example, dendritic branching is normal in

LIM kinase 1 knock-out neurons (Meng et al., 2002). Although

compensatory mechanisms may exist, this finding led us to ex-

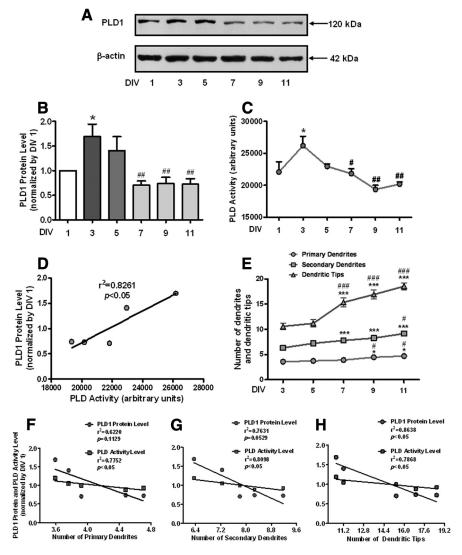
amine the role of other signaling molecules downstream of RhoA

In addition to kinases, the lipid-mediated second messenger

in the regulation of dendritic branching.

implicated in the regulation of neurite growth in neural cell lines (Hayakawa et al., 1999; Gibbs and Meier, 2000) and neural stem cells (Sung et al., 2001; Yoon et al., 2005) and the seizure-induced sprouting of mossy fibers (Zhang et al., 2004, 2005).

In the present study, we focused on PLD1 because it can be directly activated by RhoA both *in vitro* and *in vivo* (Hammond et al., 1997; Bae et al., 1998; Yamazaki et al., 1999; Du et al., 2000; Komati et al., 2005; Yoon et al., 2005; Gayral et al., 2006). Using cultured hippocampal neurons as a cellular model, we found that PLD1 negatively regulated dendritic branching. The inhibition of dendritic growth by PLD1 is consistent with the reduction of its expression and activity during dendritic maturation. Further experiments indicated that PLD1 partially contributed to the role of RhoA in dendritic arborization. PA, the enzymatic product of PLD1, appears to be important in mediating the effects of RhoA and PLD1



**Figure 1.** Protein level and activity of PLD1 during the development of hippocampal neurons. *A*, Western blot analysis of PLD1 in developing hippocampal neurons at indicated time points. *B*, Quantification of PLD1 protein levels during the development of hippocampal neurons. n=7 samples each time point. \*p<0.05, between DIV 3 and DIV 1. \*p<0.01, between DIV 7 and DIV 3, DIV11 and DIV3, and one-way ANOVA followed by Newman–Keuls multiple-comparison test. *C*, Quantification of PLD activity in developing hippocampal neurons. n=3 samples each time point. \*p<0.05, between DIV 3 and DIV 1. \*p<0.05, between DIV 7 and DIV 3. \*p<0.05, between DIV 9 and DIV3, DIV 11 and DIV 3, and one-way ANOVA followed by Newman–Keuls multiple-comparison test. *D*, The correlation between PLD1 protein levels and PLD activities ( $r^2=0.8261$ , p<0.05). *E*, Quantification of primary dendrites, secondary dendrites, and dendritic tips from DIV 3 to DIV 11. DIV 3, n=22; DIV 5, n=25; DIV 7, n=24; DIV 9, n=25; DIV 11, n=18. \*p<0.05, between DIV 9, 11 and DIV 3. \*p<0.05, between DIV 9, 11 and DIV 5 in secondary dendrites. \*\*\*\*p<0.001, between DIV 7, 9, 11 and DIV 3. \*p<0.05, between DIV 11 and DIV 5 in secondary dendrites. \*\*\*\*p<0.001, between DIV 7, 9, 11 and DIV 3. \*p<0.05, between DIV 7, 9, 11 and DIV 5 in dendritic tips, one-way ANOVA followed by Newman–Keuls multiple-comparison test. \*p<0.05, between DIV 7, 9, 11 and DIV 5 in dendritic tips, one-way ANOVA followed by Newman–Keuls multiple-comparison test. \*p<0.05, DIV 11. \*p<0.05, Detween DIV 7, 9, 11 and DIV 5 in dendritic tips one-way ANOVA followed by Newman–Keuls multiple-comparison test. \*p<0.05, Dividence DIV 7, 9, 11 and DIV 5 in dendritic tips one-way ANOVA followed by Newman–Keuls multiple-comparison test. \*p<0.05, Dividence DIV 7, 9, 11 and DIV 5 in dendritic tips, one-way ANOVA followed by Newman–Keuls multiple-comparison test. \*p<0.05, Dividence DIV 7, 9, 11 and DIV 5 in dendritic tips, one-way ANOVA followed by N

in dendritic branching. Overall, these results indicate that RhoA, PLD1, and PA may constitute one signaling pathway in the regulation of dendritic branching.

### **Materials and Methods**

Antibodies and chemicals. We used the following antibodies and chemicals: rabbit polyclonal anti-PLD1 (#3832; Cell Signaling Technology), mouse monoclonal anti- $\beta$ -actin (Sigma-Aldrich), mouse monoclonal anti-RhoA (sc-418; Santa Cruz Biotechnology), rabbit polyclonal anti-HA (sc-805; Santa Cruz Biotechnology), mouse monoclonal anti-Myc

(Zhongshan Golden Bridge Biotechnology), and 1-butanol and 2-butanol were purchased from Sigma-Aldrich.

DNA constructs. Hemagglutinin (HA)tagged wild-type PLD1 (WT-PLD1) and HAtagged dominant-negative PLD1 (DN-PLD1) were generously provided by Dr. Michael A. Frohman (Stony Brook University, Stony Brook, NY). To create siRNA-resistant PLD1 (RES-PLD1), the third nucleotide of each codon in the target sequence was mutated without changing the identity of the amino acids. cDNA of the constitutively active RhoA mutant (V14-RhoA) and the dominantnegative mutant of RhoA (N19-RhoA) were generously provided by Dr. Huaye Zhang (Department of Neuroscience and Cell Biology, Robert Wood Johnson Medical School, University of Medicine and Dentistry of New Jersey). The cDNA of V14-RhoA and N19-RhoA were subcloned to pcDNA3.1/Myc-His C (Clontech) to generate the Myc-His-RhoA fusion constructs, which were verified by DNA sequencing.

Neuronal culture and transfection. Hippocampal explants isolated from embryonic day 18 rat embryos of either sex were digested with 0.25% trypsin for 30 min at 37°C followed by trituration with a pipette in plating medium (DMEM with 10% fetal bovine serum[FBS]). Dissociated neurons were plated onto 35 mm dishes coated with poly-D-lysine (Sigma-Aldrich) at a density of  $5 \times 10^5$  cells per dish. After culturing for 4 h, media were changed to Neurobasal medium supplemented with 2% B27 and 0.5 mm GlutaMAX-I (Invitrogen). To test the role of PLD1 in dendritic branching, neurons were cotransfected with the DNA of interest and pEGFP-N1 at 3 d or 6 d in vitro (DIV) were fixed with 4% paraformaldehyde and subjected to analysis of dendritic branching 3 d after transfection. Lipofectamine 2000 (Invitrogen) was used for neuronal transfection following the instructions provided by the manufacturer. To address the effects of pharmacological inhibitors on dendritic numbers, DIV 6 hippocampal neurons were treated with 1-butanol or 2-butanol in the culture medium for 3 d before the analysis of dendritic branching.

RNA interference. The sequence of PLD1 small interfering RNA (siRNA) was as follows: 5'-CUGGAAGAUUACUUGACAA-3'. siRNA with the sequence of 5'-UUCUCCGAA CGUGUCACGU-3' was used as a control because it is unable to knock down the expression of any known proteins. In the RNA interference experiments, PLD1 siRNA or control siRNA was transfected together with pEGFP-N1. For the rescue experiments, PLD1 siRNA,

RES-PLD1, and pEGFP-N1 were transfected together with Lipo-fectamine 2000.

Western blot analysis. Cell cultures were washed twice with ice-cold PBS containing (in mm): 137 NaCl, 2.7 KCl, 10 Na<sub>2</sub>HPO<sub>4</sub>, and 2 KH<sub>2</sub>PO<sub>4</sub> and lysed in ice-cold lysis buffer (50 mm Tris-HCl, pH 7.4, 150 mm NaCl, 1.5 mm MgCl<sub>2</sub>, 10% glycerol, 1% Triton X-100, 5 mm EGTA, 1  $\mu$ g/ml leupeptin, 1 mm PMSF, 1 mm Na<sub>3</sub>VO<sub>4</sub>, 10 mm NaF, and proteinase inhibitor mixture). The lysates were centrifuged at 12,000 × g for 5 min to yield the total protein extract in the supernatant. The concentration of protein was measured with a BCA assay kit (Pierce). Equal amounts of samples (50  $\mu$ g) were denatured

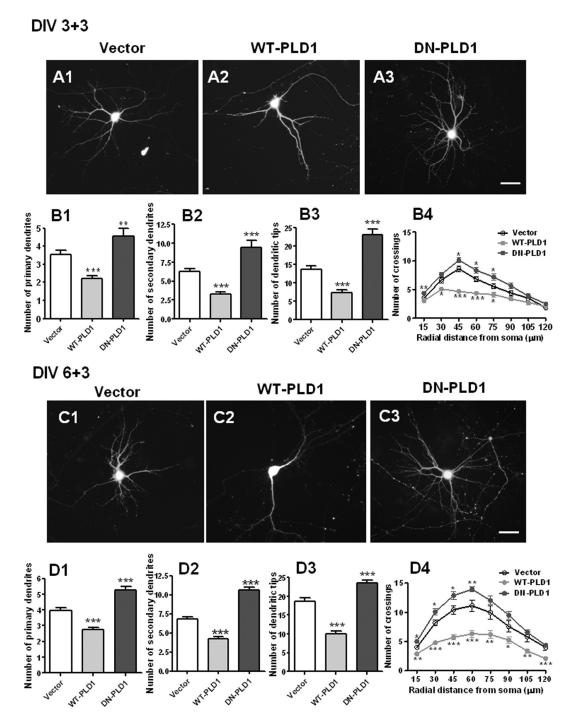
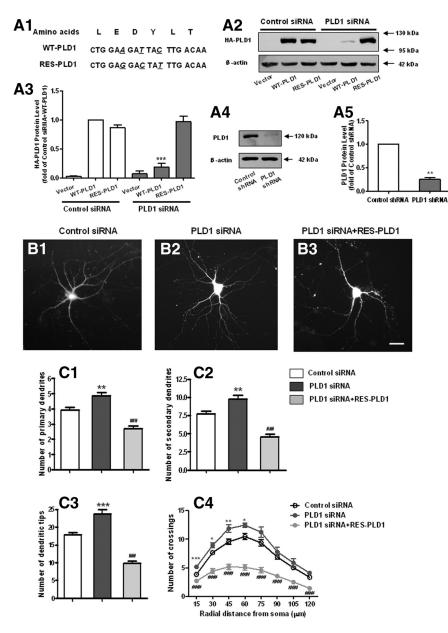


Figure 2. Effects of WT-PLD1 and DN-PLD1 on the dendritic branching of hippocampal neurons. A1-A3, Representative images of DIV 6 neurons transfected with HA-Vector (A1), HA-tagged WT-PLD1 (A2), or HA-tagged DN-PLD1 (A3) at DIV 3. Scale bar, 50  $\mu$ m. B1-B4, Quantification of primary dendrites (B1), secondary dendrites (B2), dendritic tips (B3), and Sholl analysis (B4) in A. n=22 neurons for control, n=45 neurons for WT-PLD1, n=14 neurons for DN-PLD1, n=14 neurons for DN-PLD1, n=14 neurons for DN-PLD1 (C3) at DIV 6. Scale bar, 50  $\mu$ m. D1-D4, Quantification of primary dendrites (D1), secondary dendrites (D2), dendritic tips (D3), and Sholl analysis (D4) in C. n=37 neurons for control, n=42 neurons for WT-PLD1, n=32 neurons for DN-PLD1, n=32 neurons for

and subjected to 10% SDS-PAGE. After separation, proteins were transferred to nitrocellulose membranes (Bio-Rad). The membranes were blocked with 5% nonfat milk in TBST (25 mm Tris-HCl, pH 7.4, 137 mm NaCl, 2.7 mm KCl, and 0.05% Tween 20) for 1 h at room temperature and incubated with primary antibody overnight at 4°C. After washing with TBST three times, the membranes were incubated with horseradish peroxidase (HRP)-conjugated secondary antibody for 1 h at room temperature, washed again, and finally were developed with ECL solutions (ThermoFisher Scientific). The immunoreactive bands were scanned and analyzed quantitatively by densitometry with Quantity One (Bio-Rad).

Cell culture and coimmunoprecipitation. Neuroblastoma N2a cells were cultured at 37°C in 60 mm dishes in a humidified atmosphere of 95% air and 5%  $\rm CO_2$ . The culture medium was DMEM with 10% FBS. Myc-His-V14-RhoA was coexpressed with HA-Vector, HA-WT-PLD1, or HA-DN-PLD1 in N2a cells. Thirty-six hours after transfection, cells were harvested in ice-cold lysis buffer, and the homogenates were centrifuged at  $12,000 \times g$  for 5 min to yield the total protein extract in the supernatant. Cellular extracts were mixed with anti-Myc antibody (1:100) at 4°C for 3 h before incubation with Protein A-Sepharose CL-4B resin (GE Healthcare) overnight. The



**Figure 3.** Effects of PLD1 siRNA on the dendritic branching of hippocampal neurons. **A1**, Comparison of the target sequence between WT-PLD1 and RES-PLD1. The third nucleotide in each codon was mutated in RES-PLD1 without altering the identity of amino acids. **A2**, **A3**, Knockdown effect of WT-PLD1 but not RES-PLD1 (siRNA-resistant PLD1) by PLD1 siRNA. n=5, \*\*\*p<0.001, compared with WT-PLD1 plus Control siRNA, one-way ANOVA followed by Newman–Keuls multiple-comparison test. **A4**, **A5**, Knockdown effect of endogenous PLD1 in hippocampal neurons by PLD1 shRNA. n=3, \*\*p<0.01, compared with Control shRNA, t test. **B1–B3**, Representative images of DIV 9 neurons transfected with Control siRNA (**B1**), PLD1 siRNA (**B2**), or PLD1 siRNA + RES-PLD1 (**B3**). Scale bar, 50  $\mu$ m. **C1–C4**, Quantification of primary dendrites (**C1**), secondary dendrites (**C2**), dendritic tips (**C3**), and Sholl analysis (**C4**) in **B**. n=37 for Control siRNA, n=19 for PLD1 siRNA, n=47 for PLD1 siRNA + RES-PLD1, \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, compared with Control siRNA, t test; \*##p<0.001, compared with PLD1 siRNA, t test.

beads were washed six times with TBS containing 0.1% Triton X-100 before boiling with 2  $\times$  SDS buffer for 10 min. After centrifugation at 3000  $\times$  g for 2 min, the supernatant was decanted and subjected to Western blot analysis.

*PLD activity assay.* Following the nonradioactive method, PLD-mediated PA production was analyzed with a fluorescent *in vitro* assay (Elvers et al., 2010). In this enzymatically coupled assay, PLD hydrolyzes PC in the presence of PI(4,5)P<sub>2</sub> to PA and choline, which is subsequently oxidized by choline oxidase to betaine and  $\rm H_2O_2$ . In the presence of HRP,  $\rm H_2O_2$  oxidizes Amplex red in a 1:1 stoichiometry to generate fluorescent resorufin (7-hydroxy-3H-phenoxazin-3-one). Because resorufin has absorption and fluorescence emission maxima of  $\sim$ 571 and 585 nm,

respectively, which is beyond the range of the autofluorescence from most biological samples, this fluorescent assay specifically detects PLD activity. In brief, equal amounts of cell lysates (100  $\mu$ g) were mixed with 100  $\mu$ l Amplex red reaction buffer from the Amplex Red PLD assay kit (Invitrogen). After incubation at 37°C in darkness for 45 min, the fluorescence was measured in a fluorescence microplate reader using excitation in the range of 530–560 nm and emission detection at  $\sim$ 590 nm. Each sample was duplicated to create average fluorescence values.

Measurement of PA. Neurons were harvested using a rubber policeman. After protein determination, 1.5 ml of methanol, 2.25 ml of 1 M NaCl, and 2.5 ml of chloroform were used to separate cellular lipids. The sample was centrifuged at 1500 g, and the lower phase was dried under a gentle stream of nitrogen. Lipids were resuspended with 1% Triton X-100. Following the nonradioactive method, PA concentration was analyzed with a fluorescent in vitro assay. In this enzymatically coupled assay, lipase is used to hydrolyze PA to glycerol-3phosphate, which is subsequently oxidized by glycerol-3-phosphate oxidase to generate H<sub>2</sub>O<sub>2</sub>. In the presence of peroxidase, H<sub>2</sub>O<sub>2</sub> reacts with 10-acetyl-3,7-dihydroxyphenoxazine to yield the highly fluorescent compound resorufin (Morita et al., 2009). Resorufin fluorescence can be analyzed using excitation wavelengths of 530-540 nm and emission wavelengths of 585-595 nm. The assay was performed following the instructions provided by the manufacturer (#700240; Cayman Chemical Company). Each sample was duplicated to create average fluorescence values.

Analysis of neuronal morphology. The morphologies of entire neurons were indicated by the expression of pEGFP-N1. Dissociated neurons grown at low density were used to determine morphological characteristics. Neurons treated with inhibitors were randomly chosen, and neurons transfected with DNA were selected by green fluorescent protein (GFP) expression. The neurons were photographed at 20× magnification using an Olympus fluorescent microscope. All processes and their branches were traced, and their numbers were counted using DP2-BSW microscope digital camera software (Olympus). The neurites that expressed Ankyrin G in the initial segment were taken to be the axons. To analyze dendritic branching, all nonaxonal protrusions longer than 10  $\mu$ m originating from the cell soma were defined as primary dendrites. All protrusions originating from the primary dendrites were defined as secondary

dendrites. All terminal branches of dendrites with lengths >10  $\mu$ m were counted as dendritic tips (Jaworski et al., 2005). For Sholl analysis, concentric circles with 15  $\mu$ m differences in diameter were drawn around the cell body, and the number of dendrites crossing each circle was manually counted (Sholl, 1953; Jaworski et al., 2005).

Statistical analysis. All data are represented as the mean  $\pm$  the SEM. Comparisons between two groups were made using t tests. Comparisons among three or more groups were made using one-way ANOVA analyses followed by Newman–Keuls tests. Data marked with asterisks are significantly different from the control as follows: \*\*\*p < 0.001, \*\*p < 0.01, and \*p < 0.05.

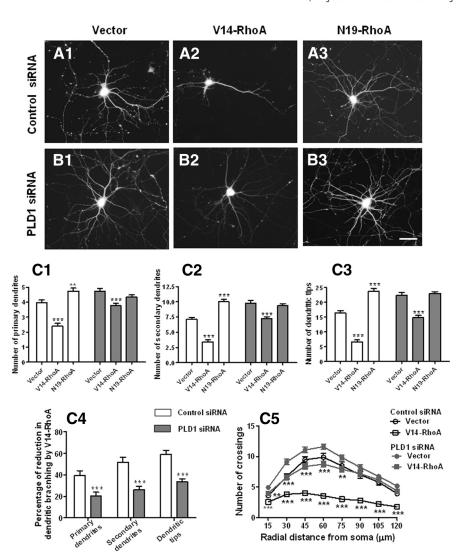
#### Results

# The expression and activity of PLD1 during the maturation of cultured hippocampal neurons

Primary culture of hippocampal neurons has long been a model system for studying the molecular mechanisms of neuronal development (Craig and Banker, 1994). After initially forming a lamellipodium (stage 1), neurons extend several short neurites that appear identical to one another (stage 2). About 2 d after plating, one of these minor neurites grows rapidly and becomes the axon (stage 3) (Dotti et al., 1988). Three or four days later, the remaining short neurites become mature dendrites (stage 4) and subsequently display dendritic spines (stage 5). To investigate the roles of PLD1 in dendritic formation, we first examined the expression and activity of PLD1 during the maturation cultured hippocampal neurons. The protein levels of PLD1 increased from DIV 1 to DIV 3, gradually decreased from DIV 3 to DIV7, and thereafter remained stable between DIV 7 and DIV 11 (Fig. 1A,B). In agreement with the protein levels, PLD activity increased from DIV 1 to DIV 3, subsequently decreased from DIV 3 to DIV 9, and finally remained stable between DIV 9 and DIV 11 (Fig. 1C). The protein level of PLD1 positively correlated with its activity (Fig. 1D). As shown in Figure 1E, both the number of primary and secondary dendrites and the number of dendritic tips increased from DIV 3 to DIV 11. Intriguingly, PLD1 protein levels and PLD activity were negatively correlated with the number of dendrites and dendritic tips, which suggests that PLD1 might be involved in the maturation of dendrites as a negative regulator (Fig. 1F-H).

# PLD1 negatively regulates dendritic branching

We cotransfected DIV 3 hippocampal neurons with GFP and WT-PLD1 or DN-PLD1 to increase the protein level or inhibit the activity of PLD1, respectively. To distinguish from endogenous PLD1, an HA tag was introduced into the N terminal of PLD1; this produced minimal interference with the normal function of PLD1 (Du et al., 2003). The coexpression of GFP and transfected PLD1 was confirmed by the immunostaining of HA in GFP-positive neurons (data not shown). To differentiate the primary dendrites and axon, we stained the neurons with anti-Ankyrin G, as Ankyrin G is specifically expressed in the initial segment of the axon (Srinivasan et al., 1988; Zhou et al., 1998). Only Ankyrin G-negative neurites were selected for further analysis of dendritic morphogenesis (data not shown). The neurons were fixed 3 d after transfection for morphological assay. We found that WT-PLD1 decreased, whereas DN-PLD1 increased, the number of primary and secondary dendrites compared with



**Figure 4.** Knockdown of PLD1 partially rescued the reduction of dendritic branching caused by overexpression of V-14 RhoA. *A1–A3*, Representative images of Control siRNA-treated neurons transfected with blank vector (*A1*), V14-RhoA (*A2*), or N19-RhoA (*A3*). *B1–B3*, Representative images of PLD1 siRNA-treated neurons transfected with blank vector (*B1*), V14-RhoA (*B2*), or N19-RhoA (*B3*). Neuronal morphology was visualized by cotransfection with GFP. Scale bar, 50  $\mu$ m. *C1–C3*, *C5*, Quantification of primary dendrites (*C1*), secondary dendrites (*C2*), dendritic tips (*C3*), and Sholl analysis (*C5*) in *A* and *B*. Control siRNA groups: n = 42 neurons for blank vector, n = 41 neurons for V14-RhoA, n = 31 neurons for N19-RhoA; PLD1 siRNA groups: n = 43 neurons for blank vector, n = 52 neurons for V14-RhoA, n = 49 neurons for N19-RhoA, \*\*\*p < 0.01, \*\*\*\*p < 0.001, compared with blank vector, one-way ANOVA followed by Newman–Keuls multiple-comparison test. *C4*, Percent reductions in dendritic branching induced by V14-RhoA in the presence of Control siRNA or PLD1 siRNA. \*\*\*\*p < 0.001, compared with Control siRNA, t test.

the control HA vector (Fig. 2*A*, *B1*,*B2*). Because the dendritic tips were at the terminal branches and reflected the complexity of dendrites, we next analyzed whether the number of dendritic tips was altered by the overexpression of WT-PLD1 and DN-PLD1. Consistent with the results from primary and secondary dendritic branching, the number of dendritic tips was decreased in neurons overexpressing WT-PLD1, whereas it was increased when the activity of PLD1 was inhibited by overexpression of DN-PLD1 (Fig. 2*A*,*B3*). Similar results were obtained when we overexpressed WT-PLD1 and DN-PLD1 in DIV 6 neurons (Fig. 2*C*,*D1*–*D3*). Finally, the effects of WT-PLD1 and DN-PLD1 on dendritic branching were confirmed by Sholl analysis (Fig. 2,*B4*,*D4*), which is the standard assay of the dendritic complexity (Sholl, 1953; Jaworski et al., 2005).

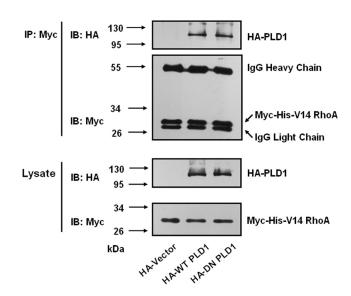
To further study the role of endogenous PLD1 in dendritic branching of hippocampal neurons, a strategy of knockdown by RNA interference was adopted. The efficiency of PLD1 siRNA was confirmed by the Western blot results from neuroblastoma N2a cells that were cotransfected with HA-WT-PLD1 and PLD1 siRNA (Fig. 3A2,A3). To test the efficiency of PLD1 RNA interference in neurons, we made a lentiviral PLD1 shRNA with the same target sequence as PLD1 siRNA. The PLD1 shRNA caused an 80% reduction in the expression of PLD1 in hippocampal neurons (Fig. 3A4,A5). We cotransfected GFP and Control siRNA or PLD siRNA in hippocampal neurons for 3 d before the analysis of dendritic branching. We found that PLD1 siRNA significantly increased dendritic branching in comparison with Control siRNA (Fig. 3B1,B2,C1–C4).

To exclude the off-target effects of PLD1 siRNA, rescue experiments were performed. An HA-PLD1 that is resistant to siRNA (hereafter referred to as RES-PLD1) was generated by replacing the nucleotides in the codon without altering the identity of the amino acids (Fig. 3A1). Western blot results showed that the PLD1 siRNA did not knock down the expression of RES-PLD1 (Fig. 3A2,A3); this result enabled us to use RES-PLD1 to rescue the dendritic phenotypes caused by PLD1 siRNA. As shown in Figure 3B3 and 3C1-C4, the increase in dendritic branching due to knockdown of PLD1 was corrected by RES-PLD1, confirming the specificity of the PLD1 siRNA. Together with the results from the DN-PLD1 (Fig. 2), these data indicate that PLD1 acts as a negative regulator of dendritic branching, which is consistent with the reduction of PLD1 expression and activity during the maturation of dendrites (Fig. 1).

## PLD1 acts as downstream target of RhoA to suppress dendritic branching

RhoA, one of the small Rho GTPases, has been implicated in the inhibition of dendritic growth and branching (Threadgill et al., 1997; Lee et al., 2000; Li et al., 2000; Nakayama et al., 2000; Chen and Firestein, 2007). Recent studies have suggested that RhoA directly interacts with and activates PLD1 both *in vitro* and *in vivo* (Hammond et al., 1997; Bae et al., 1998; Yamazaki et al., 1999; Du et al., 2000; Komati et al., 2005; Yoon et al., 2005; Gayral et al., 2006). In this study, we showed that PLD1 has the same effects as RhoA in reducing the number of dendritic branches. Together, these observations raise the possibility that PLD1 might be the downstream target of RhoA in the regulation of dendritic branching. In the following experiments, we tested this hypothesis in cultured hippocampal neurons.

Consistent with previous reports (Threadgill et al., 1997; Luo, 2000; Nakayama et al., 2000; Chen and Firestein, 2007), we found that overexpression of active RhoA (V14-RhoA) decreased dendritic branching, while N19-RhoA increased it (Fig. 4A, C1-C3, left three blank columns). However, there were no additive effects between N19-RhoA and PLD1 siRNA in promoting dendritic branching (Fig. 4A1,A3,B1,B3,C1-C3), suggesting that RhoA and PLD1 may share the same signaling pathway. Intriguingly, the inhibition of dendritic branching by V14-RhoA was partially rescued by knockdown of PLD1 (Fig. 4A1,A2,B1,B2,C1-C3). The percentage of reduction in dendritic branching by V14-RhoA was significantly decreased when PLD1 was knocked down (Fig. 4C4). As shown by the Sholl analysis, V14-RhoA had fewer effects in reducing dendritic complexity after knockdown of PLD1 (Fig. 4C5). These results indicate that PLD1 may act as the downstream target of RhoA in the regulation of dendritic branching, which is supported by the observations that RhoA interacts with PLD1 (Fig. 5) and activates PLD1 both in vitro and in vivo (Hammond et al., 1997; Bae et al., 1998; Yamazaki et al., 1999; Du et al., 2000; Komati et al., 2005; Yoon et al., 2005; Gayral et al., 2006).



**Figure 5.** Coimmunoprecipitation (IP) of HA-WT-PLD1, HA-DN-PLD1, and Myc-His-V14-RhoA in N2a cells.

### PLD1 reduces dendritic branching through PA

The enzymatic action of mammalian PLD1 produces PA and choline. Because free choline is not thought to fulfill any intracellular signaling role (Pelech and Vance, 1984), most of the biological function of PLD1 is thought to be mediated by PA, which is a bioactive lipid (Frohman et al., 1999; Jenkins and Frohman, 2005; Cockcroft, 2009). Next, we tested whether the effects of PLD1 on dendritic branching depend on PA. To this end, we overexpressed PLD1 in hippocampal neurons with the application of 0.5% 1-butanol in the culture medium. We found that 1-butanol can block the production of PA catalyzed by PLD (Fig. 6 C6) (Chen et al., 1997; Siddhanta et al., 2000). Interestingly, the neurons treated with 1-butanol had more dendritic branching than the neurons treated with 2-butanol (the inactive analog of 1-butanol) (Fig. 6*A1*,*B1*,*C1*–*C3*, vector columns), which suggests a role of endogenous PA and PLD activity in the regulation of dendritic branching. Moreover, 1-butanol rescued the reduction of dendritic branching caused by overexpression of WT-PLD1 (Fig. 6A1, A2, B1, B2, C1–C3). The suppression of dendritic branching and complexity by overexpression of WT-PLD1 were significantly restored in 1-butanol-treated neurons compared with control neurons (Fig. 6C4,C5). These results indicate that PLD1 inhibits dendritic branching through PA.

As mentioned above, PLD1 acts as one of the downstream targets of RhoA in reducing dendritic branching. If PLD1 inhibits dendritic branching through PA, we speculated that PA should also act as the downstream target of RhoA in the regulation of dendritic branching. To test this hypothesis further, we transfected hippocampal neurons with active RhoA (V14-RhoA) in the presence of 1-butanol or its inactive analog, 2-butonal. We found that the inhibition of dendritic branching and complexity by V14-RhoA was partially restored by 1-butanol in comparison with 2-butonal (Fig. 7). These results suggested that PA is also involved in the effects of RhoA on dendritic branching. Together, these data support the working model that RhoA, PLD1, and PA may constitute one signaling pathway in the regulation of dendritic branching.

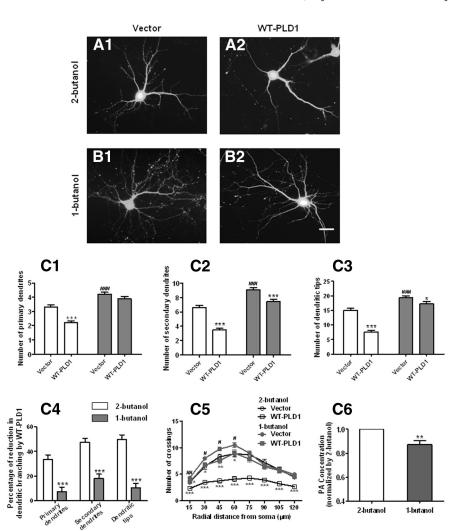
#### Discussion

In this study, we have demonstrated that PLD1 and its enzymatic product PA act downstream of RhoA in suppressing dendritic

growth and branching. Several lines of evidence support our hypothesis. First, RhoA can interact with PLD1 when they are coexpressed in N2a cells, which is consistent with previous studies (Hammond et al., 1997; Bae et al., 1998; Yamazaki et al., 1999, Du et al., 2000; Komati et al., 2005; Yoon et al., 2005; Gayral et al., 2006). Second, the reduction of dendritic branching caused by active RhoA can be partially rescued by PLD1 siRNA. Third, 1-butonal, which blocks the production of PA by PLD (Chen et al., 1997; Siddhanta et al., 2000), reduced the efficacy with which PLD1 and RhoA suppressed dendritic branching. Most of the known effectors of RhoA in dendritic morphogenesis belong to protein kinases such as Rho kinase and LIM kinase 1 (Maekawa et al., 1999; Ohashi et al., 2000; Lou et al., 2001; Lin et al., 2003; Montanez et al., 2009; Fonseca et al., 2010). In this study, we identified a lipid messenger, PA, as a novel downstream target of RhoA in the regulation of dendritic branching, which will expand our understanding of how RhoA signaling regulates dendritic morphogenesis.

We could not exclude the possibility that PLD1 regulates dendritic branching through the interaction with signaling proteins other than PA. However, most of PLD1 function identified so far depends on PA or the metabolites of PA (Cai et al., 2006; Gayral et al., 2006; Zeniou-Meyer et al., 2007; Hashimoto et al., 2008; Disse et al., 2009; Bach et al., 2010; Elvers et al., 2010), because PA is the only bioactive lipid among the enzymatic products of PLD1. Here we showed that PA was one of the critical molecules that execute the effects of PLD1 on dendritic branching. Exactly how PA is involved in dendritic branching is not clear. Several signaling pathways could mediate the effects of PA

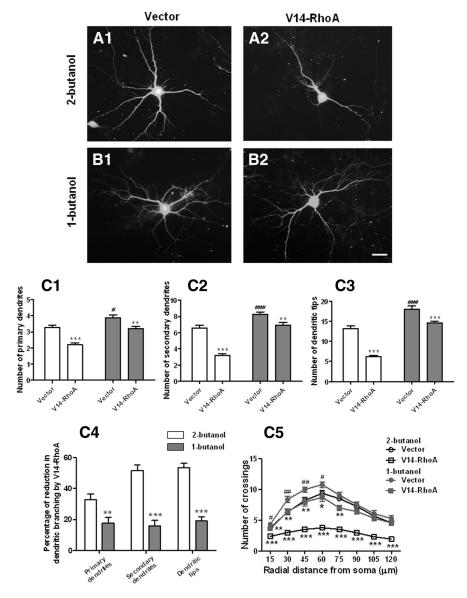
on dendritic branching. PA can be metabolized to other lipids with potent bioactivity. For instance, PA can be converted to diacylglycerol and lysophosphatidic acid (LPA). LPA has been shown to induce the retraction of neurites (Tigyi and Miledi, 1992; Jalink et al., 1993, 1994). PA can recruit some signaling molecules, such as ADP-ribosylation factor 6 (Manifava et al., 2001), which has been demonstrated to reduce dendritic branching (Hernández-Deviez et al., 2002; Sakagami et al., 2004). PA also activates certain protein kinases, such as phosphatidylinositol 4-phosphate 5-kinase (PIP5K) through direct interaction (Moritz et al., 1992; Jenkins et al., 1994). Overexpression of PIP5K was sufficient to induce neurite retraction (Halstead et al., 2010), which is similar with the dendritic phenotypes caused by overexpression of PLD1. Moreover, the activity of PIP5K is essential for neurite retraction in response to LPA (van Horck et al., 2002; Yamazaki et al., 2002), one metabolite of PA. In hippocampal neurons, the axon branching was inhibited by overexpression of PIP5K (Hernández-Deviez et al., 2004). Future



**Figure 6.** Use of 1-butanol to ameliorate the dendritic phenotypes caused by overexpression of WT-PLD1. **A1**, **A2**, Representative images of 2-butonal-treated neurons transfected with blank vector (**A1**) or WT-PLD1 (**A2**). **B1**, **B2**, Representative images of 1-butonal-treated neurons transfected with blank vector (**B1**) or WT-PLD1 (**B2**). Scale bar, 50  $\mu$ m. **C1–C3**, **C5**, Quantification of primary dendrites (**C1**), secondary dendrites (**C2**), dendritic tips (**C3**), and Sholl analysis (**C5**) in **A** and **B**. 2-butanol-treated neurons: n=41 for blank vector, n=49 for WT-PLD1; 1-butanol-treated neurons: n=48 for blank vector, n=49 for WT-PLD1, \*p<0.05, \*\*p<0.01, \*\*\*p<0.01, \*\*\*p<0.01, compared with blank vector, t test. \*\*t0 t0.00, \*\*\*t0 t0.01, compared with blank vector in 2-butanol group, t1 test. \*\*t0, Percent reductions in dendritic branching caused by overexpression of WT-PLD1 in the presence of 2-butonal or 1-butonal. \*\*\*t0 t0.001, compared with 2-butonal-treated neurons, t1 test. \*\*C6\$, Quantification of PA concentration in the presence of 1-butanol or 2-butanol. \*\*\*t0.01, compared with 2-butonal-treated neurons, t1 test.

studies will address the role of PIP5K signaling in dendritic branching and whether PIP5K is involved in the RhoA-PLD1-PA pathway.

In agreement with the role of PLD1 as a downstream target of RhoA, we found that PLD1 itself is an intrinsic suppressor of dendritic growth and branching. Knockdown of PLD1 or inhibition of its activity increases dendritic branching, whereas overexpression of PLD1 decreases the complexity of dendrites. PLD1 acts as a negative regulator of dendritic arborization. This finding is also supported by the reduction of the expression and activity of PLD1 during dendritic maturation. PLD1 siRNA could not fully rescue the effects of RhoA on dendritic branching, which suggested that RhoA has other downstream signaling pathways in addition to PLD1 and PA. Several protein kinases, such as Rho kinase and LIM kinase 1, have been implicated in mediating the biological functions of RhoA (Matsui et al., 1996; Maekawa et al., 1999; Ohashi et al., 2000). Some of these kinases might cross talk with PLD1, as it has been shown that PLD1 could be activated by Rho kinase (Schmidt et al., 1999). Interestingly, Rho kinase can also inhibit dendritic branching in hippocampal



**Figure 7.** Use of 1-butanol to partially restore the decreased dendritic branching caused by overexpression of V14-RhoA. **A1**, **A2**, Representative images of 2-butanol-treated neurons transfected with control vector (**A1**) or V14-RhoA (**A2**). **B1**, **B2**, Representative images of 1-butanol-treated neurons transfected with control vector (**B1**) or V14-RhoA (**B2**). Scale bar, 50  $\mu$ m. **C1-C3**, **C5**, Quantification of primary dendrites (**C1**), secondary dendrites (**C2**), dendritic tips (**C3**), and Sholl analysis (**C5**) in **A** and **B**. 2-butanol-treated neurons: n=44 for control vector; n=50 for V14-RhoA; 1-butanol-treated neurons: n=41 for control vector, n=49 for V14-RhoA, \*p<0.05, \*\*p<0.01, \*\*\*p<0.01, compared with control vector in 2-butanol group, t test. **C4**, Percent reductions in dendritic branching caused by V14-RhoA in the presence of 2-butonal or 1-butonal. \*\*p<0.01, \*\*\*p<0.001, compared with 2-butonal-treated neurons, t test.

neurons (Nakayama et al., 2000), which raises the possibility that PLD1 may also act downstream of Rho kinase in the regulation of dendritic branching.

Our work corroborates recent findings that PLD1 regulates neurite growth or initiation. PLD1 was involved in basic fibroblast growth factor-induced neurite outgrowth in H19–7 cells (Oh et al., 2007). PLD1 was also important in neurite initiation in neural stem cells (Sung et al., 2001; Yoon et al., 2005). In this study, we found that PLD1 specifically restricted dendritic branching when the expression or activity of PLD1 was manipulated during the maturation of dendrites, i.e., from DIV 3 to DIV 9 in cultured hippocampal neurons. The differences between the previous studies and our results may be due to the type of cells used and/or the different developmental stages or the specific

identity of the neurites (i.e., axons or dendrites). Intriguingly, PLD1 has also been implicated in calcium-dependent exocytosis (Jones et al., 1999; Bader and Vitale, 2009) and neurotransmitter release (Humeau et al., 2001). Together, these observations demonstrated the potentially important roles of PLD1 in both neural development and neurotransmission.

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