LMTK1 regulates dendritic formation by regulating movement of Rab11A-positive endosomes

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ABSTRACT Neurons extend two types of neurites—axons and dendrites—that differ in structure and function. Although it is well understood that the cytoskeleton plays a pivotal role in neurite differentiation and extension, the mechanisms by which membrane components are supplied to growing axons or dendrites is largely unknown. We previously reported that the membrane supply to axons is regulated by lemur kinase 1 (LMTK1) through Rab11A-positive endosomes. Here we investigate the role of LMTK1 in dendrite formation. Down-regulation of LMTK1 increases dendrite growth and branching of cerebral cortical neurons in vitro and in vivo. LMTK1 knockout significantly enhances the prevalence, velocity, and run length of anterograde movement of Rab11A-positive endosomes to levels similar to those expressing constitutively active Rab11A-Q70L. Rab11A-positive endosome dynamics also increases in the cell body and growth cone of LMTK1-deficient neurons. Moreover, a nonphosphorylatable LMTK1 mutant (Ser34Ala, a Cdk5 phosphorylation site) dramatically promotes dendrite growth. Thus LMTK1 negatively controls dendritic formation by regulating Rab11A-positive endosomal trafficking in a Cdk5-dependent manner, indicating the Cdk5-LMTK1-Rab11A pathway as a regulatory mechanism of dendrite development as well as axon outgrowth.

INTRODUCTION Neurons are highly polarized cells with two different types of processes called axons and dendrites. Each neuron has a single, long, unbranched process called an axon from which neurons transmit signals. Each neuron can have multiple, branched dendrites, which are the sites where neurons receive signals. Correct establishment of axons and dendrites is critical for proper neuronal network formation, and elucidating the molecular mechanisms of their formation is important for understanding neurodevelopmental disorders such as autism and schizophrenia (Bellon, 2007; Pardo and Eberhart, 2007). Neurite formation is a complicated process involving both cytoskeletal dynamics and membrane trafficking. Although axonal determination and outgrowth have been extensively studied and many critical cytoskeletal and signaling proteins have been reported (Arimura and Kaibuchi, 2007; Shelly and Poo, 2011), little attention has been paid to the mechanism of membrane transport in these processes (Sann et al., 2009; Yap and Winckler, 2012). In developing neurons, the supply of membrane to axons precedes its distribution to dendrites. Thus regulation of membrane trafficking is essential for accurate spatiotemporal delivery of membrane components.

Rab small GTPases are major regulatory proteins involved in membrane transport (Takai et al., 2001; Zerial and McBride, 2001; Stenmark, 2009). Rab small GTPases exert their regulatory control biochemically via the successive binding of GTP (active) and hydrolysis to...
yield the GDP-bound form (inactive; Takai et al. 2001; Zerial and McBride, 2001; Stenmark, 2009). More than 60 members of the Rab family have been identified, and each functions in a distinct membrane compartment. Previous analysis of Rab function has been conducted mainly in nonpolarized cells. Neurons have a specialized cell shape, and membrane trafficking must operate in a region-specific manner. However, how particular membrane proteins are delivered to axons or dendrites and whether neuronal Rabs share distinct or common functions between axons and dendrites are not known.

Rab11 is a well-known regulator of recycling endosomes (Ullrich et al., 1996; Sonnichsen et al., 2000). In nonneuronal cells, Rab11 regulates trafficking of endocytic vesicles to the pericentriolar endocytic recycling compartment (ERC) and the recycling pathway from the ERC to the plasma membrane (Ullrich et al., 1996; Sonnichsen et al., 2000; Takahashi et al., 2007). Rab11 has been proposed to be involved in several types of neuronal membrane transport, such as synaptic vesicle recycling in presynaptic regions, α-amino-3-hydroxy-5-methyl-4-isoxazole-propionic acid (AMPA) receptor recruitment in postsynaptic regions, and neuronal migration during development (Park et al., 2004; Brown et al., 2007; Sann et al., 2009; Kawachi et al., 2010; Yap and Winckler, 2012). Although several reports indicate a role for Rab11 in axonal outgrowth (Ascano et al., 2009; Eva et al., 2010; Takano et al., 2012), the role of Rab11A in dendrite outgrowth is unknown.

Lemur kinase 1 (LMTK1) is an uncharacterized Ser/Thr kinase that is highly expressed in the mammalian brain (Baker et al., 2001; Tomomura et al., 2007). We showed that LMTK1 activity is regulated by phosphorylation at Ser-34 by cyclin-dependent kinase 5 (Cdk5)-p35 and LMTK1 regulates trafficking of recycling endosomes in CHO-K1 cells (Takano et al., 2010). We recently reported that LMTK1 controls axon outgrowth in cortical neurons by modulating Rab11A function (Takano et al., 2012). Here we investigate the role of LMTK1 in Rab11A-positive endosomal trafficking during dendrite formation in vitro and in vivo and the regulation of LMTK1 by Cdk5. We show that LMTK1 negatively regulates dendrite growth and arborization by modulating Rab11A-positive endosome movement.

RESULTS
Down-regulation of LMTK1 increases dendrite growth and branching

We recently reported that LMTK1 suppresses axonal outgrowth by regulating Rab11A activity in a Cdk5-dependent manner (Takano et al., 2012). Here we investigate a possible role for LMTK1 in dendritic formation. We first examined the involvement of LMTK1 in dendritic growth of primary mouse cortical neurons by knocking down LMTK1 via a specific microRNA (miRNA) at 5 d in vitro (DIV)—when dendrites begin to elongate from cultured neurons after establishing an axon. LMTK1 was effectively knocked down at 7 DIV, as seen with immunoblotting (Supplemental Figure S1A) and immunostaining (Supplemental Figure S1B). We observed that LMTK1 knockdown neurons had more-elaborate dendritic processes than control neurons treated with scrambled-sequence (miRNA-SC; Figure 1, A–E). These neurites were identified as dendrites because they were labeled with the postsynaptic markers postsynaptic density protein 95 (PSD95) and AMPA receptor GluR2 (Supplemental Figure S1, C and D). We counted the neurites and evaluated arborization using Sholl analysis. The number of dendrites was elevated in LMTK1-knockdown neurons, 6.6 ± 0.1 and 5.7 ± 0.1, respectively, with the two different miRNAs, compared with 4.9 ± 0.2 in miRNA-SC control neurons (Figure 1B). Similarly, the number of branches in LMTK1-knockdown neurons, 10.5 ± 0.3 and 8.6 ± 0.5, respectively, was greater than the number, 6.9 ± 0.3, in control neurons (Figure 1C). Therefore LMTK1 knockdown increased the total primary dendrite length by ~1.7-fold (Figure 1D). The numbers of crossings at both proximal (40 μm) and distal (280 μm) lines from the cell body were greater in LMTK1-knockdown neurons than control neurons (Figure 1E). Thus LMTK1 plays a role in limiting dendrite growth and arborization of cortical neurons.

We noted in the previous report that when axon growth is stimulated by down-regulation of LMTK1, the length of other neurites was attenuated (Takano et al., 2012). Therefore we next investigated how overelongation of the axon in LMTK1-knockdown neurons affects dendrite outgrowth. Three possibilities exist: 1) dendrite arborization will be suppressed due to prior consumption of membrane components by overly long axons, 2) overelongation of the axon will not affect dendrite growth, and 3) LMTK1 knockdown also stimulates dendrite outgrowth as was observed in axons. To test these possibilities, we introduced knockdown vectors into cortical neurons at 0 DIV and then analyzed dendrites after growth at 5 DIV (unpublished data) and 7 DIV (Figure 1F and Table 1). Unexpectedly, we found that cortical neurons expressing LMTK1 miRNAs extended more neurites than did those expressing mi-RNA-SC at 5 DIV (unpublished data). LMTK1 knockdown increased the number of dendrites (Figure 1G), the number of branch points (Figure 1H), the total length of primary dendrites (Figure 1I), and the number of crossings at both proximal and distal regions from the cell body (Figure 1J). These data are consistent with the third possibility—dendrite formation was enhanced after down-regulation of LMTK1 independent of prior overelongation of the axon.

Dendritic arborization is enhanced in LMTK1−/− mouse neurons

We confirmed the foregoing results in neurons prepared from brains of LMTK1−/− mice. LMTK1−/− neurons exhibited significantly more and longer dendritic processes (Figure 2A), larger numbers of dendrites (Figure 2B), more branches (Figure 2C), longer total primary dendrite length (Figure 2D; see Table 1 for a detailed comparison), and increased number of crossings compared with LMTK1+/+ neurons (Figure 2E).

To confirm these findings in vivo, we compared the dendritic morphology of pyramidal neurons in the cerebral cortex of LMTK1−/− and LMTK1−/− mice with Golgi staining. We analyzed the developing brain at P7. Pyramidal neurons in layer V of the cerebral cortex have two types of dendrites—a thick apical dendrite extending toward the pial surface, and multiple basal dendrites extending basolaterally. Growth of both apical and basal dendrites was stimulated in LMTK1−/− brains (Figure 2F). Pyramidal neurons in LMTK1−/− brains had a longer apical dendrite (Figure 2G), and the number and length of the secondary dendrites increased (Figure 2, H and I). The number and length of basal dendrites also increased in LMTK1−/− brains (Figure 2, J and K). Thus LMTK1 negatively regulates dendrite growth in pyramidal neurons of the cerebral cortex in vivo.

During observation of LMTK1−/− neurons in culture, we noticed many thin protrusions along the dendrites of LMTK1−/− neurons (Figure 2L). The number of protrusions was 1.3 ± 0.1 in LMTK1+/+ neurons and 5.2 ± 0.1 in LMTK1−/− neurons (Figure 2M). These protrusions were also observed in LMTK1-knockdown neurons (Supplemental Figure S2, A and B). They might be immature dendritic branches or spines.
was found on small puncta in dendrites, which were labeled with Rab11A and partly with Rab5A but not with Rab7 (Figure 3A and Supplemental Figure S3), suggesting that LMTK1 also regulates dendrite growth via Rab11A-positive endosomes.

In the cell body, LMTK1 and Rab11A are concentrated at the ERC in the perinuclear region close to the axon, where Rab11A-positive endosomes are transported to the axon (Takano et al., 2012). It is not known whether the ERC changes its location after its delivery destination changes from the axon to dendrites. We looked carefully at the localization of endogenous LMTK1 and EGFP-Rab11A in the cell body of cultured neurons at 3 and 7 DIV. Endogenous LMTK1 localized at the perinuclear region close to the base of the axon, the longest process, at both 3 and 7 DIV (Figure 3B, arrow). Rab11A accumulated at the base of the axon in primary neurons at 3 DIV, and similar localization was observed at 7 DIV (Figure 3C, arrow). This distribution was not altered in LMTK1−/− neurons (Figure 3C). These results suggest that the pericentriolar ERC, where LMTK1 and Rab11A accumulate, retains its position at the base of the axon in developing cultured cortical neurons.

The foregoing results indicate that recycling endosomes, which have to be delivered to dendrites, would be transported in the cell body to dendrites. We next examined how recycling endosomes move in the cell body toward dendrites from the pericentriolar ERC. We observed the movements of Rab11A-positive endosomes using real-time imaging. An example is shown in Figure 3D. In fact, endosomes moved from the pericentriolar ERC toward the dendrites in the cytoplasm between the nucleus and plasma membrane (Figure 3D, yellow arrowheads; Supplemental Videos S1 and S2). No particular accumulation or retention of endosomes was observed at the base of dendrites. Vesicles appeared to follow the same path as previous vesicles. Budding of small Rab11A-positive vesicles from the pericentriolar ERC was increased in LMTK1−/− neurons compared to LMTK1+/+ neurons (Figure 3D, arrowheads in kymographs; Supplemental Video S2). These results suggest that the transport system in the cell body changes when dendrites are committed to grow.

**Increased prevalence, rate, and distance of anterograde movement of Rab11A-positive endosomes in dendrites of LMTK1−/− neurons**

We analyzed how down-regulation of LMTK1 affects the movement of Rab11A-positive endosomes in primary and secondary dendrites using real-time imaging. Because the movements were similar in
primary and secondary dendrites, we followed the movement of vesicles mainly in secondary dendrites, where Rab11A-positive endosomes were easier to see due to there being fewer endosomes and thinner processes. Approximately 81% of Rab11A-positive endosomes were mobile in dendrites of LMTK1+/− neurons (Figure 4, A and B). Among them, 37 ± 1.2% of endosomes moved from proximal to distal (anterograde), 31 ± 1.7% moved from distal to proximal (retrograde), and 32 ± 1.3% moved bidirectionally in dendrites of LMTK1+/− neurons (Figure 4C). Approximately 89% of Rab11A-positive endosomes were mobile in LMTK1−/− neurons, which was significantly higher than in LMTK1+/− neurons (Figure 4, A and B). Further, the movement was biased toward the anterograde direction. Anterograde movement increased to 47% of total, whereas retrograde movement decreased to 19% in LMTK1−/− dendrites (Figure 4C). The velocities of the endosomes were 1.24 and 1.03 μm/s for anterograde and retrograde movements, respectively, in LMTK1−/− dendrites compared with 0.94 and 0.86 μm/s, respectively, in LMTK1+/− neurons (Figure 4, D and E). The most striking difference was in the run length, which is the distance traveled by the endosome without pausing during a 20-μm observation window. Although the longest run length was 4 and 5 μm for anterograde and retrograde movements, respectively, in LMTK1+/− neurons, it was >11 μm in both directions in LMTK1−/− dendrites (Figure 4, F and G). We compared the directionality of Rab11A-positive endosomes in dendrites and axons because the polarity of microtubules differs between axons and dendrites (Baas et al., 1988; Stepanova et al., 2003). In LMTK1+/− neurons, at times when axons (3 DIV) and dendrites (7 DIV) were actively elongating, the proportion of anterograde movement in axons and dendrites was 51 and 37%, which is higher than the 45 and 31% for retrograde movement, respectively (Table 2). In LMTK1−/− neurons, the proportion of anterograde movement increased to 58% in axons at 3 DIV and 47% in dendrites at 7 DIV, whereas that of retrograde movement decreased to 40%...
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Movement of wild-type Rab11A-positive endosomes in LMTK1+/− neurons and constitutively active Rab11A-Q70L–positive endosomes in LMTK1+/− neurons

We previously showed that LMTK1 decreases the activity of Rab11A (Takano et al., 2012). To assess more carefully the interaction of LMTK1 with Rab11A in endosome movement, we used constitutively active (ca) Rab11A-Q70L, which is GTPase deficient, and dominant-negative (dn) Rab11A-S25N, which is a GDP-locked mutant (Ullrich et al., 1996; Ren et al., 1998). We compared the movement of wtRab11A-positive endosomes in LMTK1−/− neurons and constitutively active Rab11A-Q70L–positive endosomes in LMTK1+/+ neurons (Figure 6A). The caRab11A-Q70L–positive endosomes showed similar movement characteristics to wtRab11A-positive endosomes in LMTK1−/− neurons (Figure 6A). Approximately 85% of caRab11A-Q70L–labeled endosomes were mobile in dendrites of LMTK1+/+ neurons, a little higher than the 78% of wtRab11A endosomes that were mobile in LMTK1−/− neurons (Figure 6, A and B, and Supplemental Video S7). In contrast, only 22% of dn-Rab11A-S25N–positive endosomes were mobile (Figure 6, A and B, and Supplemental Video S8). Among mobile caRab11A-Q70L endosomes, ∼49% of endosomes showed anterograde movement, which was identical to the 49% of wtRab11A-positive endosomes in LMTK1−/− dendrites that were mobile but higher than the 41% of mobile wtRab11A endosomes in LMTK1+/+ dendrites (Figure 6C). The caRab11A-Q70L vesicles moved at mean velocities of 0.99 (anterograde) and 0.96 μm/s (retrograde), which were slightly faster than the corresponding velocities of 0.85 (anterograde) and 0.85 μm/s (retrograde) for wtRab11A-labeled endosomes in LMTK1+/+ neurons (Figure 6, E and F, and Table 2). The size of caRab11A-Q70L vesicles was 0.17 ± 0.01 μm², which was smaller than the 0.4 ± 0.02 μm² for wtRab11A-labeled endosomes in LMTK1+/− neurons (Figure 6D). The percentage of bidirectionally moving caRab11A-Q70L endosomes (15 ± 2.4%) was lower than the 26% for wtRab11A-labeled endosomes as estimated in the primary dendrites of LMTK1+/− neurons. The size of wtRab11A-positive endosomes in LMTK1+/− neurons was and, remarkably, 19% in axons and dendrites, respectively (Table 2).

Of interest, the proportion of bidirectional movement of Rab11A-positive endosomes was considerably higher in dendrites of both LMTK1+/+ (32%) and LMTK1+/− (34%) neurons than in axons of LMTK1+/+ (8%) and LMTK1−/− (4%) at 7 DIV (Figures 4C and 5C). The increased bidirectional movement in dendrites may reflect the mixed polarity of microtubules (Baas et al., 1988; Stepanova et al., 2003).
neurons (Figure 6, E and F). Many caRab11A-Q70L endosomes in LMTK1+/− neurons and wtRab11A-positive endosomes in LMTK1−/− neurons moved >11 μm without pausing, whereas the peak of the run length in the anterograde and retrograde directions was 5 μm for wtRab11A endosomes (Figure 6, G and H). The run length of wtRab11A-positive endosomes in LMTK1−/− neurons increased to >11 μm, compared with 5 μm of caRab11A-Q70L vesicles (Figure 6, G and H). Thus, as measured by several kinetic parameters, the movement of wtRab11A-positive endosomes in LMTK1−/− neurons was similar but not identical to that of caRab11A-Q70L-positive endosomes, suggesting that the Rab11A in LMTK1−/− neurons is largely in the active GTP form.

Movement of Rab11A-positive endosomes in growth cones of LMTK1−/− neurons

Recycling endosomes that moved to the ends of neurites may be incorporated into the plasma membrane at growth cones. However, no observations of the dynamics of Rab11A-positive endosomal movements in growth cones have been reported. Therefore we analyzed the movements of Rab11A-positive endosomes in the growth cones of dendrites in LMTK1+/+ and LMTK1−/− neurons. To visualize the entire growth cone, we cotransfected a plasmid encoding a red fluorescent protein (DsRed) with a plasmid encoding EGFP-Rab11A into primary neurons. LMTK1−/− neurons showed a growth cone area (30 ± 4.2 μm²) that was larger than that of LMTK1+/+ growth cones (17 ± 2.6 μm²; Figure 7, A and B). Rab11A-positive endosomes were smaller in LMTK1−/− growth cones than in LMTK1+/+ growth cones (Figure 7C). We observed both moving and stationary endosomes in growth cones. The percentage of moving Rab11A-positive vesicles was 84 ± 0.9% in LMTK1−/− neurons, which was higher than in LMTK1+/+ neurons (57 ± 3.4%; Figure 7, D and E, and Supplemental Videos S9 and S10). We frequently observed vesicles that moved to the tip of the growth cone and disappeared (Figure 7E). During a 160-s recording, 5.8 ± 0.6 Rab11A-positive endosomes disappeared at the tip of growth cones in LMTK1−/− neurons, whereas only 2.7 ± 0.5 were in LMTK1+/+ growth cones (Figure 7F). However, the number of vesicles returning to the dendritic shaft was lower in LMTK1−/− neurons (1.9 ± 0.3) compared with LMTK1+/+ neurons (2.8 ± 0.3; Figure 7H). The difference will result in

FIGURE 3: Localization of LMTK1 and Rab11A in the cell body and dendrites and movement of Rab11A-positive endosomes in the cell body of cortical neurons. (A) Colocalization of LMTK1 and Rab11A in the cell body and dendrites. Plasmids encoding LMTK1 and EGFP-Rab11A were cotransfected into primary cortical neurons at 0 DIV, and cells were visualized by immunostaining with anti-LMTK1 at 7 DIV (left). Middle, Rab11A. Right, merged image; dendrites in which LMTK1 also colocalized with Rab11A (arrows). Bar, 10 μm (cell body), 5 μm (dendrites). (B) Localization of endogenous LMTK1 at 3 (left) or 7 DIV (right). Endogenous LMTK1 was immunostained with anti-LMTK1. Bar, 10 μm. (C) Localization of Rab11A in LMTK1+/+ (top) or LMTK1−/− (bottom) cortical neurons at 3 DIV (left) or 7 DIV (right). Bar, 10 μm. (D) Movement of Rab11A-positive endosomes in the cell body of LMTK1+/+ (top) or LMTK1−/− (bottom) neurons. Neurons were transfected with a plasmid encoding EGFP-Rab11A at 6 DIV. Movement of Rab11A-positive endosomes was observed with real-time imaging. High-magnification images of the area indicated by the dotted line are shown for five frames with 1- to 3-s intervals. The movement of Rab11A-positive endosomes is labeled by arrowheads. Right, kymographs of Rab11A-positive endosomes budding from pericentriolar ERC and moving toward the dendrites. Arrowheads indicate budding of endosomes. N, nucleus. Bar, 20 μm.
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Cdk5-p35 can phosphorylate LMTK1 at Ser-34, thereby modulating LMTK1’s ability to regulate axonal outgrowth (Takano et al., 2012). We therefore examined whether the function of LMTK1 on dendrite arborization is also regulated by Cdk5-p35-mediated phosphorylation. The phosphorylation of LMTK1 at Ser-34 increased gradually from 3 to 9 DIV in cultured cortical neurons and then remained constant until 13 DIV, in parallel with the expression of the Cdk5 activator p35 (Supplemental Figure S4). Because our phospho-Ser-34 antibody was not sensitive enough to detect endogenous LMTK1 phosphorylation, we examined the cellular localization of phospho–Ser-34 by expressing exogenous LMTK1 in cortical neurons. Phospho-LMTK1 was abundantly expressed in the perinuclear region of the cell soma (Figure 8A, top, arrow) and was seen on small puncta in dendrites (Figure 8A, bottom). We next examined the effect of overexpressing two myc-labeled LMTK1 mutants—the S34A nonphosphorylatable form and the S34D phosphorylation mimic form—on dendritic arborization. Neurons were visualized with DsRed cotransfected with LMTK1-myc mutants. Neurons expressing LMTK1-S34A showed remarkably well-developed dendritic arborization compared with neurons expressing LMTK1-S34D or LMTK1-WT, the dendrites of which were similar to those of control neurons expressing LMTK1-WT, the dendrites of which were similar to those of control neurons expressing LMTK1-WT. These results suggest that LMTK1 also suppresses the fusion of Rab11A-positive endosomes into the tip of neurites, the final step of the recycling pathway.

**Effect of phosphorylation of LMTK1 at Ser-34 on dendritic arborization**

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**FIGURE 4:** Dynamics of Rab11A-positive endosomes in dendrites of LMTK1−/− neurons. (A) Movement of Rab11A-positive endosomes in a dendrite of a LMTK1+/+ or LMTK1−/− neuron. Neurons were transfected with a plasmid encoding EGFP-Rab11A. Movement of Rab11A-positive endosomes in secondary dendrites was observed with real-time imaging. Sequential frames for 12 s with a 4-s interval are shown for LMTK1+/+ and LMTK1−/−. The cell body is on the left side. The movement of a typical Rab11A-positive endosome is indicated with arrows. Bottom, kymographs of Rab11A-positive endosomes moving in dendrites. Bar, 5 μm. (B) Percentage of moving (light gray) or stationary (dark gray) Rab11A-positive endosomes in dendrites. (C) Percentage of Rab11A-positive endosomes moving anterogradely (gray), retrogradely (light gray), or bidirectionally (dark gray) in dendrites (n = 381 endosomes in LMTK1+/+ neurons and n = 432 endosomes in LMTK1−/− neurons). (D, E) Velocity of Rab11A-positive endosomes in dendrites. Relative frequency of Rab11A-positive endosomes moving at the indicated velocities in the anterograde (D) or retrograde (E) direction in dendrites of LMTK1+/+ (light gray) or LMTK1−/− (dark gray) neurons (n = 150 endosomes each for LMTK1+/+ and LMTK1−/−). (F, G) Run length of Rab11A-positive endosomes in dendrites. The distance that Rab11A-positive endosomes moved without pausing was measured in a 20-μm observation window. Relative frequency of Rab11A-positive endosomes moving at the indicated run length in the anterograde (F) or retrograde (G) direction in dendrites of LMTK1+/+ (light gray) or LMTK1−/− (dark gray) neurons (n = 100 endosomes each for LMTK1+/+ and LMTK1−/−).
observed for endosomes that have incorporated caRab11A-Q70L; that is, Rab11A is highly active in the absence of LMTK1, and thus LMTK1 seems to maintain somewhat low levels of Rab11A activity. This is consistent with our previous results showing that LMTK1−/− mouse brains have higher levels of active Rab11A (Takano et al., 2012). Indeed, premature outgrowth of dendrites in the LMTK1−/− brain was observed at early postnatal days, and thus LMTK1 could prevent dendritic hyperplasia by regulating membrane supply. LMTK1 function may therefore be important for establishing proper neuronal connections during brain development. It may be worth mentioning the current idea that premature or accelerated neuronal growth could be a cellular or molecular mechanism of autism spectrum disorder (Chomiak and Hu, 2013). It would be interesting to analyze the cytoarchitecture of autism-related brain regions, as well as behaviors, of LMTK1−/− mice.

Rab11 is a general marker of recycling endosomes (Ullrich et al., 1996; Sonnichsen et al., 2000). In neurons, Rab11 is suggested to regulate membrane supply (Figure 8B). The number of dendrites and branches and the total primary dendrite length were significantly increased in neurons expressing LMTK1-S34A (Figure 8, C–E). Sholl analysis revealed an increased number of crossings at both proximal (40 μm) and distal (280 μm) sites in LMTK1-S34A–expressing neurons compared with those expressing LMTK1-S34D or LMTK1-WT (Figure 8F). These results indicate that Cdk5, by phosphorylating Ser-34 of LMTK1, also regulates the effect of LMTK1 on dendritic arborization.

**DISCUSSION**

Down-regulation of LMTK1 induced longer dendrites with increased numbers of branches, indicating that LMTK1 suppresses dendrite growth and arborization. This activity of LMTK1 was mediated by Rab11A. Rab11A movements correlated well with the growth and branching of dendrites. The prevalence, speed, and run length of the outward movements of Rab11A-positive endosomes increased upon knockdown or knockout of LMTK1 in neurons to levels observed for endosomes that have incorporated caRab11A-Q70L; that is, Rab11A is highly active in the absence of LMTK1, and thus LMTK1 seems to maintain somewhat low levels of Rab11A activity. This is consistent with our previous results showing that LMTK1−/− mouse brains have higher levels of active Rab11A (Takano et al., 2012). Indeed, premature outgrowth of dendrites in the LMTK1−/− brain was observed at early postnatal days, and thus LMTK1 could prevent dendritic hyperplasia by regulating membrane supply. LMTK1 function may therefore be important for establishing proper neuronal connections during brain development. It may be worth mentioning the current idea that premature or accelerated neuronal growth could be a cellular or molecular mechanism of autism spectrum disorder (Chomiak and Hu, 2013). It would be interesting to analyze the cytoarchitecture of autism-related brain regions, as well as behaviors, of LMTK1−/− mice.

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**FIGURE 5:** Dynamics of Rab11A-positive endosomes in axons of LMTK1−/− neurons. (A) Movement of Rab11A-positive endosomes in axons of LMTK1+/+ or LMTK1−/− neurons. Typical anterograde movement of a Rab11A-positive endosome is indicated with arrows. The cell body is on the left side. Bottom, kymographs of Rab11A-positive endosomes moving in axon. Bar, 5 μm. (B) Percentage of moving (light gray) or stationary (dark gray) Rab11A-positive endosomes at 3 DIV (3) and 7 DIV (7) in axons. (C) Percentage of Rab11A-positive vesicles moving anterogradely (gray), retrogradely (light gray), and bidirectionally (dark gray) in axons of LMTK1+/+ or LMTK1−/− neurons (n = 305 endosomes for LMTK1+/+ and n = 301 endosomes for LMTK1−/−). (D, E) Velocity of Rab11A-positive endosomes in axons. Relative frequency of Rab11A-positive endosomes moving at the indicated velocities in the anterograde (D) or retrograde (E) direction (n = 150 endosomes each for LMTK1+/+ and LMTK1−/−). (F, G) Run length of Rab11A-positive endosomes in LMTK1+/+ or LMTK1−/− neurons. Relative frequency of Rab11A-positive endosomes moving at the indicated run length in the anterograde (F) or retrograde (G) direction (n = 100 endosomes each for LMTK1+/+ and LMTK1−/−). Data are the mean ± SEM of three independent experiments (*p < 0.05, **p < 0.01, Student’s t test).
participate in the trafficking of synaptic vesicles in presynaptic regions, endocytic vesicles containing AMPA receptors in postsynaptic regions, and recycling endosomal vesicles in axons (Park et al., 2004; Brown et al., 2007; Ascano et al., 2009; Takano et al., 2012). Here we added dendritic growth and arborization as functions of Rab11A. The dynamics of Rab11-positive endosomes was only recently analyzed in detail in axons (Ascano et al., 2009; Eva et al., 2010), and no data are available for other regions of neurons. We have now analyzed trafficking of Rab11A-positive endosomes by real-time imaging in three regions of neurons, namely the cell body, dendritic shaft, and growth cone; in each region, LMTK1 down-regulation increased the dynamics of Rab11A-positive endosomes.

The movement of Rab11-positive endosomes in dendrites was compared with that in axons. This was particularly interesting because the polarity of microtubules, along which endosomal vesicles are transported, differs between axons and dendrites—all microtubules are plus-end distal in axons, and microtubules of opposite polarity run parallel to each other in dendrites (Baas et al., 1988; Stepanova et al., 2003). It is unclear which polarity of microtubules serves as the track for vesicle transport in dendrites. We attempted to determine the types of movements in dendrites by comparing to those in axons. Rab11-positive endosomes moved distally (anterogradely) and proximally (retrogradely) at a rate of 0.7–1.2 μm/s in dendrites, which was similar to movement in axons in this study and comparable to the rates of 0.4–1.2 μm/s in axons or neurites reported by others (Ascano et al., 2009; Eva et al., 2010; Power et al., 2012). The rate was also similar to that estimated with caRab11A-Q70L-positive endosomes. These results indicate that moving endosomes are mostly loaded with the active Rab11A-GTP, but the data do not provide insight into the polarity of microtubules used for dendritic transport. One of the main differences we observed was in the percentage of bidirectional movement, which was 32–34% in dendrites and 4–8% in axons. A large portion of the bidirectional movement in dendrites may reflect frequent exchange of Rab11A-positive endosomes between microtubules having opposite polarity. In addition, these results may suggest that Rab11A-positive endosomes use mainly plus-end-distal microtubules for vesicle transport even in dendrites, because LMTK1 knockout did not increase the prevalence of retrograde transport of Rab11A-positive endosomes in dendrites (Table 2).

The prevalence of anterograde movement of wtRab11A-positive endosomes in LMTK1+/− neurons increased in LMTK1−/− neurons to levels observed with caRab11A-Q70L-labeled vesicles in LMTK1+/− neurons. How does reduced LMTK1 activity increase the prevalence of anterograde movement? Anterograde movement in axons should be driven by the plus-end motor, kinesin. Rab11 effectors such as Rp11/FIP5, Rab11-FIP3, and protrudin interact with the kinesins KIF3 and KIF5 (Schonteich et al., 2008; Simon and Prekeris, 2008; Matsuzaaki et al., 2011). Down-regulation of LMTK1 up-regulates Rab11A activity (Takano et al., 2012), which in turn would be expected to increase the number of interactions between the Rab11A effectors and kinesin, leading to anterograde-biased movement. Another issue is how recycling endosomes travel longer distances in LMTK1−/− axons and dendrites. Up-regulation of Rab11A activity increases endosomal vesicle budding (Jedd et al., 1997; Takahashi et al., 2012), which would be expected to result in smaller endosomal vesicles. The caRab11A-Q70L-positive endosomes were indeed smaller than wtRab11A endosomes. We observed greater dynamics among smaller endosomes compared with larger endosomes. This was also the case in LMTK1−/− neurons. We observed frequent budding of Rab11A-positive endosomes at the perinuclear endosomal compartment, and those endosomes were smaller than those in LMTK1+/− neurons. Small endosomes may experience less resistance than larger endosomes when they pass through bundles of microtubules, similar to mitochondrial transport in axons (Shahpasand et al., 2012). Further, compared with larger vesicles, these smaller endosomal vesicles would have a greater density of active Rab11A on their surface, implying that the consequent greater

| Axons or dendrites | Mobile (%) | Immobile (%) | Anterograde movement (%) | Retrograde movement (%) | Bidirectional movement (%) | Velocity (μm/s) | Anterograde | Retrograde |
|-------------------|------------|--------------|--------------------------|-------------------------|---------------------------|-----------------|-------------|------------|
| LMTK1+/+a         |            |              |                          |                         |                           |                 |             |            |
| Axons (3 D)       | 71 ± 2.5   | 29 ± 2.5     | 51 ± 1.2                 | 45 ± 10.2               | 4 ± 0.1                   | 0.75 ± 0.03     | 0.71 ± 0.03 |
| Axons (7 D)       | 60 ± 3.4   | 40 ± 2.2     | 45 ± 0.5                 | 47 ± 10.5               | 8 ± 1.3                   | 0.85 ± 0.04     | 0.92 ± 0.03 |
| Dendrites         | 78 ± 1.8   | 22 ± 1.8     | 37 ± 1.2                 | 31 ± 1.7                | 32 ± 1.3                  | 0.94 ± 0.03     | 0.86 ± 0.03 |
| LMTK1−/−b         |            |              |                          |                         |                           |                 |             |            |
| Axons (3 D)       | 80 ± 3.4   | 20 ± 3.4     | 58 ± 10.9                | 40 ± 10.9               | 2 ± 0.3                   | 1.2 ± 0.02      | 1.11 ± 0.01 |
| Axons (7 D)       | 67 ± 1.3   | 33 ± 1.3     | 48 ± 0.1                 | 48 ± 0.3                | 4 ± 0.3                   | 1.18 ± 0.02     | 1.15 ± 0.03 |
| Dendrites         | 89 ± 10.2  | 11 ± 10.2    | 47 ± 0.5                 | 19 ± 0.8                | 34 ± 0.9                  | 1.24 ± 0.17     | 1.03 ± 0.05 |
| wtRab11A          | 81 ± 2.0   | 19 ± 2.0     | 41 ± 0.6                 | 33 ± 2.0                | 26 ± 1.4                  | 0.85 ± 0.04     | 0.85 ± 0.02 |
| caRab11A<sup>d</sup> | 85 ± 1.7   | 15 ± 1.7     | 49 ± 2.7                 | 35 ± 0.7                | 15 ± 2.4                  | 0.99 ± 0.02     | 0.96 ± 0.02 |
| dnRab11A<sup>e</sup> | 22 ± 4.2   | 78 ± 4.2     | ND                       | ND                      | ND                        | ND              |             |            |

TABLE 2: The movements of Rab11A-positive endosomes in axons or dendrites.
FIGURE 6: Movement of caRab11A-Q70L–positive endosomes in LMTK1<sup>+/+</sup> neurons and wtRab11A-positive endosomes in LMTK1<sup>−/−</sup> neurons. (A) A plasmid encoding EGFP-wtRab11A, EGFP-caRab11A-Q70L, or EGFP-dnRab11A-S25N was transfected into LMTK1<sup>+/+</sup> neurons, and movement of the Rab11A-positive endosomes was analyzed. Sequential frames over 10 s are shown in dendrites of LMTK1<sup>+/+</sup> neurons. Right, EGFP-Rab11A transfected into LMTK1<sup>−/−</sup> neurons. The cell body is to the left. The movement of typical Rab11A-positive endosomes is indicated with arrows. Bottom, kymographs of Rab11A-positive endosomes moving in dendrites. (B) Percentage of moving (light gray) or stationary (dark gray) Rab11A-positive endosomes in dendrites. (C) Percentage of Rab11A-positive endosomes moving anterogradely (gray), retrogradely (light gray), or bidirectionally (dark gray) in dendrites (n = 347 endosomes for LMTK1<sup>+/+</sup> neurons expressing wtRab11A, n = 349 endosomes for LMTK1<sup>+/+</sup> neurons expressing caRab11A-Q70L, n = 174 endosomes for LMTK1<sup>+/+</sup> neurons expressing dnRab11A-S25N, and n = 329 endosomes for LMTK1<sup>−/−</sup> neurons expressing wtRab11A). (D) The size of Rab11A-positive endosomes in primary dendrites. Relative frequency of Rab11A-positive endosomes of the indicated sizes. (E, F) The velocity of Rab11A-positive endosomes in primary dendrites.
FIGURE 7: Movement of Rab11A-positive endosomes in growth cones of LMTK1+/+ or LMTK1−/− neurons. (A) Growth cone of cortical neurons, which were prepared from LMTK1+/+ (left) or LMTK1−/− (right) mouse brains cotransfected with a plasmid encoding EGFP-Rab11A (top) or DsRed (bottom) at 6 DIV. Rab11A-positive endosomes were observed at 7 DIV. Bar, 2.5 μm. (B) The growth cone area of LMTK1+/+ or LMTK1−/− neurons (n = 30 each for LMTK1+/+ or LMTK1−/− neurons). (C) The size of Rab11A-positive recycling endosomes in growth cones of LMTK1+/+ or LMTK1−/− neurons (n = 30 endosomes each). (D) Percentage of mobile (light gray) or stationary (dark gray) Rab11A-positive endosomes in growth cones (n = 321 endosomes in LMTK1+/+ neurons and n = 250 endosomes in LMTK1−/− neurons). (E) Movement of Rab11A-positive endosomes in growth cones of LMTK1+/+ or LMTK1−/− neurons. Neurons prepared from LMTK1+/+ (top) or LMTK1−/− (bottom) mouse brains were cotransfected with a plasmid encoding EGFP-Rab11A or DsRed. Movements of Rab11A-positive endosomes were analyzed with real-time imaging. Sequential frames are shown over 84 s for LMTK1+/+ (top) and LMTK1−/− (bottom). The movement of a typical Rab11A-positive endosome is indicated with arrows. (F–H) The number of Rab11A-positive endosomes that were inserted into the plasma membrane at the tip of the growth cone (F), entered the growth cone from the dendritic shaft (G), or exited the growth cone (H) over 160 s (n = 165 endosomes in LMTK1+/+ neurons and n = 185 endosomes in LMTK1−/− neurons). Data are the mean ± SEM of three independent experiments (*p < 0.05; **p < 0.01; n.s., not significant, Student’s t test).
Cdk5-p35 phosphorylates LMTK1 at Ser-34 upstream of the kinase domain (Takano et al., 2010; Tsutsumi et al., 2010), and expression of the nonphosphorylatable mutant of LMTK1, S34A, led to increased Rab11A-positive endosome movement, axonal number of kinesin molecules on these small Rab11A-positive vesicles biases vesicle movement toward the anterograde direction. In this way, active Rab11A may sustain movement for a longer time, resulting in preferential anterograde movement.

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outgrowth, and dendrite growth and arborization. These results indicate that the Cdk5-LMTK1-Rab11A pathway operates during the process of supplying membrane materials to neurites. Indeed, Cdk5 has been shown to have a function in membrane transport. In axons of Caenorhabditis elegans, the trafficking of synaptic vesicles is stimulated in Cdk5 mutants (Ou et al., 2010; Goodwin et al., 2012), consistent with our results. Although there are no reports on Cdk5 function in endosomal transport in mammalian neurons, involvement of Cdk5 in dendrite formation has been documented. Layer V pyramidal neurons in cerebral cortex fail to develop a normal apical dendrite and instead extend multiple lateral dendrites in Cdk5−/− mouse brain (Ohshima et al., 2007). Knockdown or knock-out of Cdk5 in hippocampal neurons suppresses the brain-derived neurotrophic factor-induced increase in primary dendrites (Cheung et al., 2007). Cdk5 also affects dendrite formation by phosphorylating cytoskeletal proteins, different from that observed here through membrane traffic. Collapsin response mediator proteins (CRMPs) 1 and 2 are microtubule-associated proteins that are phosphorylated at Ser-522 by Cdk5 (Uchida et al., 2005). Double-mutant crmp1−/− (knock-out mutant of Ser522Ala) and crmp1−/− induce an irregular curling dendritic phenotype without changing the number and branching of primary dendrites (Yamashita et al., 2012). Cdk5 is a multifunctional protein kinase with action points in several signaling pathways, including neurotrophic factor–dependent signal transduction, cytoskeletal reorganization, and membrane trafficking (Dhavan and Tsai, 2001; Cheung and Ip, 2007; Hisanaga and Endo, 2010). Results different from the foregoing may be a consequence of differences in experimental systems. Of importance, axon and dendrite formation may be a result of integration of multiple Cdk5 actions.

MATERIALS AND METHODS
Antibodies
Anti-p35 (C-19), anti-Cdk5 (DC17), and monoclonal anti-myc (9E10) were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Anti-Rab11 was from Invitrogen (Carlsbad, CA). Monoclonal antineuron-specific class III β-tubulin was from Techno Corporation (Minneapolis, MN). Anti-actin was from Sigma-Aldrich (Saint Louis, MO). Anti-PSD95 was from Thermo Scientific (Manor Park Runcom, Cheshire, United Kingdom). Anti-GluR2 was from Chemicon (Temecula, CA). Anti-LMTK1 and anti-phospho-Ser34 of LMTK1 (pS34) have been described (Tsutsumi et al., 2008, 2010). Alexa-conjugated secondary antibodies were from Invitrogen.

Plasmid construction
EmGFP-miRNAs against LMTK1 have been described (Takano et al., 2012). The LMTK1 alanine (A) mutant and the phosphorylation-mimic aspartic acid (D) mutant at Ser-34 have been described (Takano et al., 2010; Tsutsumi et al., 2010). pEgFP-Rab5A, pEgFP-Rab7, pEgFP-Rab11A, pEgFP-Rab11A-Q70L, and pEgFP-Rab11A-S25N have been described (Fukuda, 2003; Fukuda et al., 2008).

Primary cortical neuron cultures and transfection
ICR mice were obtained from Sankyo Laboratory Service (Tokyo, Japan). LMTK1−/− mice have been described (Takano et al., 2012). Mice were housed in a temperature-controlled room under a 12-h light/12-h dark cycle with free access to food and water. All experiments were performed according to the guidelines for animal experimentation of Tokyo Metropolitan University. Brain cortices from embryonic day 17 or 18 mice were dissected and dissociated, and neurons were plated in polyethyleneimine (PE)-coated dishes or PE-coated glass coverslips at a density of 2.0 × 10^4 neurons/cm² (Saito et al., 2007). Plasmid vectors were introduced into neurons with electroporation using an Amaxa Nucleofector apparatus (Lonza, Tokyo, Japan; Takano et al., 2012).

Immunofluorescence staining and statistical analysis
Neurons were fixed with 4% paraformaldehyde in phosphate-buffered saline (PBS) for 20 min at room temperature and permeabilized in PBS containing 0.1% (wt/vol) Triton X-100 and 5% normal goat serum for 30 min. The cells were incubated with the indicated primary antibodies followed by secondary antibodies conjugated to Alexa 488, Alexa 546, or Alexa 647 (Invitrogen). Fluorescence images were acquired with an LSM 5 EXCITER confocal microscope or Zeiss LSM 710 confocal microscope (Carl Zeiss, Jena, Germany). The statistical significance of the data was evaluated with the Student’s t test, *p < 0.05 and **p < 0.01.

Analysis of Rab11A-positive endosome dynamics in neurons
Mouse brain cortical neurons were plated on a polyethyleneimine-coated glass-bottom dish (CELLview; Greiner, Frickenhuisen, Germany). Three or seven days after seeding, the culture medium was replaced with Neurobasal medium without phenol red and supplemented with B-27 (Invitrogen). The culture dish was mounted on a temperature-controlled stage on a Zeiss confocal microscope (LSM 710) at 37°C in 5% CO₂. Live images of vesicle movement were captured in dendrites or axons for at least 1 min with a 1-s intervals using a 63× oil-immersion lens. Rab11A-positive endosome movement was analyzed for the proportion of moving or stationary endosomes, the direction of movement, and speed. Rab11A-positive endosomes were classified as mobile or immobile according to criteria described by Eva et al. (2010). Briefly, Rab11A-positive endosomes were classified as immobile when their movement was <2 μm during observation for 100 s. Rab11A-positive endosomes moving in both directions but with net movement of <2 μm were classified as bidirectional. Rab11A-positive endosomes with net movement >2 μm in either direction were classified as anterograde or retrograde. Rab11A-positive endosomes were tracked for 100 s, and the maximum distance moved during three scans (3 s) was measured using Zen software and used to calculate the speed. The run length covered by Rab11A-positive endosomes was measured as a single continuous movement over 20 scans (20 s). Apparent sizes of Rab11A-positive endosomes were estimated with Zen software based on the area of EGFP fluorescence. Kymographs were made from serial frames of the same area using MetaMorph software (Molecular Devices, Sunnyvale, CA).

Preparation of cell extracts and immunoblotting
Neurons were disrupted in RIPA buffer (20 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1 mM ethylene glycol tetraacetic acid, 1 mM EDTA, 1% [wt/vol] Nonidet P-40, 0.1% SDS, 0.1% sodium deoxycholate, 0.4 mM Pefabloc, 10 μg/ml leupeptin, 10 mM NaF, 10 mM β-glycerophosphate, 1 mM Na₂VO₄). After centrifugation at 10,000 × g for 20 min at 4°C, the extract was suspended in SDS sample buffer and boiled for 5 min. SDS-PAGE and immunoblotting were performed as described (Takano et al., 2010, 2012).

Golgi staining
LMTK1−/− or LMTK1−/− brain at postnatal day 7 or 3 mo of age was stained with reagents in an FD Rapid GolgiStain kit (FD Neurotechnologies, Columbia, MD).
ACKNOWLEDGMENTS

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