



## RESEARCH ARTICLE

### ENHANCEMENT OF PKC $\epsilon$ ACTIVITY THROUGH SERINE AUTOPHOSPHORYLATION

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#### ABSTRACT

The linoleic acid derivative 8-[2-(2-pentyl-cyclopropylmethyl)-cyclopropyl]-octanoic acid (DCP-LA), with cyclopropane rings instead of *cis*-double bonds, serves as a selective activator of PKC $\epsilon$ . The present study investigated whether PKC $\epsilon$ , activated by DCP-LA, phosphorylates PKC $\epsilon$  itself and how phosphorylated PKC $\epsilon$  affects its own activity. DCP-LA phosphorylated PKC $\epsilon$  at the serine residues, but not the threonine residues, in PC-12 cells, which is abolished the PKC inhibitor GF109203X. The plasmids for wild-type and mutant PKC $\epsilon$  were transfected into MSTO-211H human malignant mesothelioma cells with very little expression of the PKC $\epsilon$  mRNA. DCP-LA enhanced PKC $\epsilon$  activity approximately 7 folds in MSTO-211H cells transfected with the wild-type rat PKC $\epsilon$  plasmid as compared with that in non-transfected control cells. Such effect was not obtained with cells transfected with the plasmid for mutant rat PKC $\epsilon$  replacing Ser234, Ser316, Ser368, or Ser729 by Ala. Taken together, these results indicate that PKC $\epsilon$ , activated by DCP-LA, enhances its own activity through serine autophosphorylation

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## INTRODUCTION

PKC isozymes are classified into conventional PKCs such as PKC $\alpha$ , - $\beta$ I, - $\beta$ II, and - $\gamma$ , novel PKCs such as PKC $\delta$ , - $\epsilon$ , - $\eta$ , - $\theta$ , and - $\mu$ , and atypical PKCs such as PKC $\lambda$ /I for mouse/human, - $\zeta$ , and - $\nu$ . PKCs are activated through several pathways linked to phospholipase C (PLC), phospholipase A<sub>2</sub> (PLA<sub>2</sub>), phospholipase D (PLD), and phosphatidylcholine-specific PLC (Isakov, 1993; Nishizuka, 1992; Nishizuka, 1995). PLC hydrolyzes phosphatidylinositol 4,5-bisphosphate into diacylglycerol and inositol 1,4,5-trisphosphate (IP<sub>3</sub>), the latter activating IP<sub>3</sub> receptors to release Ca<sup>2+</sup> from intracellular calcium stores, and conventional PKCs are activated by diacylglycerol and Ca<sup>2+</sup> (Nishizuka, 1992; Nishizuka, 1995). Phosphatidylcholine-specific PLC produces diacylglycerol by hydrolysis of phosphatidylcholine, thereby activating PKC (Isakov, 1993). *cis*-Unsaturated free fatty acids such as arachidonic, oleic, linoleic, linolenic, and docosahexaenoic acid, that are produced by PLA<sub>2</sub>-catalyzed hydrolysis of phosphatidylcholine, activate novel PKCs in a Ca<sup>2+</sup>-independent manner (Nishizuka, 1992; Nishizuka, 1995). The free fatty acids, alternatively, are implicated in synergistic activation of conventional PKCs or sustained activation of conventional PKCs activated (Nishizuka, 1992; Nishizuka, 1995). The linoleic acid derivative 8-[2-(2-pentyl-cyclopropylmethyl)-cyclopropyl]-octanoic acid (DCP-LA),

with cyclopropane rings instead of *cis*-double bonds, activates PKC $\epsilon$  selectively (Tanaka and Nishizaki, 2003; Kanno *et al.*, 2006). DCP-LA binds to the phosphatidylserine binding/associating sites Arg50 and Ile89 in the C2-like domain of PKC $\epsilon$  at the carboxyl-terminal end and the cyclopropane rings, respectively, which are distinct from the phorbol 12-myristate 13-acetate (PMA) binding site in the C1 domain (Kanno *et al.*, 2015). PKC $\epsilon$  consists of the regulatory domain including the C2-like domain and the C1A/1B domain, the V3 domain (hinge domain), and the kinase (catalytic) domain including the C3 domain, the V4 domain, the C4 domain, and the V5 domain (Steinberg, 2008). PKC $\epsilon$  as well as other PKC isozymes contain several serine/threonine phosphorylation sites. Little, however, is known about the functional role of PKC $\epsilon$  autophosphorylation. To address this question, the present study examined how phosphorylated PKC $\epsilon$  affects the PKC $\epsilon$  activity. The results show that PKC $\epsilon$ , activated by DCP-LA, enhances its own activity through serine autophosphorylation.

## MATERIALS AND METHODS

**Cell culture:** PC-12 cells, that were obtained from RIKEN Cell Bank (Tsukuba, Japan), were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% (v/v) heat-inactivated fetal bovine serum, 10% (v/v) heat-

inactivated horse serum, penicillin (100 U/mL), and streptomycin (0.1 mg/mL) in a humidified atmosphere of 5% CO<sub>2</sub> and 95% air at 37 °C. PC-12 cells were differentiated by treatment with nerve growth factor (100 ng/mL) for 5 days.

Human malignant mesothelioma cell lines MSTO-211H and NCIH-28 cells were purchased from American Type Culture Collection (Manassas, VA, USA). Cells were grown in RPMI-1640 medium supplemented with 10% heat-inactivated FBS, 0.003% L-glutamine, penicillin (100 U/mL), and streptomycin (0.1 mg/mL) in a humidified atmosphere of 5% CO<sub>2</sub> and 95% air at 37 °C.

**Immunoprecipitation and Western Blotting:** PC-12 cells, treated with dimethyl sulfoxide (DMSO) or DCP-LA (100 nM) for 10 min, were homogenized by sonication in TBS-T [150 mM NaCl, 0.1% (v/v) Tween-20 and 20 mM Tris, pH 7.5] containing 1% (v/v) phosphatase inhibitor cocktail and subsequently, homogenates were centrifuged at 3,000 rpm for 5 min at 4 °C. The supernatants (200  $\mu$ g of protein) were incubated with an antibody against PKC $\epsilon$  (BD Biosciences, San Jose, CA, USA) overnight at 4 °C. Then, 20  $\mu$ L of protein G sepharose (GE healthcare, Piscataway, NJ, USA) was added to the extracts and incubated for 60 min at 4 °C. Pellets were washed three times with TBS-T and dissolved in 30  $\mu$ L of a sodium dodecyl sulfate (SDS) sample buffer [0.2 mM Tris, 0.05% (w/v) SDS, and 20% (v/v) glycerol, pH 6.8]. After boiling for 5 min, proteins were separated by SDS-polyacrylamide gel electrophoresis (SDS-PAGE) using a TGX gel (BioRad, Hercules, CA, USA) and then transferred to polyvinylidene difluoride membranes. Blotting membranes were blocked with TBS-T containing 5% (w/v) bovine serum albumin and subsequently incubated with an anti-phospho-Ser (pSer) antibody (QIAGEN, Hilden, Germany) or an anti-phospho-Thr (pThr) antibody (QIAGEN). After washing, membranes were reacted with a horseradish peroxidase-conjugated goat anti-mouse IgG antibody. Immunoreactivity was detected with an ECL kit (GE Healthcare) and visualized using a chemiluminescence detection system (GE Healthcare). Protein concentrations for each sample were determined with a BCA protein assay kit (Pierce, Rockford, IL, USA).

**Real-time reverse transcription-polymerase chain reaction (RT-PCR):** Real-time RT-PCR was carried-out by the method previously described (Tsuchiya *et al.*, 2012). Briefly, total RNAs from MSTO-211H and NCIH-28 cells were purified using a Sepasol-RNA I Super kit (Nacalai, Kyoto, Japan) and treated with RNase-free DNase I (2 units) for 30 min at 37 °C to remove genomic DNAs. RNAs purified was incubated in a RT buffer containing random primers, dNTP, and Multiscribe Reverse Transcriptase for 10 min at 25 °C and in turn, for 120 min at 37 °C to synthesize the first-strand cDNA. Real-time RT-PCR was performed using a SYBR Green Realtime PCR Master Mix (Takara Bio, Otsu, Japan) and the Applied Biosystems 7900 real-time PCR detection system (ABI, Foster City, CA). Thermal cycling conditions were as follows: first step, 94 °C for 4 min; the ensuing 40 cycles, 94 °C for 1 s, 65 °C for 15 s, and 72 °C for 30 s. The mRNA quantity for each PKC isozyme was calculated from the standard curve made by amplifying different amount of the GAPDH mRNA. Primers used for real-time RT-PCR are shown in Table I.

**Construction of the plasmid for PKC $\epsilon$  and transfection:** Nucleotide sequence coding for wild-type rat PKC $\epsilon$  was cloned into pcDNA6/V5-His A vector (Invitrogen) at the KpnI-XhoI site.

For mutant rat PKC $\epsilon$  the serine residues at 234, 316, 368, and 729 were replaced by alanine (mS234A, mS316A, mS368A, and mS729A, respectively). The plasmids for wild-type and mutant PKC $\epsilon$  were constructed, and transfected into MSTO-211H cells using a Lipofectamine reagent (Invitrogen). Cells were used for experiments 48 h after transfection.

### **In situ PKC assay**

PKC activity in cells was assayed by the method as previously described (Kanno *et al.*, 2006). Cells were treated with DMSO or DCP-LA (100 nM) at 37 °C for 10 min in an extracellular solution [137 mM NaCl, 5.4 mM KCl, 10 mM MgCl<sub>2</sub>, 5 mM ethylene glycol-bis(2-aminoethyl ether)-N,N,N',N'-tetraacetic acid, 0.3 mM Na<sub>2</sub>HPO<sub>4</sub>, 0.4 mM K<sub>2</sub>HPO<sub>4</sub>, and 20 mM 4-(2-hydroxyethyl)-1-piperazine-ethanesulfonic acid, pH 7.2]. Then, cells were rinsed with 100  $\mu$ L of Ca<sup>2+</sup>-free phosphate-buffered saline (PBS) and incubated at 30 °C for 15 min in 50  $\mu$ L of the extracellular solution containing 50  $\mu$ g/mL digitonin, 25 mM glycerol 2-phosphate, 200  $\mu$ M ATP, and 100  $\mu$ M synthetic PKC substrate peptide (Pyr-Lys-Arg-Pro-Ser-Gln-Arg-Ser-Lys-Tyr-Leu) (Peptide Institute Inc., Osaka, Japan). The supernatants were collected and boiled at 100 °C for 5 min to terminate the reaction. Aliquot of the solution (20  $\mu$ L) was loaded onto a reversed phase high performance liquid chromatography (HPLC) (LC-10ATvp, Shimadzu Co., Kyoto, Japan). A substrate peptide peak and a new product peak were detected at an absorbance of 214 nm (SPD-10Avp UV-VIS detector, Shimadzu Co., Kyoto, Japan). It was confirmed that each peak corresponds to non-phosphorylated and phosphorylated substrate peptide in the analysis of matrix-assisted laser desorption ionization time of flight mass spectrometry (Voyager DE-STR, PE Biosystems Inc., Foster City, USA). Molecular weights were calibrated from the two standard spectrums, bradykinin (MW 1060.2) and neurotensin (MW 1672.9). Areas for non-phosphorylated and phosphorylated PKC substrate peptide were measured (total area corresponds to the concentration of PKC substrate peptide used here). The quantity of phosphorylated substrate peptide (pmol/min/cell protein weight) was calculated and used as an index of PKC activity.

**Statistical analysis:** Statistical analysis was carried out using analysis of variance (ANOVA) followed by a Bonferonni correction.

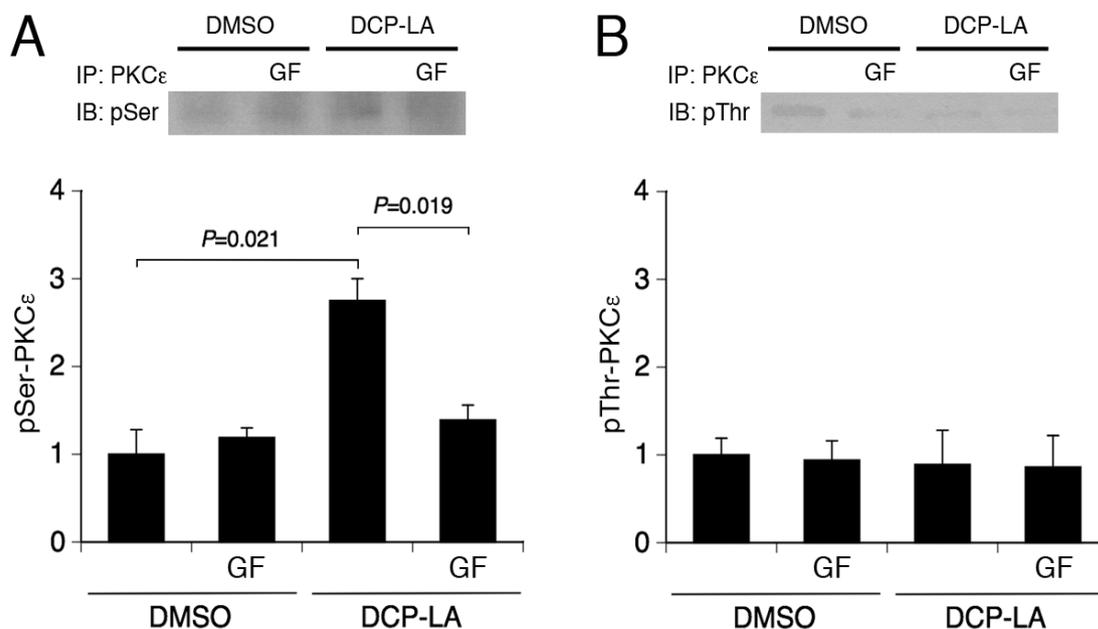
## **RESULTS**

**PKC $\epsilon$  phosphorylates PKC $\epsilon$  itself at the serine residues:** DCP-LA is recognized to activate PKC $\epsilon$  selectively (Kanno *et al.*, 2006). The signal intensity for pSer-PKC $\epsilon$  in the immunoprecipitants from PC-12 cells treated with DCP-LA using an anti-PKC $\epsilon$  antibody was significantly enhanced as compared with that in the immunoprecipitants from untreated control cells (Figure 1A). The effect of DCP-LA was clearly inhibited by the PKC inhibitor GF109203X (Figure 1A). In contrast, DCP-LA had no effect on the signal intensity for pThr-PKC $\epsilon$  (Figure 1B). Collectively, these results indicate that DCP-LA activates PKC $\epsilon$ , to phosphorylate its own PKC $\epsilon$  or each other's PKC $\epsilon$  at the serine residues, but not the threonine residues.

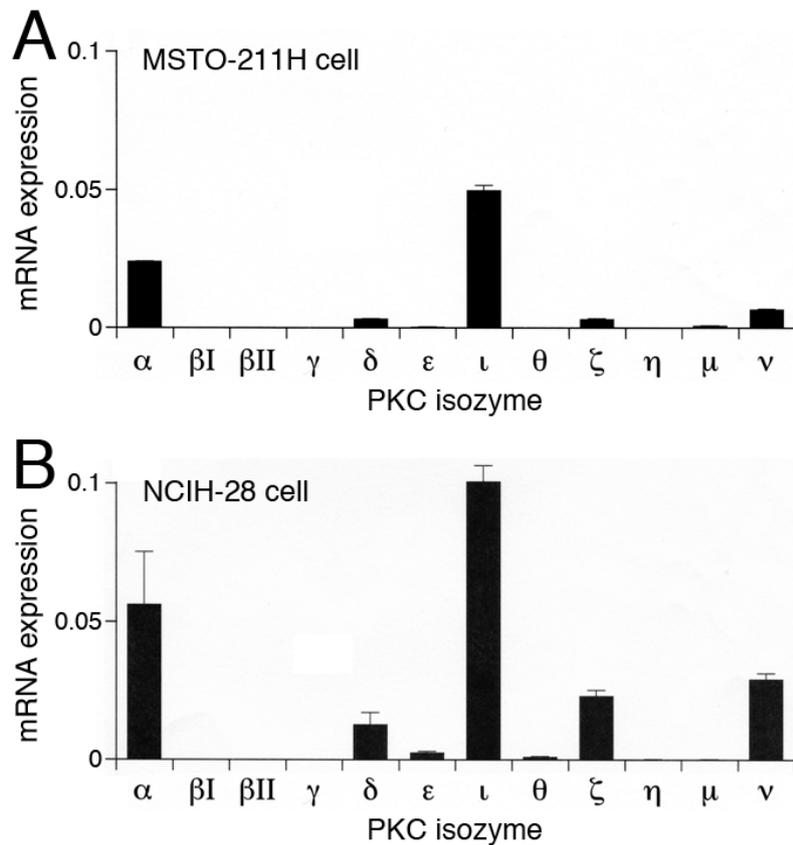
**PKC $\epsilon$  enhances its activity through its own serine phosphorylation:** The next attempt was to understand the effect of PKC $\epsilon$  phosphorylation on the PKC $\epsilon$  activity.

**Table 1.** Primers used for real-time RT-PCR.

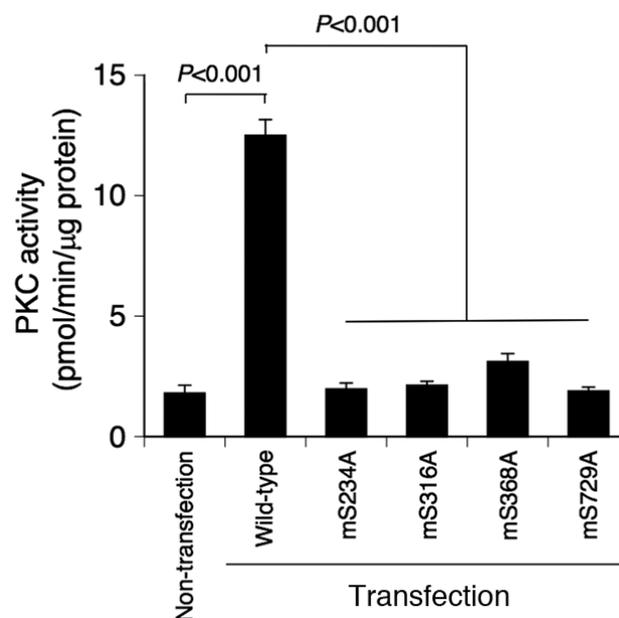
<i>PKC<math>\alpha</math></i>	Sense	GCCGTATGGAAAATCTGTGGACTGG
	Anti-sense	TGGGCTTGAATGGTGGCTGGAT
<i>PKC<math>\beta</math>I</i>	Sense	TTTGAAGGGGAGGATGAAGATGAACTC
	Anti-sense	TGAAGAGTTTATCAGTGGGGGTCAGTTC
<i>PKC<math>\beta</math>II</i>	Sense	TTTGAAGGGGAGGATGAAGATGAACTC
	Anti-sense	TGAATGACAGAAATGAAGGACGGAGAT
<i>PKC<math>\gamma</math></i>	Sense	GCAGCCCCACTTCACCCCC
	Anti-sense	CCAGAAATCCCCAGAGCACAGCA
<i>PKC<math>\delta</math></i>	Sense	CAACAGTGGGACCTACGGCAAGAT
	Anti-sense	CCTGTAAATGATGCCCTTGCTGTGTAG
<i>PKC<math>\epsilon</math></i>	Sense	ATGCCCCACAAGTTCGGTATCCAC
	Anti-sense	GGTGCTCCTCTCCTCGGTTGTCA
<i>PKC<math>\iota</math></i>	Sense	TAGATGAGGAAGGAGACCCGTGTACAGTAT
	Anti-sense	CTTCCCTGGTGTTCATTGCCTCTT
<i>PKC<math>\theta</math></i>	Sense	GGACTGGCACGGCAAGGACTC
	Anti-sense	GCAGAGATGGTCTTTCTTTGTTTCAGTTCA
<i>PKC<math>\zeta</math></i>	Sense	GCCTCCAGTAGACGACAAGAACGAGG
	Anti-sense	CGTAGAACCTGGCGTGCTCCTCA
<i>PKC<math>\eta</math></i>	Sense	CTATGCCAAGGCGAGTCCACCAG
	Anti-sense	TCCTTGTCGCATTATTCCCCAGAG
<i>PKC<math>\mu</math></i>	Sense	AGTGCCCCTGATGAGCCCCTT
	Anti-sense	CCCCTGTTTCCTTCACTATCATTGTCA
<i>PKC<math>\nu</math></i>	Sense	ACTCTGCCCCGACTCTCTAATGGAAG
	Anti-sense	CCACAGCCTTACATTTTCAGTCCTTG



**Figure 1.** PKC $\epsilon$ , activated by DCP-LA, phosphorylates PKC $\epsilon$  itself at the serine residues. Lysates from PC-12 cells treated with DMSO or DCP-LA (100 nM) for 10 min in the presence and absence of GF109203X (GF) (100 nM), were immunoprecipitated with an anti-PKC $\epsilon$  antibody, followed by Western blotting using antibodies against phospho-serine (pSer) and phospho-threonine (pThr). IP, immunoprecipitation; IB, immunoblot. In the graphs, each column represents the mean ( $\pm$  SEM) signal intensity for pSer-PKC $\epsilon$  or pThr-PKC $\epsilon$  relative to that for cells treated with DMSO in the absence of GF109203X (n=4 independent experiments). *P* values, ANOVA followed by a Bonferroni correction



**Figure 2.** Expression of the PKC isozyme mRNAs in human malignant mesothelioma cell lines. Real-time RT-PCR was carried out in MSTO-211H and NCIH-28 cells. In the graphs, each point represents the mean ( $\pm$  SEM) mRNA quantity ( $n=4$  independent experiments).



**Figure 3.** PKC activity. MSTO-211H cells, non-transfected and transfected with the wild-type and mutant rat PKC $\epsilon$  plasmids as indicated, were treated with DCP-LA (100 nM) for 10 min, and PKC activity was assayed. In the graph, each column represents the mean ( $\pm$  SEM) PKC activity ( $n=6$  independent experiments).  $P$  values, ANOVA followed by a Bonferroni correction.

In the real-time RT-PCR, the PKC $\epsilon$  mRNA was very little expressed in MSTO-211H human malignant mesothelioma cells (Figure 2A), while expression of the PKC $\epsilon$  mRNA was detected at the low levels in NCIH-28 cells (Figure 2B). DCP-LA enhanced the PKC activity in MSTO-211H cells transfected with the wild-type rat PKC $\epsilon$  plasmid to more than 7 folds than that in non-transfected control cells (Figure 3).

In contrast, no enhancement of the PKC activity was obtained with cells transfected with the plasmid for mutant rat PKC $\epsilon$  replacing Ser234, Ser316, Ser368, or Ser729 by Ala (Figure 3). Overall, these results indicate that PKC $\epsilon$ , activated by DCP-LA, phosphorylates PKC $\epsilon$  itself at the serine residues, to enhance the PKC $\epsilon$  activity.

## DISCUSSION

As is the case with other PKC isozymes, PKC $\epsilon$  is inactivated in the closed conformation at the hinge domain, and when the closed conformation is opened by binding phosphatidylserine, diacylglycerol, or free fatty acids, PKC $\epsilon$  is activated. The linoleic acid derivative DCP-LA activates PKC $\epsilon$  selectively by binding to the phosphatidylserine binding/associating sites in the C2-like domain (Kanno *et al.*, 2006; Kanno *et al.*, 2015). PKC contains the phosphorylation sites on PKC itself (Steinberg, 2008). In the present study, DCP-LA significantly increased serine phosphorylation of PKC $\epsilon$ , that is abolished by the PKC inhibitor GF109203X. This indicates that PKC $\epsilon$ , activated by DCP-LA, phosphorylates PKC $\epsilon$  at the serine residues by itself or each other. Intriguingly, DCP-LA had no effect on threonine phosphorylation of PKC $\epsilon$ . This explains that PKC $\epsilon$  preferentially phosphorylates its own serine residues. DCP-LA markedly enhanced the PKC $\epsilon$  activity in human MSTO-211H cells with very poor expression of PKC $\epsilon$  by transfecting with the wild-type rat PKC $\epsilon$  plasmid. DCP-LA, however, had no effect on the PKC activity in MSTO-211H cells transfected with the plasmid for the mutant rat PKC $\epsilon$  mS234A, mS316A, mS368A, or mS729A. These results indicate that the PKC $\epsilon$  activity is enhanced through its own serine phosphorylation. PKC $\epsilon$ -Ser729 belongs to the hydrophobic motif located in the V5 domain (Steinberg, 2008). The V5 domain is critical for the PKC $\epsilon$  catalytic activity and function; ATP binding pocket is formed between the kinase domain and the V5 domain (Steinberg, 2008). PKC $\epsilon$ -Ser729 phosphorylation is supposed to sustain the PKC $\epsilon$  activity by dissociating Ser/Thr protein phosphatase. The present results suggest that other serine phosphorylation sites PKC $\epsilon$ -Ser234 in the C2-like domain, PKC $\epsilon$ -Ser316 in the C1A/1B domain, and PKC $\epsilon$ -Ser368 in the C1A/1B domain as well as PKC $\epsilon$ -Ser729 in the V5 domain regulate the PKC $\epsilon$  catalytic activity.

## Conclusion

The results of the present study show that PKC $\epsilon$ , activated by DCP-LA, phosphorylates its own serine residues, to enhance the PKC $\epsilon$  activity. This may extend our understanding about the mechanism of PKC $\epsilon$  activation.

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