

# Synaptic removal of diacylglycerol by DGKζ and PSD-95 regulates dendritic spine maintenance

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Diacylglycerol (DAG) is an important lipid signalling molecule that exerts an effect on various effector proteins including protein kinase C. A main mechanism for DAG removal is to convert it to phosphatidic acid (PA) by DAG kinases (DGKs). However, it is not well understood how DGKs are targeted to specific subcellular sites and tightly regulates DAG levels. The neuronal synapse is a prominent site of DAG production. Here, we show that DGK<sup>\(\zeta\)</sup> is targeted to excitatory synapses through its direct interaction with the postsynaptic PDZ scaffold PSD-95. Overexpression of DGKζ in cultured neurons increases the number of dendritic spines, which receive the majority of excitatory synaptic inputs, in a manner requiring its catalytic activity and PSD-95 binding. Conversely, DGKC knockdown reduces spine density. Mice deficient in DGK expression show reduced spine density and excitatory synaptic transmission. Time-lapse imaging indicates that DGK is required for spine maintenance but not formation. We propose that PSD-95 targets DGK<sub>2</sub> to synaptic DAG-producing receptors to tightly couple synaptic DAG production to its conversion to PA for the maintenance of spine density. The EMBO Journal (2009) 28, 1170-1179. doi:10.1038/ emboj.2009.44; Published online 19 February 2009 Subject Categories: signal transduction; neuroscience Keywords: DGKζ; diacylglycerol kinase; phosphatidic acid; PSD-95; spine

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# Introduction

Activation of various surface receptors including G proteincoupled receptors (GPCRs) and receptor tyrosine kinases leads to the hydrolysis of phosphatidylinositol 4,5-bisphosphate (PIP<sub>2</sub>) (Sternweis *et al*, 1992; Rhee, 2001). PIP<sub>2</sub> hydrolysis is mediated by phospholipase C (PLC) enzymes and produces diacylglycerol (DAG) and inositol 1,4,5-triphosphate (IP<sub>3</sub>). Although IP<sub>3</sub> induces Ca<sup>2+</sup> release from intracellular stores, DAG binds and stimulates various C1 domain-containing effector proteins including protein kinase C (PKC).

The generation and removal of DAG at various subcellular sites occur rapidly (within seconds), indicating a tight regulation of signal termination. The main mechanism for the termination of DAG signalling is to phosphorylate DAG and convert it to phosphatidic acid (PA) by diacylglycerol kinases (DGKs), enzymes conserved in most eukaryotic species (Topham, 2006; Sakane *et al*, 2007). However, it is not well understood how DGKs are targeted to subcellular sites and involved in the tight regulation of local DAG levels.

The excitatory neuronal synapse is a prominent site of receptor activation and signal transduction. Production of DAG at synapses occurs through the activation of PLC-coupled GPCRs, including metabotropic glutamate receptors (mGluRs) (Conn and Pin, 1997). Strong synaptic PLC activation is also triggered by non-GPCRs such as NMDA receptors (NMDARs), a key mediator of excitatory synaptic transmission (Reyes-Harde and Stanton, 1998; Choi *et al*, 2005; Horne and Dell'Acqua, 2007).

Excitatory synapses are mainly localized in spines, which are actin-rich tiny protrusions on neuronal dendrites that exhibit structural and functional interplay with excitatory synaptic transmission (Nimchinsky et al, 2002; Tsay and Yuste, 2004; Carlisle and Kennedy, 2005; Ethell and Pasquale, 2005; Govek et al, 2005; Hayashi and Majewska, 2005; Matus, 2005; Calabrese et al, 2006; Tada and Sheng, 2006; Alvarez and Sabatini, 2007; Schubert and Dotti, 2007; Sekino et al, 2007; Bourne and Harris, 2008; Cingolani and Goda, 2008). Because dendritic spines are spatially isolated from dendrites through a narrow neck and have a relatively small volume compared with dendrites, the removal of synaptic signalling molecules may be a key mechanism determining the intensity and time course of postsynaptic responses. In this context, although the mechanisms underlying synaptic Ca<sup>2+</sup> removal are relatively well understood (Sabatini et al, 2001), little is known about how synaptic DAG is efficiently removed.

PSD-95 is an abundant scaffolding protein at excitatory synapses implicated in the regulation of synapse formation, maturation, and structural and functional plasticity (Kennedy, 2000; Scannevin and Huganir, 2000; Funke *et al*, 2004; Elias and Nicoll, 2007; Sheng and Hoogenraad, 2007). These PSD-95 functions are thought to be mediated in part by its interaction with various synaptic signalling proteins and

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organization of efficient signalling pathways. For instance, PSD-95 simultaneously binds NMDARs and neuronal nitric oxide synthase to promote the functional coupling between activated receptors and downstream effectors (Brenman *et al*, 1996; Sattler *et al*, 1999). However, involvement of PSD-95 in the termination of synaptic signalling, which is as important as signal activation, is not well understood.

Here, we present data supporting that PSD-95 promotes synaptic recruitment of DGK $\zeta$ , a DGK isoform enriched at excitatory synapses. Spine density in cultured neurons is regulated by overexpression and knockdown of DGK $\zeta$  in a manner requiring the catalytic activity and PSD-95 binding. Moreover, deficiency of DGK $\zeta$  expression in mice leads to reductions in spine density and excitatory synaptic transmission. These results suggest that the conversion of synaptic DAG to PA by the concerted action of DGK $\zeta$  and PSD-95 is important for the maintenance of normal synaptic density and transmission.

## Results

# *PSD-95 interacts with and promotes synaptic localization of DGK*ζ

To better understand the PSD-95-dependent molecular organization of excitatory synapses, we searched for PSD-95-binding proteins by yeast two-hybrid screen. The screen identified DGK $\zeta$ , 1 of the 10 mammalian DGKs (type IV) with a unique C-terminal PDZ domain-binding motif (Figure 1A) (Goto and Kondo, 1996). Additional two-hybrid assays confirmed a specific interaction between the DGK $\zeta$  C terminus and PDZ domains of PSD-95 family proteins (PSD-95, SAP97, PSD-93/chapsyn-110, and SAP102) (Supplementary Figure S1A). DGK $\zeta$  did not interact with other PDZ domains from Shank1, GRIP2, PICK1, and nNOS, suggesting that DGK $\zeta$  and PSD-95 family proteins. DGK $\zeta$  and rat brain (Supplementary Figure S1B; Figure 1B and C).



Figure 1 PSD-95 interacts with and promotes synaptic localization of DGKζ. (A) Domain structure of DGKζ. C1, protein kinase C conserved region 1; MARCKS, myristoylated alanine-rich C-kinase substrate domain; PB, PDZ domain-binding motif. (B, C) Association of DGK<sup>L</sup> with PSD-95 family proteins in vivo. Adult rat brain lysates, solubilized by 1% sodium deoxycholate, were immunoprecipitated with DGKÇ antibodies (1519) and immunoblotted with the indicated antibodies. Input, 10%; IP, immunoprecipitation. (D) Widespread distribution of DGKζ in various brain regions. α-tubulin was used for normalization. (E) DGKζ expression increases during postnatal rat brain development. E, embryonic day; P, postnatal day. PSD-95 was visualized for comparison. (F) Distribution patterns of DGK in subcellular fractions of adult rat brain. H, homogenates; P2, crude synaptosomes; S2, supernatant after P2 precipitation; S3, cytosol; P3, light membranes; LP1, synaptosomal membranes; LS2, synaptosomal cytosol; LP2, synaptic vesicle-enriched fraction. PSD-95 and synaptophysin were visualized for comparison. (G) Enrichment of DGK (in postsynaptic density (PSD) fractions; extracted with Triton X-100 once (PSD I), twice (PSD II), or with Triton X-100 and Sarkosyl (PSD III). (H-J) Wild-type (WT) DGK shows a greater spine localization compared with a mutant DGK that lacks the PSD-95binding C terminus ( $\Delta$ C). Cultured hippocampal neurons transfected with HA-DGK $\zeta$  constructs and PSD-95-EGFP (days *in vitro*, DIV 17–18) were stained with HA and EGFP antibodies and used to measure the ratio of DGKζ signals at a spine versus an adjacent dendritic trunk. PSD-95-EGFP was co-transfected with DGK $\zeta$  so that the amount of PSD-95 does not become a limiting factor. Data represent mean ± s.e.m. (WT, n = 18neurons;  $\Delta C$ , n = 10 neurons). \*\*\*P < 0.001 (Student's *t*-test). Scale bar, 10  $\mu$ m. (K-M) Knockdown of PSD-95 reduces spine localization of DGKζ. Cultured neurons were doubly transfected with HA-DGKζ and a PSD-95 knockdown construct (sh-PSD-95), or HA-DGKζ and empty sh-RNA vector and (sh-vec; DIV 14-20), and the spine/dendrite ratio of DGK<sup>z</sup> signals were measured. Data represent mean ± s.e.m. (sh-vec, n = 13 neurons; sh-PSD-95, n = 19 neurons). \*\*\*P < 0.001 (Student's *t*-test). Scale bar, 10 µm.

Before we explore the functions of the interaction between PSD-95 and DGKζ, we first determined the expression patterns of DGK<sup>2</sup> proteins in rat brain, using a DGK<sup>2</sup>-specific polyclonal antibody that we generated (Supplementary Figure S2). In immunoblot analysis, DGK<sup>C</sup> protein expression was detected in various rat brain regions (Figure 1D). Consistently, DGKζ immunofluorescence was observed in various brain regions, including hippocampus and cerebellum (Supplementary Figure S3). Minimal signals were detected in brain slices from DGKζ-deficient mice (Supplementary Figure S3), indicating the specificity of the antibody. DGKζ expression gradually increased during postnatal rat brain development (Figure 1E). In subcellular fractions of the rat brain, DGKζ was detected in synaptic fractions, including crude synaptosomal, synaptic membrane, and synaptic vesicle fractions (Figure 1F). Notably, DGKζ was highly enriched in postsynaptic density (PSD; electron-dense postsynaptic multiprotein complexes) fractions, to an extent comparable to PSD-95 (Figure 1G), indicating that DGK is tightly associated with the PSD.

The subcellular fractionation patterns of PSD-95 and DGK $\zeta$ (Figure 1F) indicate that PSD-95 is mainly localized at synaptic fractions, whereas DGK $\zeta$  is not. We thus reasoned that DGK $\zeta$  may depend on PSD-95 interaction for its synaptic localization. Indeed, wild-type (WT) DGK $\zeta$  expressed in cultured hippocampal neurons showed a greater localization at dendritic spines, compared with a mutant that lacks PSD-95 binding (DGK $\zeta\Delta$ C) (Figure 1H–J), suggesting that PSD-95 promotes synaptic localization of DGK $\zeta$ . Of note, some of DGK $\zeta\Delta$ C signals colocalized with PSD-95 in dendrites (Figure 1I). This is most likely to be caused by the association of DGK $\zeta\Delta$ C with endogenous DGK $\zeta$ , which is capable of binding to PSD-95, rather than a C terminus-independent DGK $\zeta$  binding to PSD-95 (Supplementary Figure S4A). In support of this possibility, DGK $\zeta$  could form self-multimers, as shown by the co-immunoprecipitation of differentially tagged DGK $\zeta$  proteins (Supplementary Figure S4B). In further support of the role for PSD-95 in synaptic DGK $\zeta$  localization, knockdown of PSD-95 significantly decreased spine localization of DGK $\zeta$  (Figure 1K–M; Supplementary Figure S5A). DGK $\zeta$  has also been shown to interact with another PDZ protein,  $\gamma$ 1-syntrophin (Hogan *et al*, 2001). However, the role for  $\gamma$ 1-syntrophin is synaptic DGK $\zeta$  localization is unclear because  $\gamma$ 1-syntrophin is detected mainly in the endoplasmic reticulum of neuronal cell bodies (Alessi *et al*, 2006).

# DGK $\zeta$ is detected mainly in dendrites and postsynaptic sites

We next determined the subcellular and ultrastructural localization of DGK $\zeta$ . In cultured hippocampal neurons, DGK $\zeta$ was detected in MAP2 (a dendritic marker)-positive dendrites, although relative weak signals were also detected in MAP2-negative axons (Figure 2A). In dendrites, DGK $\zeta$ showed a widespread subcellular distribution, with some of the DGK $\zeta$  signals being colocalized with PSD-95 (an excitatory postsynaptic marker) and synapsin I (a presynaptic marker) (Figure 2B and C).

Ultrastructural localizations of DGK $\zeta$  were determined by electron microscopic analysis. DGK $\zeta$  was detected mainly in the cell body and proximal dendrites (Figure 2D) as well as distal dendrites (Figure 2E). In contrast, DGK $\zeta$  was rarely detected in axonal compartments (Figure 2F). The majority of dendritic spines contained DGK $\zeta$  signals in regions including the PSD, whereas axon terminals in contact with dendritic spines were usually devoid of detectable DGK $\zeta$  signals



**Figure 2** Subcellular and ultrastructural localization of DGK $\zeta$  in neurons. (**A**) DGK $\zeta$  localizes to MAP2 (a dendritic marker protein)-positive dendrites as well as MAP2-negative axons (arrowheads). Cultured hippocampal neurons (DIV 21) were stained with DGK $\zeta$  (1521) and MAP2 antibodies. Scale bar, 50 µm. (**B**, **C**) DGK $\zeta$  colocalizes with PSD-95 (an excitatory postsynaptic marker) and synapsin I (a presynaptic marker). Note that DGK $\zeta$  shows widespread distribution patterns including the dendritic trunk and PSD-95/synapsin I-positive synaptic sites. Scale bar, 10 µm. (**D**–**G**) Ultrastructural localization of DGK $\zeta$  in somatodendritic regions and dendritic spines of CA1 pyramidal neurons. Note that DGK $\zeta$  signals, shown as dark DAB precipitates (arrowheads), are present mainly in the cell body and proximal dendrites (D) as well as distal dendrites (E; indicated by an arrow), whereas axons rarely contain DGK $\zeta$  signals (F; an arrow indicates an axon terminal). In addition, DGK $\zeta$  is detected in the majority of dendritic spines in regions including the PSD, whereas axon terminals in contact with dendritic spines rarely contain DGK $\zeta$  signals (G; an arrow indicates a spine synapse). Scale bars; 2 µm (D), 1 µm (E), and 0.2 µm (F, G).

(Figure 2G). These results suggest that DGK $\zeta$  is present mainly in postsynaptic compartments.

# DGK<sup>C</sup>-dependent spine regulation requires its catalytic activity and PSD-95 binding

Production of DAG at neuronal synapses can occur through the activation of PLC-coupled GPCRs, including mGluRs (Conn and Pin, 1997) and non-GPCRs such as NMDARs, a key mediator of excitatory synaptic transmission (Reves-Harde and Stanton, 1998; Choi et al, 2005; Horne and Dell'Acqua, 2007). Synaptic proteins that exert an effect in the downstream of DAG such as α1-chimaerin and MARCKS have been implicated in the regulation of dendritic spines (Calabrese and Halpain, 2005; Buttery et al, 2006), which are actin-rich tiny protrusions on neuronal dendrites (Calabrese et al, 2006; Tada and Sheng, 2006; Bourne and Harris, 2008). In addition, PSD-95, which is in direct association with DGKC, is a key regulator of spine morphogenesis (Tada and Sheng, 2006). We thus tested whether DGK $\zeta$  is involved in the regulation of dendritic spines. DGKζ overexpression in cultured hippocampal neurons (days in vitro or DIV 15-22) significantly increased the density, but not length and width, of dendritic spines (Figure 3A-E). In contrast, DGK mutants ( $\Delta C$  and kinase-dead) that lack PSD-95 binding and kinase activity, respectively, had no effect on spine density (Figure 3A-E). These results suggest that DGK<sup>\(\zeta\)</sup> increases spine density through mechanisms requiring its synaptic localization and catalytic activity.

Because DGK $\zeta$  overexpression increases spine density, we reasoned that a reduced expression of DGK $\zeta$  may negatively regulate spine density. Indeed, cultured neurons with DGK $\zeta$  knocked down by RNAi (DIV 15–19) displayed a significantly reduced spine density, compared with controls (Figure 4A and B; Supplementary Figures S5B and S5C). Spine length and width, however, were unaffected (Figure 4C and D). The reduced spine density could be rescued by co-transfection of a WT DGK $\zeta$  expression construct resistant to sh-DGK $\zeta$ , but not by a kinase-dead construct (Figure 4A–D; Supplementary Figure S5D). DGK $\zeta$  knockdown also reduced the number of

PSD-95-positive dendritic spines (Figure 4E and F). However, DGK $\zeta$  knockdown did not reduce spine localization of PSD-95 (Figure 4G), suggesting that DGK $\zeta$  knockdown does not affect synaptic PSD-95 localization. These results suggest that DGK $\zeta$  is important for the maintenance of dendritic spines in a manner requiring its catalytic activity.

# Reduced PA production and spine density in $\textit{DGK}\zeta-/-$ mice

To determine whether the results obtained from cultured neurons have relevance in vivo, we analysed DGK(-/mice, which exhibit enhanced T-cell responses caused by reduced DAG conversion to PA (Zhong et al, 2003). We observed a similar reduction in PA production in DGK $\zeta$ -/hippocampal slices, when synaptic DAG production was stimulated with DHPG, a group I mGluR agonist (Figure 5A). In contrast, basal PA levels in DGK<sup>2</sup>/- slices were not significantly different from WT mice, although there was a tendency for a decrease (Figure 5A). In addition to measuring PA production, we compared expression levels of DGK proteins and other synaptic proteins in the whole brain of DGK $\zeta$ -/- and WT mice. Although DGK $\zeta$  expression could not be detected in DGK $\zeta$ -/- brain, expression levels of other synaptic proteins were unchanged (Supplementary Figure S6).

For spine analysis, hippocampal CA1 pyramidal neurons were visualized by ballistic delivery of DiI-coated particles (Gan *et al*, 2000). Consistent with the results from cultured neurons, spine density, but not spine head area, was significantly reduced ( $\sim$ 30%) in DGK $\zeta$ -/- pyramidal neurons, compared with WT controls (Figure 5B–D), suggesting that DGK $\zeta$  is required for the maintenance of dendritic spines *in vivo*.

# Reduced excitatory synaptic transmission in $DGK\zeta$ -/- mice

We next tested whether the reduction of spine density in DGK $\zeta$ -/- mice is accompanied by functional reduction in excitatory synaptic transmission. The frequency, but not



**Figure 3** Overexpression of DGK $\zeta$  increases spine density in cultured neurons. (**A**) Domain structure of DGK $\zeta$  variants. KD, kinase-dead mutant. (**B**) Increased spine density by overexpression of WT DGK $\zeta$ , but not by  $\Delta$ C and KD mutants. Cultured hippocampal neurons transfected with HA-DGK $\zeta$  (WT and mutants) and EGFP, or EGFP alone (control; DIV 15–22), were stained with EGFP antibodies. Scale bar, 10 µm. (**C**–**E**) Quantification of spine density, length, and width from the results in (B). Data represent mean ± s.e.m. (Control, 5.06 ± 0.28 spines per 10 µm dendrite. n = 37; WT, 7.27 ± 0.26, n = 31;  $\Delta$ C, 5.33 ± 0.45, n = 22; KD, 4.46 ± 0.28, n = 16). \*\*\*P<0.001 (Student's *t*-test).

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**Figure 4** Knockdown of DGK $\zeta$  decreases spine density. (**A**) Reduced spine density by DGK $\zeta$  knockdown, and rescue of the effect requiring catalytic activity. Neurons were transfected with an shRNA DGK $\zeta$  knockdown construct (sh-DGK $\zeta$ ), or vector alone (sh-vec; DIV 15–19), and stained with EGFP antibodies for spine visualization. Rescue expression constructs were doubly transfected with sh-DGK $\zeta$ . Scale bar, 10 µm. (**B**–**D**) Quantification of the results in (A). Mean ± s.e.m. (sh-vec, 4.74 ± 0.26, *n* = 19; sh-DGK $\zeta$ , 2.56 ± 0.21, *n* = 18; WT rescue, 4.67 ± 0.42, *n* = 8; KD rescue, 2.36 ± 0.19, *n* = 26). \*\*\**P*<0.001 (Student's *t*-test). (**E**) DGK $\zeta$  knockdown reduces the number of PSD-95-positive dendritic spines. Neurons were transfected with sh-DGK $\zeta$  or sh-vec (DIV 15–19), and stained for PSD-95 and EGFP. Scale bar, 10 µm. (**F**) Quantification of the results in (E). Mean ± s.e.m. (sh-vec, 4.27 ± 0.31, *n* = 24; sh-DGK $\zeta$ , 1.42 ± 0.2, *n* = 25). \*\*\**P*<0.001 (Student's *t*-test). (**G**) Quantification of the spine/dendrite ratio of PSD-95 from the results in (E). Mean ± s.e.m. (sh-vec, 4.88 ± 0.89, *n* = 16; sh-DGK $\zeta$ , 4.19 ± 0.68, *n* = 16).



**Figure 5** Reduced PA production and spine density in DGK $\zeta$ -/mice. (**A**) Reduced PA production in DGK $\zeta$ -/- hippocampus. PA production was induced by the treatment of DHPG (a group I mGluR agonist) on WT and DGK $\zeta$ -/- hippocampal slices. Data represent mean±s.e.m. (basal-WT, 2.57±0.48, *n*=3; basal-KO, 1.05±0.25, *n*=3; DHPG-WT, 7.06±0.28, *n*=3; DHPG-KO, 3.96±0.44, *n*=3). \**P*<0.05 (Student's *t*-test). (**B**) Reduced spine density in DGK $\zeta$ -/- CA1 pyramidal neurons. Spines in WT and DGK $\zeta$ -/- pyramidal neurons were labelled by ballistic delivery of Dil-coated particles to hippocampal slices (3 weeks old). Scale bar, 10 µm. (**C**, **D**) Quantification of the results from (B) reveals a reduction in spine density, but not in spine head area, in DGK $\zeta$ -/neurons. Mean±s.e.m. (WT, 12.41±0.87 spines per 10 µm dendrite, *n*=27 neurons from three mice; KO, 8.79±0.51, *n*=30, three mice). \*\*\**P*<0.01 (Student's *t*-test).

amplitude, of miniature excitatory postsynaptic currents (mEPSCs) was reduced ( $\sim$ 44%) in DGK $\zeta$ -/- CA1 pyramidal neurons (Figure 6A and B), suggesting that DGK is required for the maintenance of AMPA receptor-mediated spontaneous excitatory synaptic transmission. Miniature inhibitory postsynaptic currents (mIPSCs), however, were unaffected (Figure 6C and D). The slopes of AMPA receptor-mediated evoked fEPSPs plotted against the stimulation intensity were reduced at DGKζ-/- Schaffer collateral-CA1 pyramidal synapses (Figure 6E), consistent with the reduction in mEPSC frequencies. The paired-pulse facilitation (PPF) ratio was not changed at DGKζ-/- Schaffer collateral-CA1 pyramidal synapses (Figure 6F), suggesting that the reductions in AMPA receptor-mediated spontaneous and evoked responses were not caused by a reduction in presynaptic release probability. The ratio of AMPA receptor and NMDAR-mediated evoked EPSCs in DGKζ-/- CA1 pyramidal neurons was comparable to that of WT mice (Figure 6G and H), suggesting that DGK<sup>\(\)</sup> deficiency did not cause a change in the AMPA/ NMDA ratio of excitatory transmission in synapses that escaped the reduction in number.

### DGKζ regulates the maintenance but not formation of dendritic spines

The reduced spine density induced by DGK $\zeta$  knockdown may be caused by either reduced spine formation and/or increased spine elimination. To address this issue, we monitored the dynamics of dendritic spines in live cultured neurons transfected with the DGK $\zeta$  knockdown construct. DGK $\zeta$  knockdown markedly increased the number of disappearing dendritic spines during the period of 2-h live imaging, compared with control neurons transfected with the empty knockdown vector (Figure 7A and B). Accordingly, the num-



**Figure 6** Reduced excitatory synaptic transmission in DGK $\zeta$ -/- mice. (**A**, **B**) Reduced frequency, but not amplitude, of mEPSCs in DGK $\zeta$ -/- CA1 pyramidal neurons. Mean ± s.e.m. (WT, 0.17 ± 0.02 Hz, n = 27 cells from seven mice; KO, 0.09 ± 0.01 Hz, n = 26, four mice). \*\*P<0.01 (Student's *t*-test). (**C**, **D**) Normal frequency and amplitude of mIPSCs in DGK $\zeta$ -/- CA1 neurons. (WT, n = 10 cells from three mice; KO, n = 9, three mice). (**E**) Reduced AMPA receptor-mediated evoked synaptic transmission at DGK $\zeta$ -/- Schaffer collateral-CA1 pyramidal synapses. The synaptic input-output relationship was obtained by plotting fEPSP slopes against stimulation intensities. n = 27 slices from 10 KO mice and 31 slices from 11 WT mice. Scale bar, 4 ms/0.4 mV. (**F**) Normal paired-pulse facilitation ratio at DGK $\zeta$ -/- Schaffer collateral-CA1 pyramidal synapses. Facilitation ratios were plotted against interstimulus intervals.  $n = 38 \sim 39$  slices from 13  $\sim$ 14 mice. Scale bar, 10 ms/0.5 mV. (**G**, **H**) Normal ratio of AMPA and NMDA receptor-mediated EPSCs. Mean ± s.e.m. (WT, n = 11 cells from seven mice; KO, n = 9, seven mice). AMPA and NMDA receptor-mediated responses were recorded under voltage clamp with a holding potential of -70 and +40 mV, respectively.



**Figure 7** DGK $\zeta$  regulates the maintenance but not formation of dendritic spines. (**A**) DGK $\zeta$  knockdown markedly increases the number of disappearing spines. Cultured neurons were transfected with sh-DGK $\zeta$  or control vector (sh-vec), at DIV 13, followed by time-lapse imaging of EGFP-positive live neurons at DIV 17 for 2h. Open arrowheads, disappearing spines; filled arrowheads, appearing spines; arrows: spines remained after the 2-h imaging. Scale bar, 10 µm. (**B**) Quantification of the results in (A). Percentages of remaining, appearing, and disappearing protrusions are indicated (the sum of remained, appeared, and disappeared spines was used as the total number of spines). Mean ± s.e.m. sh-vec (remain, 84.6±4.16; appear, 7.25±2.86; disappear, 8.14±2.38; *n*=264 spines from five neurons); sh-DGK $\zeta$  (remain, 45.14±5.57; appear, 1.95±1.22; disappear, 52.9±6.35; *n*=204 spines from six neurons). \*\*\**P*<0.001 (Student's *t*-test).

ber of remaining dendritic spines after 2-h imaging was significantly decreased in DGKζ knockdown neurons (Figure 7A and B). In contrast, the number of appearing spines was not significantly decreased in DGKζ knockdown neurons, although there was a tendency for a decrease (Figure 7A and B). Because the overall number of disappearing spines is much greater than that of appearing spines in DGK $\zeta$  knockdown neurons, DGK $\zeta$  knockdown seems to reduce spine density mainly through spine elimination.

### PKC inhibition does not block DGKζ knockdowninduced spine density reduction

Decreased expression of DGK $\zeta$  in cultured neurons may lead to both an increase in DAG levels and a decrease in PA levels.

A well-known downstream effector of DAG is PKC. In addition, MARCKS, a substrate of PKC, is required for the spine shrinkage induced by phorbol ester (a PKC activator) in cultured neurons (Calabrese et al, 2006). These results suggest that DGK<sup>\(\)</sup> knockdown may reduce spine density through the PKC signalling pathway. To this end, we tested whether PKC inhibition reduces the spine reduction induced by DGK knockdown. However, incubation of cultured neurons with Gö6976, a specific inhibitor of PKC, during the period of DGKζ knockdown (DIV 15-19) did not significantly reduce the effect of DGK<sup>(</sup> knockdown (Figure 8A and B). The effectiveness of Gö6976 as a PKC inhibitor was verified in heterologous cells and neurons stimulated with PMA (a PKC activator) (Supplementary Figure S7). These results indicate that PKC is minimally involved in DGK knockdown-induced reduction in spine density.

# Discussion

Our results suggest that PSD-95 recruits DGK $\zeta$  to excitatory synapses to tightly control synaptic DAG conversion to PA. Given the direct interaction of PSD-95 with NMDARs, a strong stimulator of PLC activation and DAG production (Horne and Dell'Acqua, 2007), PSD-95 may target DGK $\zeta$  close to NMDARs to tightly couple DAG production to its conversion to PA. Consistently, a recent study has demonstrated that DGKs are targeted to GPCRs by their interactions with  $\beta$ -arrestin, a signalling adaptor/scaffolding protein (Nelson *et al*, 2007).

PSD-95 is known to be linked to other proteins in the DAG signalling pathway in the brain. Shank, a synaptic scaffold linked to PSD-95 through GKAP, interacts directly with PLC $\beta$ and indirectly with mGluRs (a stimulator of PLC $\beta$ ) (Tu *et al*, 1999; Hwang et al, 2005). These interactions along with our results suggest that PSD-95 may function as a scaffold to organize the synaptic DAG signalling pathway into a macromolecular complex, which contains proteins involved in PLC stimulation (NMDARs and mGluRs), DAG generation (PLC $\beta$ ), and DAG-to-PA conversion (DGKζ). This is reminiscent of the INAD (a multi-PDZ protein)-dependent protein complex in the Drosophila phototransduction pathway that contains PLC, TRP channels (a DAG effector), and ePKC (a negative regulator of TRP channel), and which is thought to promote the high sensitivity, fast activation/deactivation, and feedback modulation of the signalling pathway (Montell, 1999).



**Figure 8** PKC inhibition does not block DGK $\zeta$  knockdown-induced spine density reduction. (A) PKC inhibition does not affect DGK $\zeta$  knockdown-induced spine reduction. Cultured neurons transfected with sh-DGK $\zeta$  or sh-vec (DIV 15–19), were treated with Gö6976 (200 nM) or vehicle (DMSO; DIV 15–19). (B) Quantification of the results in (B). Mean ± s.e.m. (sh-vec/vehicle, 4.72 ± 0.18, *n* = 39; sh-DGK $\zeta$ /vehicle, 2.71 ± 0.17, *n* = 34; sh-vec/Gö6976, 4.62 ± 0.16, *n* = 60; sh-DGK $\zeta$ /Gö6976, 3.32 ± 0.33, *n* = 33).

Such examples, however, have not been identified in higher organisms.

Our data indicate that DGK<sup>\(\zeta\)</sup> forms a complex with all four known members of the PSD-95 family in brain (Figure 1), consistent with the in vitro results (Supplementary Figure S1). PSD-95 family proteins have been suggested to have overlapping as well as distinct functions with regard to their spatiotemporal expression patterns, protein interactions, and regulation of synaptic transmission (Sans et al, 2000, 2001; Valtschanoff et al, 2000; Rumbaugh et al, 2003; Townsend et al, 2003; Kim and Sheng, 2004; Elias et al, 2006; Fitzjohn et al, 2006; Schluter et al, 2006; Elias and Nicoll, 2007). For instance, targeted truncation of PSD-95 in mice does not affect AMPA receptor-mediated synaptic transmission (Migaud et al, 1998), most likely due to the compensation by other PSD-95 family members. Therefore, the promiscuous interaction of DGKζ with PSD-95 family proteins may contribute to the stable maintenance of dendritic spines and excitatory transmission.

Our results indicate that DGK $\zeta$  is detected mainly in dendrites and postsynaptic sites, consistent with the DGK $\zeta$ -dependent spine regulation. However, a small portion of DGK $\zeta$  proteins is also detected in axon terminals, consistent with the localization of DGK $\zeta$  in the LP2 (synaptic vesicleenriched) fraction. Therefore, we cannot exclude the possibility that the reductions in spine density and synaptic transmission at DGK $\zeta$ -/- synapses are caused, at least in part, by presynaptic defects, although the presynaptic release probability, as measured by PPF, appears normal in the knockout animals.

Our data from cultured neurons and DGK<sup>L</sup>/- mice suggest that DGKζ-mediated synaptic conversion of DAG to PA is required for the maintenance of dendritic spines. The reduced spine density may result from enhanced actions of DAG on synaptic effectors such as PKC. However, inhibition of PKC during DGKC knockdown did not significantly reduce the effect of DGK<sup>\z</sup> knockdown. This suggests that PKC may not be the major pathway mediating the spine-reducing effect of DGK knockdown. However, it is possible that, even in the absence of the influences of PKC pathway, other DAG pathways may still be sufficient to induce spine reduction under the condition of elevated DAG levels. A possible non-PKC DAG effector is α1-chimaerin, a Rac1 GTPase-activating protein known to mediate DAG-dependent spine pruning (Buttery et al, 2006). Alternatively, because DGK deficiency leads to a decrease in PA production (Figure 5), insufficient activation of downstream PA effectors may inhibit spine maintenance. Previous studies have identified a large number of PA effectors, including Rac1, PAK1, PIP5K (phosphatidylinositol-4-phosphate 5-kinase), RasGAP, sphingosine kinase, atypical PKC, Raf-1, PKC<sub>E</sub>, PP1c (the catalytic subunit of protein phosphatase-1), mTOR, Opi1p, p47<sup>phox</sup>, n-chimaerin, and B2-chimaerin. Of these proteins, Rac1 and PAK1 have been strongly implicated in the regulation of dendritic spines and F-actin, the main cytoskeleton of dendritic spines (Calabrese et al, 2006; Tada and Sheng, 2006; Bourne and Harris, 2008). Therefore, our study suggests the PA effectors listed above as potential regulators of dendritic spines.

In conclusion, our results suggest that synaptic DAG-to-PA conversion promoted by PSD-95-dependent synaptic targeting of DGK $\zeta$  tightly regulates local DAG levels for the maintenance of normal spine density and excitatory

synaptic transmission. Our study suggests a novel mechanism by which local DAG concentration can be efficiently regulated by scaffolding protein-dependent localization of DAG-converting enzymes to the vicinity of DAG-generating proteins, which may be employed by various other subcellular sites.

### Materials and methods

#### Constructs

The PDZ2 domain of PSD-95 (aa 89–299) in pBHA was used to screen a human brain cDNA library in pACT2 (Clontech). DGK $\zeta$  (aa 923–929), wild type, and mutant (V929A) were annealed into pBHA. Rat DGK $\zeta$  expression constructs (HA-DGK $\zeta$  WT and  $\Delta$ C; WT, full length;  $\Delta$ C, deletion of the last four residues) were subcloned into pcDNA3.1-HA. EGFP-DGK $\zeta$  was generated by subcloning full-length DGK $\zeta$  into pEGFP-C1, kinase-dead (G354D) DGK $\zeta$  was generated using QuickChange kit (Stratagene). The DGK $\zeta$  knock-down construct was generated by subcloning nt 159–177 (CAG-GAAAGCCATCACCAAG) into pSUPER.gfp/neo (OligoEngine). The PSD-95 knockdown construct was generated by subcloning nt 415–433 (GAGGCAGGTTCCATCGTTC) into pSUPER.gfp/neo (OligoEngine). GW1-PSD-95, GW1-Myc-PSD-95, and GW1-PSD-95-EGFP have been described (Kim *et al.*, 1995; Hsueh *et al.*, 1997; Arnold and Clapham, 1999).

#### Antibodies

GST-DGK $\zeta$  (aa 661–929) was used to generate rabbit (1519) and guinea-pig (1521) polyclonal antibodies. The following antibodies have been reported: PSD-95 (SM55), PSD-95 (1402), EGFP (1167), PSD-93/chapsyn-110 (B9594), SAP97 (B9591), SAP102 (1445), and GRIP2 (1757). The other antibodies were purchased: Myc, HA, and synaptophysin (Santa Cruz), synapsin I (Chemicon), phospho-(Ser) PKC substrate (Cell Signaling Technology), and  $\alpha$ -tubulin and MAP2 (Sigma, St Louis, MO).

#### Fractionation of rat brain lysates

Subcellular fractions of rat brains (6 weeks) were prepared as described (Huttner *et al*, 1983). PSD fractions were prepared as described (Cho *et al*, 1992).

#### Electron microscopy

Two male Sprague–Dawley rats (280–300 g) were deeply anaesthetized with sodium pentobarbital (80 mg/kg, i.p.) and intracardially perfused with a mixture of 4% paraformaldehyde and 0.01% glutaraldehyde in phosphate buffer (PB, pH 7.4). Hippocampus was removed and cut at 60 µm on a vibratome and then cryoprotected in 30% sucrose in PB overnight. Cryoprotected sections were frozen and thawed, and then treated with  $3\,\%$   $H_2O_2$  and  $10\,\%$  normal donkey serum. The sections were incubated overnight in the guinea-pig DGKζ antibodies (1521; 4 µg/ml) at room temperature and then with a biotinylated donkey anti-guinea-pig antibody (1:200; Jackson Immunoresearch) for 2 h. After washing, the sections were incubated with ExtrAvidin peroxidase (1:5000; Sigma), and the immunoperoxidase was visualized by nickelintensified 3,3'-diaminobenzidine tetrahydrochloride. Sections were osmicated, dehydrated in graded alcohols, embedded with Durcupan ACM (Fluka, Buchs, Switzerland), and cured for 48 h at 60°C. Area containing pyramidal cell layer and stratum radiatum of the CA1 region was trimmed. Thin sections were cut, mounted on nickel grids, and stained with uranyl acetate and lead citrate. Grids were examined on a Hitachi H 7500 electron microscope at 80 kV accelerating voltage.

#### Neuron culture and transfection

Cultured neurons prepared from rat hippocampi at E18 were maintained in neurobasal medium supplemented with B27 (Life

Technologies). Neurons were transfected using the calcium phosphate method.

#### PA analysis

Hippocampal slices were incubated in phosphate-free artificial cerebrospinal fluid for 1 h, followed by metabolic labelling with  $10 \,\mu$ Ci/ml of <sup>32</sup>P-orthophosphate (GE Healthcare Life Sciences) for 30 min in the presence of 1% ethanol at 30°C. Labelled slices were incubated with  $100 \,\mu$ M DHPG for 20 min. Lipid extraction and thin layer chromatography were carried out as described previously (Nelson *et al*, 2007).

### DiOlistic spine labelling and image analysis

Transcardially perfused 3-week-old mice brain slices were labelled by the ballistic delivery of the lipophilic dye DiI as described (Gan *et al*, 2000). After dye delivery, brain slices were kept in phosphatebuffered saline for 4 h to allow dye diffusion. *Z*-stacked images from the proximal stratum radiatum region of hippocampus were used for blind image analysis. Spines were defined by protrusions with a bulbous head wider than the neck and with length longer than  $0.5 \,\mu$ m. Spine density was measured from 40- to 80- $\mu$ m segments of secondary dendrites on ~1-2 dendrites per neuron. The narrowest point in the spine neck was defined as the border between spine head and neck. MetaMorph software (Molecular Devices) was used for quantitative analysis.

#### Electrophysiology

For whole-cell recordings, CA1 pyramidal cell was held at -70 mV using a MultiClamp 700B amplifier (Axon Instruments). Pipette solutions contained (in mM) 110 Cs-gluconate, 30 CsCl, 20 HEPES, 4 MgATP, 0.3 NaGTP, 4 NaVitC, 0.5 EGTA (10 EGTA for AMPA versus NMDAR ratio) for mEPSC, or 145 CsCl, 20 HEPES, 4 MgATP, 0.3 NaGTP, 10 EGTA for mIPSC. TTX (0.5  $\mu$ M) and bicuculline (20  $\mu$ M) were added to ACSF for mEPSC measurements, and bicuculline was replaced with CNQX (10  $\mu$ M) and AP5 (50  $\mu$ M) for mIPSC measurements. To obtain AMPA receptor-mediated EPSCs, neuron in ACSF with 100  $\mu$ M picrotoxin was held at -70 mV and stimulated at 0.066 Hz. NMDAR-mediated EPSC was isolated by changing holding potential to +40 mV and adding 10  $\mu$ M CNQX to ACSF. EPSC samples ( $\sim$ 25–30) were averaged to obtain the AMPA/NMDA EPSC ratio.

#### Time-lapse imaging

Cultured neurons expressing sh-DGK $\zeta$ , or control vector (sh-vec), were mounted in a chamber (Zeiss) where temperature (37°C) and CO<sub>2</sub> concentration (10%) were maintained during the recording. Images were acquired with an LSM510 confocal microscope (Zeiss) using a C-Apochromat × 63, 1.20 NA water-immersion objective. The laser power was attenuated to less than 3% to avoid fluorescence bleaching and neuronal toxicity, and images were taken every 10 min for 2 h.

#### Supplementary data

Supplementary data are available at *The EMBO Journal* Online (http://www.embojournal.org).

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