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Upregulation of PKC η by PKC ϵ and PDK1 involves two distinct mechanisms and promotes breast cancer cell survival

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Abstract

Background—Protein kinase C (PKC) serves as the receptor for tumor-promoting phorbol esters, which are potent activators of conventional (c) and novel (n) PKCs. We recently showed that these activators induced selective upregulation of PKC η in breast cancer cells. The objective of this study is to understand unique regulation of PKC η and its importance in breast cancer.

Methods—The levels of PKC isozymes were monitored in breast cancer cells following treatment with inhibitors of kinases, proteasome and proteases by Western blotting. PKC ϵ was introduced by adenoviral delivery. PKC η and PDK1 were depleted by siRNA silencing. Cell growth was determined by the MTT or clonal assay.

Results—The general PKC inhibitors Gö 6983 and bisindolylmaleimide but not cPKC inhibitor Gö 6976 led to substantial PKC η downregulation, which was partly rescued by the introduction of nPKC ϵ . Inhibition of phosphoinositide-dependent kinase-1 (PDK1) by Ly294002 or knockdown of PDK1 also led to downregulation of basal PKC η but had no effect on PKC activator-induced upregulation of PKC η . Proteasome inhibitors blocked PKC η downregulation triggered by PDK1 inhibition/depletion but not by Gö 6983. PKC η level increased in malignant but not in non-tumorigenic or pre-malignant cells in the progressive MCF-10A series associated with activated PDK1, and knockdown of PKC η inhibited breast cancer cell growth and clonogenic survival.

Conclusion—Upregulation of PKC η contributes to breast cancer cell growth and targeting either PKC ϵ or PDK1 triggers PKC η downregulation but involves two distinct mechanisms.

General significance—The status of PKC η may serve as a potential biomarker for breast cancer malignancy.

Keywords

PKC; PDK1; downregulation; breast cancer

1. Introduction

Protein kinase C, a family of phospholipid-dependent serine/threonine kinases, plays a critical role in growth factor signal transduction pathways [1, 2]. The PKC family is classified into conventional (α , β I, β II, γ), novel (δ , ϵ , η , θ) and atypical PKCs (ζ , λ / τ)

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based on their structural features and cofactor sensitivity [1, 2]. Diacylglycerol (DAG) and Ca^{2+} are required for the activity of conventional PKCs whereas novel PKCs are insensitive to Ca^{2+} but dependent on DAG; atypical PKCs are insensitive to both Ca^{2+} and DAG [1, 2].

Tumor-promoting phorbol esters are potent activators of PKCs and can substitute for DAG. However, sustained activation by phorbol esters leads to downregulation of PKCs causing termination of the PKC signaling. Calcium-activated proteases, such as calpains and ubiquitin proteasome-mediated pathways have been implicated in PKC activator-induced downregulation of PKCs. Calpains are believed to downregulate cPKCs [3, 4] whereas PKC α , - δ and - ϵ were shown to be degraded via proteasome-mediated pathway [5-7]. Downregulation of PKCs has important implications in regulating long-term cellular responses such as cell proliferation, differentiation and tumor promotion [1, 8, 9].

PKC η is a member of the novel PKC family and shares highest homology with PKC ϵ [10]. However, the regulation of PKC η appears to be unique. We have recently shown that several different PKC activators, including tumor-promoting phorbol esters, induce upregulation rather than downregulation of PKC η [11]. Moreover, phosphorylation of PKC η by novel PKC ϵ appears to be responsible for PKC activator-induced upregulation of PKC η [11]. Depending on the cellular context, inhibition of PKC η could either promote or suppress tumor promotion [12-19]. Since PKC η is upregulated by tumor promoters, we examined the mechanism of PKC η downregulation. Our results indicate that inhibition of either PKC or PDK1 induces PKC η downregulation but involves two distinct mechanisms. While inhibition of PKC resulted in PKC η downregulation via proteasome-independent pathway, inhibition of PDK1 led to PKC η downregulation via proteasome-dependent pathway. We also showed that PKC η level is increased in malignant but not in non-tumorigenic or pre-malignant cells in the progressive MCF-10A series and knockdown of PKC η inhibited breast cancer cell growth.

2. Materials and Methods

2.1. Materials

PDBu was purchased from Alexis Biochemicals (San Diego, CA). Ly294002 was purchased from Calbiochem (San Diego, CA) and Cell Signaling Technology, Inc (Danvers, MA). Gö 6983, Gö 6976, KT5720, Rottlerin, PD98059, BIM, U0126, MG-132, lactacystin, calpeptin and cathepsin inhibitor I were obtained from Calbiochem (San Diego, CA). Polyclonal antibodies to PKC η , PKC δ , PKC ϵ and monoclonal antibody to GAPDH were obtained from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA). Monoclonal antibody to PKC α was obtained from Upstate Biotechnology (Lake Placid, NY). Polyclonal antibody against PDK1 was purchased from Cell Signaling Technology, Inc. (Danvers, MA). Monoclonal antibody against actin was obtained from Sigma (St. Louis, MO). Horseradish-peroxidase-conjugated donkey anti-rabbit and goat anti-mouse secondary antibodies were purchased from Jackson ImmunoResearch Laboratories, Inc. (West Grove, PA). Control non-targeting siRNA and siRNA specific for PDK1 and PKC η were obtained from Dharmacon (Lafayette, CO). Poly(vinylidenedifluoride) membrane was from Millipore (Bedford, MA) and enhanced chemiluminescence detection kit was from Amersham (Arlington Heights, IL).

2.2. Cell culture

MCF-7 and T47D cells were maintained in RPMI 1640 medium supplemented with 10% fetal bovine serum and 2 mM glutamine. Cells were kept in a humidified incubator at 37°C with 95% air and 5% CO_2 . The MCF-10A series developed by Dr. Fred Miller and colleagues [20], was obtained from the Barbara Ann Karmanos Cancer Institute (Detroit, MI). They were cultured as described previously [21].

2.3. Transfection

Control non-targeting siRNA or SMARTpool siRNA against PDK1 or PKC η were introduced into cells using Lipofectamine RNAiMax (Invitrogen, Carlsbad, CA) and manufacturer's protocol. Cells were treated as indicated in the text and processed for Western blot analysis.

2.4. Immunoblot analysis

Cells were lysed in extraction buffer containing 1 mM DTT, protease inhibitors and phosphatase inhibitors. Equal amounts of protein were separated by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred on PVDF membranes. Western blot analysis was performed as described previously [22].

2.5. Clonogenic assay

Cells transfected with or without control non-targeting or PKC η siRNA were cultured at 37°C in a humidified incubator with 5% CO₂ until there were at least 50 cells per colony. At the end of the incubation, the cells were washed with PBS and incubated with 0.025% crystal violet solution for 15 minutes. Colonies were counted using ImageJ software (<http://rsbweb.nih.gov/ij/>), and the plate was photographed using the BioChemi System (BioImaging System, UVP, Upland, CA).

2.6. MTT assay

Cells transfected with or without control non-targeting or PKC η siRNA were plated in microtiter plates and incubated at 37°C and 5% CO₂. After 4 days, the number of viable cells was determined using 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT) as described previously [23].

2.7. Statistical analysis

Statistical significance was determined by Student's *t* test by using PASW Statistics (SPSS, Inc.). *p* values < 0.05 was considered statistically significant.

3. Results

3.1. Inhibitors of PKC and PI3K induce downregulation of PKC η

We have previously shown that the regulation of PKC η is unique since it is upregulated rather than downregulated by PKC activators in breast cancer cells [11]. In the present study, we examined if inhibition of PKC induces its downregulation. Figure 1A shows that the general PKC inhibitor bisindolylmaleimide caused substantial downregulation of PKC η whereas rottlerin, which inhibits novel PKC δ , had no effect; the conventional PKC inhibitor Gö 6976 caused only a modest decrease in PKC η level in MCF-7 cells. The PI3K inhibitor Ly294002 also induced PKC η downregulation whereas rapamycin, PD98059 and KT5720, which inhibit mTOR, MAPK and PKA, respectively had little or no effect (Fig. 1A). To determine if the ability of PKC or PI3K inhibitor to downregulate PKC η is a general phenomenon, we tested several different breast cancer cell lines. Figure 1B shows that similar to MCF-7 cells, bisindolylmaleimide and Ly294002 induced substantial downregulation of PKC η in T47D cells whereas mTOR inhibitor rapamycin had a modest effect and MEK inhibitor U0126 had no effect. These results suggest that both PKC and PI3K pathways are involved in regulating PKC η level.

3.2. PKC and PI3K inhibitors induce PKC η downregulation via two distinct pathways

Since PKC activators induce upregulation of PKC η , we examined if the general PKC inhibitor Gö 6983 or the PI3K inhibitor Ly294002 could block PKC activator-induced

upregulation of PKC η . Figure 2 shows that while the PKC activator PDBu induced downregulation of PKC α , - δ and - ϵ , it caused upregulation of PKC η . Consistent with the notion that PKC activity is required for PKC activator-induced downregulation of PKCs [5, 6, 24], the general PKC inhibitor Gö 6983 prevented PDBu-induced downregulation of PKC α and PKC δ . However, Gö 6983 was able to induce PKC η downregulation in PDBu-treated cells. The PI3K inhibitor Ly294002 caused a substantial decrease in basal PKC η level but the constitutive levels of PKC α , - δ and - ϵ were not altered by the Ly294002 treatment. In contrast to Gö 6983, Ly294002 was unable to induce PKC η downregulation in PDBu-treated cells. These results suggest that Gö 6983 and Ly294002 induced PKC η downregulation via two distinct pathways.

3.3. Ly294002 induces PKC η downregulation via ubiquitin proteasome-mediated pathway

Both calcium-activated proteases, such as calpains as well as ubiquitin proteasome-mediated pathway have been implicated in the activation-induced downregulation of PKCs [4-6]. Therefore, we examined the ability of calpain inhibitor calpeptin, cathepsin inhibitor I and proteasome inhibitor MG-132 or lactacystin in preventing downregulation of PKC η by Gö 6983 or Ly294002. Figure 3 shows that neither calpain inhibitor nor cathepsin inhibitor blocked PKC η downregulation induced by either Gö 6983 or Ly294002. Although the proteasome inhibitor MG-132 prevented PKC η downregulation by Ly294002, it had only a modest effect on Gö 6983-mediated downregulation of PKC η (Figs. 4A & 4B). Similar results were observed in T47D cells where two different proteasome inhibitors MG-132 and lactacystin blocked Ly294002-mediated but not Gö 6983-mediated downregulation of PKC η (Fig. 4C). These results demonstrate that Ly294002 but not Gö 6983 induces PKC η downregulation via the proteasome-mediated pathway.

3.4. Inhibition of PDK1 and PKC ϵ triggers PKC η downregulation

Since PDK1 is believed to phosphorylate PKC η at the activation loop [25] and Ly294002 inhibits PDK1 which acts downstream of PI3K, we examined if knockdown of PDK1 induces downregulation of PKC η . Figure 5A shows that similar to Ly294002, depletion of PDK1 by siRNA decreased basal PKC η level. The proteasome inhibitors MG-132 and lactacystin caused a modest increase in PKC η level in cells transfected with control non-targeting siRNA or in untransfected cells, and blocked PKC η downregulation by both Ly294002 and PDK1 knockdown (Fig. 5A). These results suggest that inhibition of PDK1 by Ly294002 targets PKC η to proteasome-mediated downregulation.

Since we have previously shown that novel PKC ϵ was responsible for PKC activator-induced upregulation of PKC η [11], we speculated that the general PKC inhibitor Gö 6983 triggers PKC η downregulation by inhibiting PKC ϵ . Since T47D cells were more dependent on PKC ϵ -mediated upregulation of PKC η [11], we infected T47D cells with adenoviral vector expressing either GFP or PKC ϵ . Figure 5B shows that adenoviral delivery of PKCs but not GFP increased basal PKCs level and partially blocked Gö 6983-induced downregulation of PKC η . Taken together, these results suggest that inhibition of PDK1 and novel PKC ϵ by Ly294002 and Gö 6983, respectively triggers PKC η downregulation via two distinct pathways.

3.5. PKC η is progressively increased in MCF-10A series and promotes breast cancer cell survival

Since PKC η level is regulated by PKC and PI3K/PDK1 pathways that have been implicated in breast cancer [26-28], we compared PKC η level in non-tumorigenic MCF-10A cells and breast cancer cells. The level and activation status of PDK1 was increased in the progressive MCF-10A series [21] that includes spontaneously immortalized non-tumorigenic mammary epithelial MCF-10A cells, premalignant MCF-10AT cells, ductal carcinoma in situ (DCIS)

and highly malignant MCF-10CA1d cells [20]. Figure 6A shows that PKC η is expressed only in malignant breast cancer cells but not in non-tumorigenic MCF-10A or premalignant MCF-10AT cells. The levels of other PKCs, such as PKC α , PKC δ , PKC ϵ or PKC ζ were not increased with the malignancy of the MCF-10A series. Similar to MCF-7 and T47D cells, both Gö 6983 and Ly294002 led to PKC η downregulation in CA1d cells (Fig. 6B).

To determine the functional significance of PKC η upregulation in breast cancer cells, we determined the consequence of PKC η depletion on the viability of breast cancer cells. Depletion of PKC η by siRNA (Fig. 7A) decreased MCF-7 cell survival in a long-term clonogenic assay (Figs. 7B & 7C). Knockdown of PKC η also inhibited the growth of both T47D (Fig. 7D) and DCIS (Fig. 7E) cells. These results suggest that PKC η contributes to the viability of breast cancer cells.

4. Discussion

Because of the pivotal role of the PKC family members in signal transduction and cell regulation, there have been significant efforts in understanding their function and regulation. Among the novel PKCs, most of the studies thus far have been focused on PKC δ and ϵ but little is known about the regulation of novel PKC η . Our results suggest that the regulation of PKC η is unique. We recently reported that persistent treatment with PKC activators induce upregulation of PKC η rather than downregulation which is seen with other phorbol ester-sensitive conventional and novel PKCs. In the present study, we show that downregulation of PKC η can be achieved via two distinct pathways. While inhibition of PKC induces PKC η downregulation via a proteasome-independent pathway, inhibition of PDK1 results in PKC η downregulation via a proteasome-dependent pathway. We also made a novel observation that PKC η is the only PKC isozyme which is upregulated in breast cancer cells compared to non-malignant cells and downregulation of PKC η inhibits breast cancer cell growth.

PKC η has been implicated in both tumor promotion and tumor suppression. For example, PKC η expression was decreased in hepatocellular carcinoma [16], and PKC η deficiency enhanced susceptibility to tumor formation in a two-stage skin carcinogenesis model [12, 13]. Moreover, overexpression of PKC η induced differentiation in keratinocytes [17, 19]. In contrast, PKC η has been implicated in breast cancer [29], glioblastoma [15], lung cancer [14] and renal cell carcinoma [18]. In addition, overexpression of PKC η induced anchorage-independent growth in NIH3T3 cells [30] and provided proliferative advantage in astrocytic tumor cells [31]. The contrasting function of PKC η is not unique and several other PKC isoforms have been shown to exert opposite effects depending on the cell type [1, 9, 27, 32]. Thus, it is important to understand how PKC η is regulated in a particular cellular context in order to understand its function.

We recently reported that the regulation of PKC η in breast cancer cells is distinct from other PKC isozymes. While brief exposure to PKC activators, such as tumor-promoting phorbol esters leads to activation of conventional and novel PKCs, persistent treatment with these activators leads to their downregulation [1, 2]. We showed that prolonged treatment with several structurally and functionally distinct PKC activators caused upregulation rather than downregulation of PKC η and a general PKC inhibitor led to PKC η downregulation [11]. When we compared the effects of pharmacological inhibitors of several different kinases, we found that not only PKC inhibitors but also the PI3K inhibitor Ly294002 induced PKC η downregulation.

PKCs are phosphorylated by autophosphorylation and transphosphorylation by other members of the PKC family as well as PDK1 [33-35]. We recently reported that transphosphorylation of PKC η by novel PKCs is responsible for its upregulation [11]. Since two different general PKC inhibitors bisindolylmaleimide and Gö 6983 induced PKC η

downregulation whereas the conventional PKC inhibitor Gö 6976 had only a modest effect, it is likely that inhibition of PKC η phosphorylation by novel PKC ϵ leads to its downregulation. Our observation that adenoviral delivery of PKC ϵ increased PKC η level both in control and Gö 6983-treated cells supports this notion. We, however, found that overexpression of PKC ϵ was unable to completely prevent Gö 6983-mediated downregulation of PKC η , presumably because Gö 6983 could inhibit PKC ϵ in PKC ϵ -overexpressing cells albeit to a lesser extent compared to endogenous PKC ϵ .

It has been reported that PDK1 directly phosphorylates PKC η resulting in its activation [25] but knockdown of PDK1 had little effect on PKC activator-induced upregulation of PKC η [11]. Although PDK1 was not required for PKC activator-induced upregulation of PKC η , depletion of PDK1 by siRNA or its inhibition by the PI3K inhibitor Ly294002 decreased basal PKC η level, suggesting that phosphorylation of PKC η by PDK1 also protects it from downregulation. However, the mechanism by which Gö 6983 and Ly294002 induced PKC η downregulation was distinct. While both Gö 6983 and Ly294002 decreased basal PKC η level, only Gö 6983 but not Ly294002 could induce PKC η downregulation in the presence of PDBu. Moreover, two distinct proteasome inhibitors MG-132 and lactacystin could prevent PKC η downregulation caused by Ly294002 or PDK1 knockdown but not by Gö 6983. These results suggest that while inhibition of PDK1 triggers PKC η downregulation via proteasome-dependent pathway, inhibition of PKC triggers PKC η downregulation via proteasome-independent pathway.

Both PKC and PI3K/PDK1/Akt signaling pathways play important roles in the development and progression of breast cancer [1, 26-28]. It has been reported that the level and activation status of PDK1 is progressively increased in the MCF-10A series [21] which includes spontaneously immortalized non-tumorigenic mammary epithelial MCF-10A, premalignant MCF-10AT and MCF10AT3G, ductal carcinoma in situ DCIS and fully malignant MCF10CA1a and MCF10CA1d cells [20, 21]. The MCF-10A series provides a unique cell culture model system to study the alterations in signaling molecules leading to breast cancer since these cells share common genetic background. We made a novel observation that PKC η is the only PKC isozyme that is progressively increased in the MCF-10 cell series but there was no correlation between the levels of conventional PKC α , novel PKC δ and ϵ and atypical PKC ζ and the malignant status of MCF10A-derived cells. While PKC η level correlated with the level/activation status of PDK1 in the MCF-10 cell series, there was no correlation between PKC η level and hormone receptor status. For example, it was difficult to detect PKC η in triple-negative MDA-MB-231 cells (data not shown). However, PKC η is regulated by similar mechanisms regardless of the hormone receptor status. Both PKC inhibitor Gö 6983 and PI3K/PDK1 inhibitor Ly294002 induced downregulation of PKC η in basal type MCF-10CA1d (Fig. 6B) and MCF-10CA1a cells (data not shown).

To determine the functional significance of PKC η in breast cancer cells we depleted PKC η using siRNA since a selective inhibitor of PKC η is currently not available. Silencing of PKC η by siRNA decreased the growth of both DCIS and T47D cells. Knockdown of PKC η also reduced the long-term clonogenic survival of MCF-7 cells, which express very high level of PKC η . Moreover, since PKC η is regulated by PKC ϵ and PDK1 that play critical roles in breast cancer, PKC η may serve as an important target for breast cancer therapy.

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Abbreviations

BIM	Bisindolylmaleimide
DAG	diacylglycerol
ILV	indolactam V
MAPK	mitogen-activated protein kinase

MTT	3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide
PKA	protein kinase A
PI3K	phosphatidylinositol 3-kinase
PDK1	phosphoinositide-dependent kinase-1
PKC	protein kinase C
aPKC	atypical PKC
cPKC	conventional PKC
nPKC	novel PKC
PDBu	Phorbol 12,13-dibutyrate

Highlights

- Unique regulation of PKC η by nPKC and PDK1 involves two distinct mechanisms.
- Upregulation of PKC η is associated with breast cancer malignancy.
- PKC η contributes to breast cancer cell growth.



Fig. 1. Effects of kinase inhibitors on PKC η levels. MCF-7 (A) or T47D (B) cells were treated with or without 10 μ M BIM, 10 μ M rottlerin, 1 μ M Gö 6976, 25 μ M Ly294002, 50 nM rapamycin, 50 μ M PD98059, 10 μ M U0126 or 2 μ M KT5720 for 15 h. Western blot analysis was performed with total cellular extracts using the indicated antibodies. Actin was used as a loading control.

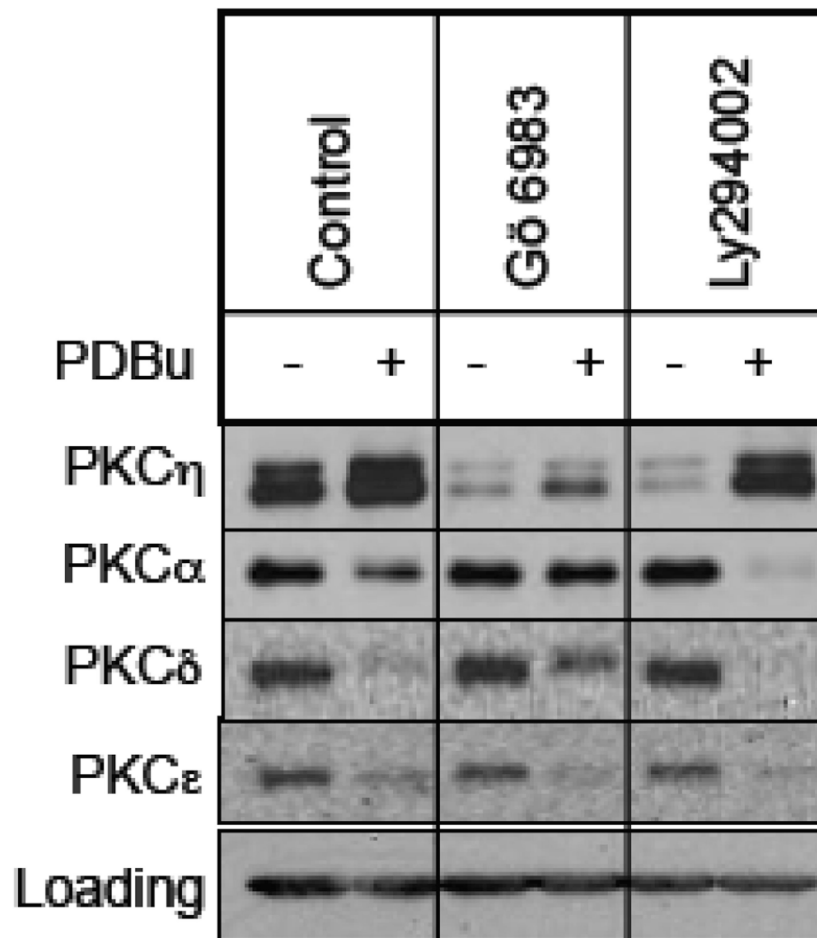


Fig. 2. Effect of PKC activators on PKC η downregulation by PKC and PI3K inhibitors. MCF-7 cells were pretreated with 1 μ M PDBu for 15 min, followed by treatment with or without 1 μ M Gö 6983 or 25 μ M Ly294002 for 15 h. Western blot analysis was performed with total cellular extracts using the indicated antibodies. Actin was used as a loading control.

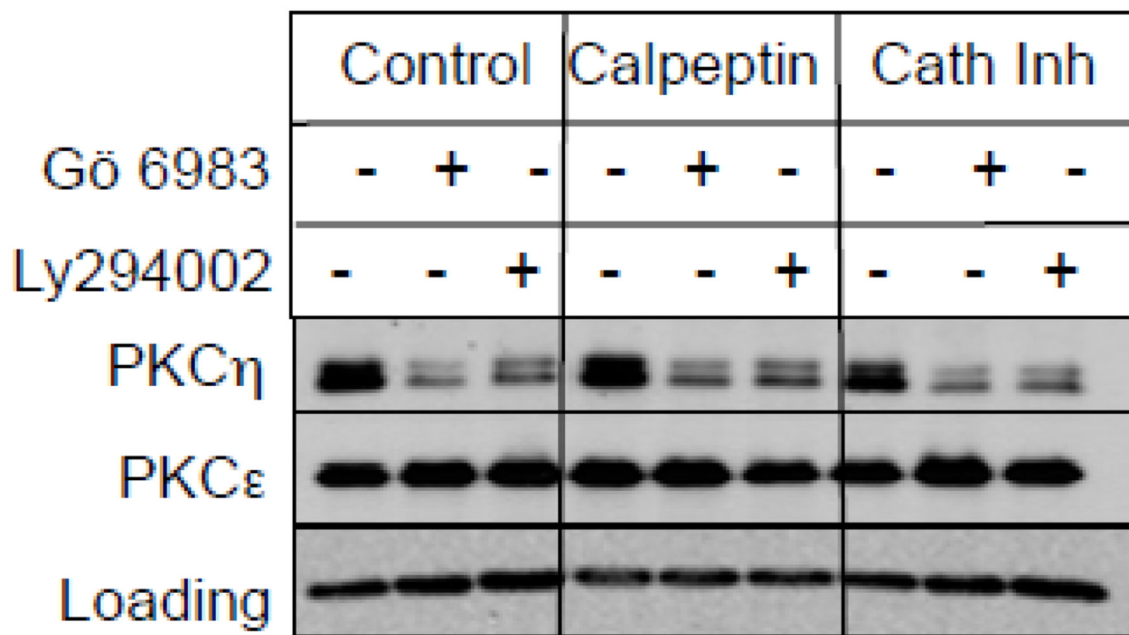
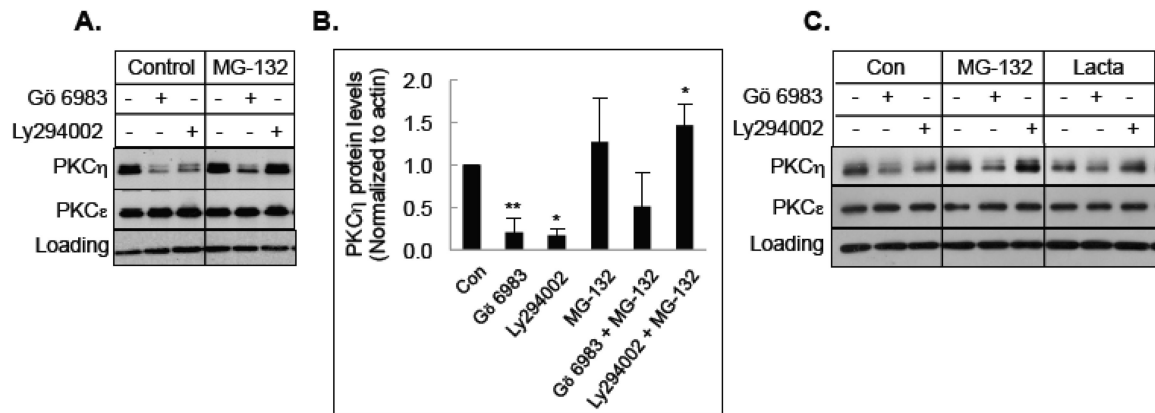


Fig. 3. Effect of protease inhibitors on PKC η downregulation in MCF-7 cells. MCF-7 cells were pretreated with or without 50 μ M calpeptin or 50 μ M cathepsin inhibitor I for 30 min, followed by treatment with 1 μ M Gö 6983 or 25 μ M Ly294002 for 12 h. Western blot analysis was performed with total cellular extracts using the indicated antibodies. GAPDH was used as a loading control. Results are representative of 2 independent experiments.

**Fig. 4.**

Effect of proteasome inhibitors on PKC η downregulation. A, MCF-7 cells were pretreated with 10 μ M MG-132 for 30 min, followed by treatment with 1 μ M Gö 6983 or 25 μ M Ly294002 for 12 h. Western blot analysis was performed with total cellular extracts using the indicated antibodies. GAPDH was used as a loading control. B, Densitometric quantification of PKC η protein levels from 3 separate experiments corrected for loading. Data represents the mean \pm s.e.m. The asterisk (*) indicates significant difference of MG-132 treated cells from control or treatment with the inhibitors alone using Student's *t*-test. *, $P < 0.05$; **, $P < 0.005$. C, T47D cells were pretreated with 10 μ M MG-132 or 10 μ M lactacystin for 30 min, followed by treatment with 1 μ M Gö 6983 or 10 μ M Ly294002 for 15 h. Western blot analysis was performed with total cellular extracts using the indicated antibodies. Actin was used as a loading control.

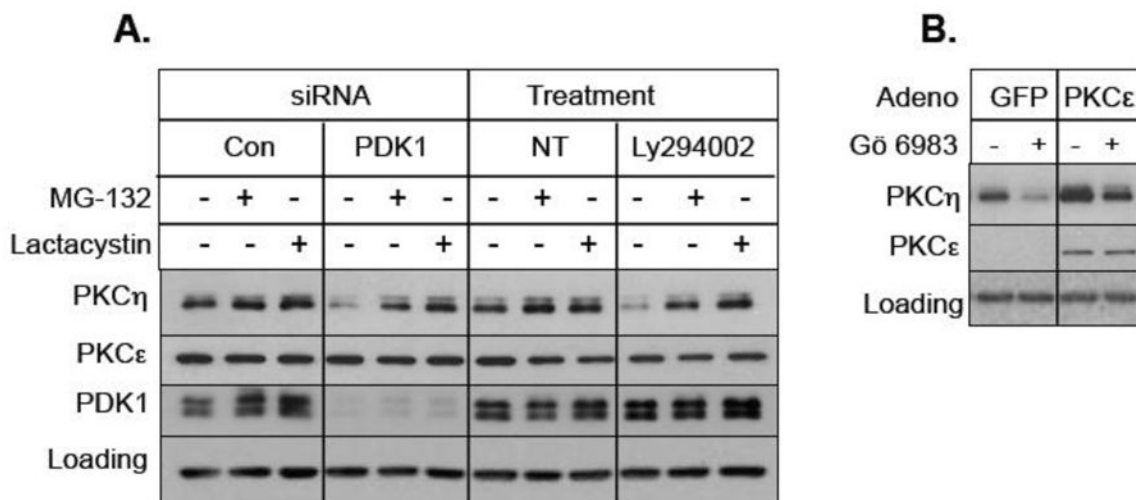


Fig. 5. Effect of PDK1 and PKC ϵ on PKC η downregulation. A, MCF-7 cells were transfected with control non-targeting siRNA, PDK1 siRNA or were left non-transfected (NT). Cells were then pre-treated with either 10 μ M MG-132 or 10 μ M lactacystin, followed by treatment with 25 μ M Ly294002. Western blot analysis was performed with indicated antibodies. Actin was used as a loading control. B, T47D cells were infected with adenovirus vector containing GFP or PKC ϵ construct and then treated with 1 μ M Gö 6983 for 15 h. Western blot analysis was carried out with indicated antibodies.

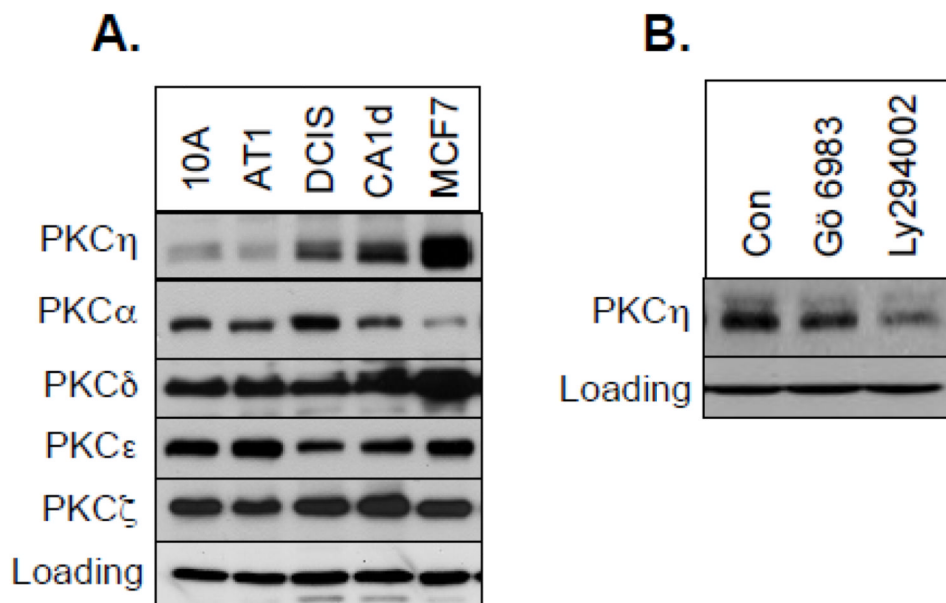
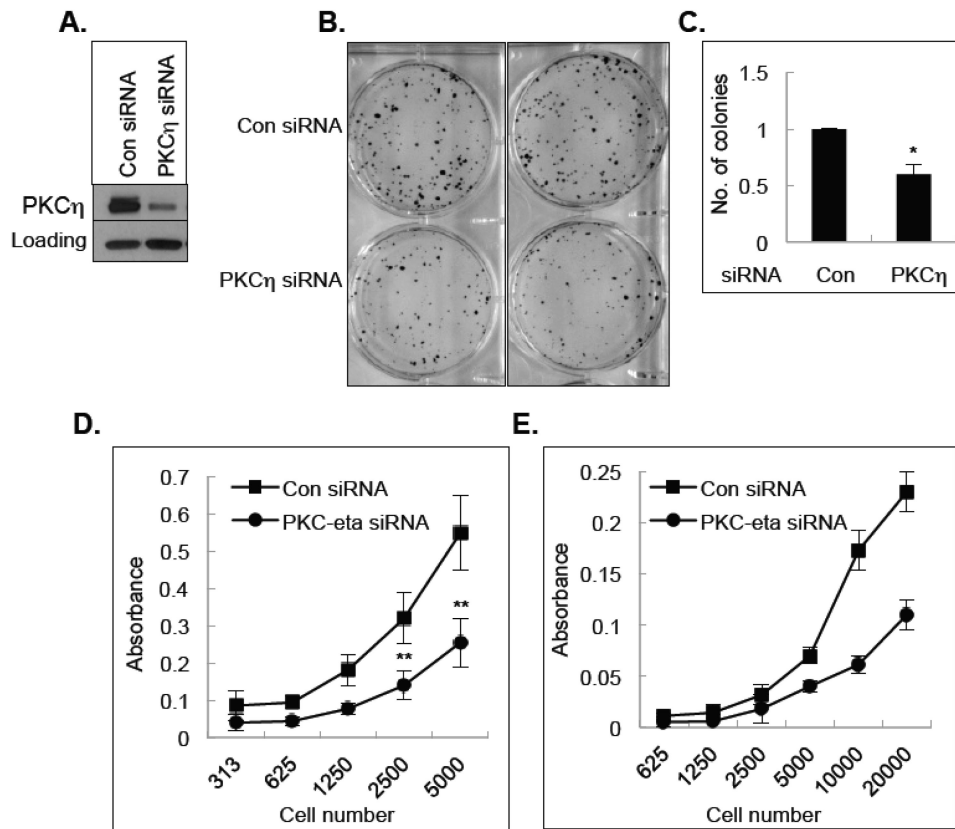


Fig. 6. Comparison of PKC η levels in MCF10A series. A, Western blot analysis was performed with total cellular extracts from 10A, AT1, DCIS, CA1d and MCF-7 cells and probed with the indicated antibodies. B, CA1d cells were treated with or without 1 μ M Gö 6983 or 25 μ M Ly294002 for 16 h. Western blot analysis was performed with total cellular extracts using the indicated antibodies. Actin was used as a loading control.

**Fig. 7.**

Effect of PKC η on cell growth. A, MCF-7 cells were transfected with control non-targeting siRNA or PKC η siRNA and total cell extracts were used for Western blot analysis. Actin was used as a loading control. B, Clonogenic assay was performed with MCF-7 cells transfected with either control non-targeting siRNA or PKC η siRNA as described in the Materials and Methods. C, Quantification of the number of colonies as determined by the clonogenic assay. The results are representative of 3 independent experiments. * $P < 0.05$ using paired Student t test. D, MTT assay was performed with T47D cells transfected with control, non-targeting siRNA or PKC η siRNA as described under Materials and Methods. The results are representative of 3 independent experiments. ** $P < 0.01$ using paired Student t test. E, MTT assay was performed with DCIS cells transfected with control non-targeting siRNA or PKC η siRNA as described under Materials and Methods.