

# Atypical Protein Kinase C- $\zeta$ Modulates Clonogenicity, Motility, and Secretion of Proteolytic Enzymes in Murine Mammary Cells

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In this paper, we investigated whether protein kinase C- $\zeta$  (PKC $\zeta$ ), a member of the atypical PKC family, induces phenotypic alterations associated with malignant transformation and tumor progression in mammary cells. The stable overexpression of PKC $\zeta$  in immortalized mammary epithelial cells (NMuMG), activates the mitogenic extracellular signal-regulated kinase (ERK) pathway, enhanced clonal cell growth and exerts profound effects on proteases secretion. The effect on proteases expression seems to be specific for urokinase-type plasminogen activator and metalloproteinase-9 (MMP-9) because no modulation in MMP-2 and MMP-3 production could be detected. In addition, our experiments demonstrated that PKC $\zeta$  overexpression markedly altered the adhesive, spreading, and migratory abilities of NMuMG cells. The overexpression of this enzyme was not sufficient to confer an anchorage-independent growth capacity. An extensive mutational analysis of PKC $\zeta$  revealed that the effects observed in NMuMG cells were strictly dependent on the kinase (catalytic) domain of the enzyme. Taken together, these results suggest that in mammary cells PKC $\zeta$  modulates several of the critical events involved in tumor development and dissemination through the activation of mitogen activated protein kinase (MAPK)/ERK pathway. © 2004 Wiley-Liss, Inc.

**Key words:** PKC $\zeta$ ; metalloproteinases; urokinase; ERK1/2 MAPK pathway

## INTRODUCTION

Mammary epithelial cells undergo several changes during mammary gland growth and involution cycles [1]. These events require the activation of mitogenic and/or apoptotic signals, the expression of extracellular matrix molecules (ECM) and also the degradation of this ECM molecules. ECM degradation is dependent on proteolytic enzymes such as matrix metalloproteinases (MMPs) [2] and urokinase-type plasminogen activator (uPA) [3]. The alteration of the signaling pathways involved in the control of all these molecules may contribute to malignant transformation. An altered expression of some protein kinase C (PKC) isoforms, including PKC $\alpha$ ,  $\delta$ ,  $\epsilon$ , and  $\zeta$ , has been associated with an increase in protease levels, found during mammary gland differentiation and involution [4].

PKC comprises a family of at least ten phospholipid-dependent serine-threonine kinases that play central roles in signal transduction, governing gene expression, proliferation, apoptosis, differentiation, and malignant transformation [5–7]. PKC isozymes are grouped into three subclasses: the "classical" (PKC $\alpha$ ,  $\beta$ , and  $\gamma$ ), which can be stimulated by Ca<sup>2+</sup> and diacylglycerol or phorbol esters; the "novel" (PKC $\delta$ ,  $\epsilon$ ,  $\eta$ , and  $\theta$ ), which can be activated by diacylglycerol or phorbol esters but are Ca<sup>2+</sup> independent, and the "atypical" (PKC $\zeta$  and  $\lambda/\iota$ ), which

are unresponsive to Ca<sup>2+</sup>, diacylglycerol, and phorbol esters. PKCs possess a regulatory N-terminal domain and a catalytic C-terminal (kinase) domain, presenting conserved regions (C1 to C4) and variable regions (V1 to V5) with specific functional roles. A distinctive feature of the atypical PKC $\zeta$  is that its regulatory region possesses a single phorbol ester unresponsive C1 domain instead of two. It also lacks a C2 domain, which is involved in calcium and/or phospholipid recognition in classical and novel PKCs. The regulatory regions of PKC isozymes play an essential role in allosteric modulation and intracellular targeting, though in some cases, regulatory regions have been shown to trigger biological

Abbreviations: ECM, extracellular matrix; MMPs, matrix metalloproteinases; uPA, urokinase-type plasminogen activator; PKC, protein kinase C; MEK, mitogen-activated protein kinase/ERK kinase; ERK, extracellular signal-regulated kinase; MEM, minimum essential medium; FCS, fetal calf serum; PBS, phosphate buffered saline; SDS, sodium dodecyl sulfate; PAGE, polyacrylamide gel electrophoresis; CM, conditioned media; PAI, plasminogen activator inhibitor; MAPK, mitogen activated protein kinase.

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responses, as shown for the PKC $\delta$  regulatory domain in breast cancer cells [7,8]. Individual PKC isozymes exhibit different tissue distribution, subcellular localization, and biochemical properties, an indication that they play specialized biological roles [9–11].

Deregulation of PKC isozymes has been often associated with malignant transformation and metastatic dissemination [10]. There is strong evidence for the involvement of PKC isozymes in breast cancer progression. Indeed, PKC $\alpha$  overexpression correlates with the lack of estrogen receptor expression in mammary tumor cells, an indicator of poor prognosis [12,13]. In addition, several classical and novel PKC isoforms have been associated with an enhanced expression of MMPs and uPA [14–16], enzymes that play crucial roles in tissue remodeling as well as in cancer initiation and progression [17]. Furthermore, increase of uPA activity functionally correlates with highly aggressive mammary cancers [18]. On the contrary, very little information is available regarding the functionality of the atypical PKC $\zeta$  in the modulation of proteolytic enzymes [19].

Upregulation of PKC $\zeta$  has been detected in various cancer models, including bladder carcinomas [20], hepatocellular carcinomas [21], prostate cancer [22], and melanoma metastasis [23]. Moreover, some studies suggest that atypical PKCs play a functional role in tumor dissemination, and PKC $\zeta$  has been involved in the control of cell motility in pancreatic adenocarcinoma and melanoma cells [24,25]. Recently, it has been reported that PKC $\zeta$  is overexpressed in breast carcinomas [26], though the functional implications of this phenomenon are not known.

The main goal of the present work was to analyze whether an aberrantly activated PKC $\zeta$  pathway facilitates the acquisition of phenotypic characteristics associated with malignancy and invasiveness through the regulation of proteolytic enzymes. As an experimental approach we overexpressed this atypical PKC isozyme in NMuMG cells, a well-established model of normal mouse mammary cells that has been widely used to study mammary carcinogenesis [27–29]. Our experiments revealed that PKC $\zeta$  activates the mitogen-activated protein kinase (MEK)/extracellular signal-regulated kinase (ERK) pathway, leading to profound effects on the ability of NMuMG cells to proliferate, adhere, migrate, and secrete proteases. These findings were substantiated by an extensive mutational analysis of PKC $\zeta$  which revealed that the kinase domain is essential for such effects.

## MATERIALS AND METHODS

### Reagents and Antibodies

Media for cell culture (MEM), agarose, geneticin (G418), and the transfection reagent Lipofectamine Plus were from Life Technologies (Rockville, MD).

Fetal calf serum (FCS) was from GEN (Buenos Aires, Argentina). Plasminogen was purchased to Chromogenix (Molndal, Sweden). Human urokinase was a gift from Serono (Buenos Aires, Argentina). Triton X-100 was obtained from J.T. Baker (Phillipsburg, NJ). Gelatin,  $\alpha$ -casein, PD98059, and acrylamide were from Sigma Co. (St. Louis, MO). All other reagents for polyacrylamide gel electrophoresis and zymography were obtained from Bio-Rad (Richmond, CA). The anti-PKC $\epsilon$  (anti- $\epsilon$ -tag) antibody was obtained from Life Technologies. Antitotal-ERK, anti-phospho-ERK, anti-PKCs ( $\alpha$ ,  $\beta$ ,  $\delta$ , and  $\zeta$ ), and antiactin antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Horseradish peroxidase conjugated antirabbit and antimouse antibodies, Hybond-P membranes and chemiluminescence reagents (ECL) were from Amersham (Aylesbury, UK).

### Cell Line

NMuMG cells, a normal immortalized mammary cell line derived from NAMRU mice [30] were used for the stable transfection assays. Cells were maintained in MEM supplemented with 10% FCS and 80  $\mu$ g/mL gentamycin in humidified atmosphere with 5% CO<sub>2</sub>.

### Expression Vectors, Transfection, and Selection

For the generation of stable transfectants, NMuMG cells were transfected using Lipofectamine Plus with 5  $\mu$ g of expression vectors encoding for wild-type PKC $\zeta$  or various PKC $\zeta$  deletion mutants (PS-cat, Zn-cat, V3-cat, and REG) in pCR3 $\epsilon$  vectors. The generation of these PKC $\zeta$  mutants is described elsewhere [25]. All the expression constructs possess an  $\epsilon$ -tag insert sequence that encodes for a KGFSYFGEDLMP peptide, derived from the last 12 amino acids of PKC $\epsilon$  [31]. The  $\epsilon$ -tagged proteins can be readily detected with a commercially available antibody specific for the  $\epsilon$ -peptide. NMuMG cells transfected with the empty vector (pCR3 $\epsilon$ ) were used as controls (NMuMG-vector). Forty-eight hours after transfection, cells were selected by adding 500  $\mu$ g/mL of G418. After selection, approximately forty G418-resistant clones corresponding to each transfectant were pooled to avoid clonal variations. Transfection assays were carried out in at least two independent experiments for PKC $\zeta$  mutants and three times for PKC $\zeta$  wt and the vector alone. Transfected cell lines were maintained for 10–12 passages before use.

### Western Blot

Semiconfluent monolayers were washed twice with ice-cold phosphate buffered saline (PBS) and then lysed with 1% Triton X-100 in PBS by scraping with a teflon scrapper. Samples were denatured by boiling in sample buffer with 5%  $\beta$ -mercaptoethanol and run in 9% sodium dodecyl sulfate (SDS)-polyacrylamide gel electrophoresis (PAGE). Gels

were blotted to Hybond-P membranes. Membranes were incubated for 1 h with PBS containing 5% skim-milk and 0.1% Tween-20. Membranes were incubated with the first antibodies overnight at 4°C, and then with a secondary antibody coupled to horseradish peroxidase. Detection was performed by chemiluminescence. Bands were quantified by scanning on a digital GS-700 densitometer and the resulting data processed with Molecular Analyst software (Bio-Rad).

#### Cell Proliferation

Doubling time was determined by assessing cell number during the exponential growth phase of unsynchronized monolayer cultures. Briefly,  $1 \times 10^5$  cells were seeded onto 60 mm Petri dishes in MEM supplemented with 10% FCS, 80  $\mu\text{g}/\text{mL}$  gentamycin, and 250  $\mu\text{g}/\text{mL}$  G418. At different times after seeding, cells from triplicate wells were washed twice with PBS, trypsinized, and counted using a hemocytometer and trypan blue exclusion.

#### Clonogenic Assay

Clonal cell growth was studied as previously described [32]. Briefly, single-cell suspensions containing 500–1000 cells were seeded onto 60 mm plastic dishes in MEM with 10% of FCS either in the presence or absence of 50  $\mu\text{M}$  PD98059. Medium was changed every 72 h. After 10 d of culture, cells were washed, fixed with methanol, and stained with hematoxylin. The number of colonies was determined using an inverted microscope. Plating efficiency was defined as the percentage of cells able to grow as colonies having more than ten cells.

#### Anchorage-Independent Growth

For soft agar assays, 60 mm Petri dishes were prepared with 2 mL base feeder layer of 0.6% agar in complete medium and a semisolid top layer (0.4% agar) containing log phase growing monodispersed cells ( $10^5$  cells/dish). Fifteen days after seeding, cultures were fixed by adding 10% formaldehyde in PBS, and the number of colonies was determined using an inverted microscope. Colony efficiency was determined as the percentage of cells able to form colonies in soft agar.

#### Apoptosis Susceptibility

Subconfluent monolayers growing in 96-well plates were extensively washed with PBS and incubated with MEM without FCS for 48 h or treated with 0.5–2  $\mu\text{M}$  of doxorubicin for 2 h. In this last case, the drug was removed, and the cells were further incubated in MEM plus 10% FCS for an additional 24 h. Cell viability was evaluated with an MTS assay (Celltiter 96<sup>TM</sup> Nonradioactive Proliferation Assay, Promega) as described by the manufacturer.

#### Adhesion Assay

Suspensions of  $4 \times 10^5$  cells were seeded onto 35 mm Petri dishes in MEM supplemented with 10% FCS and incubated at 37°C. After 90 minutes the medium was removed and adherent cells were washed twice with PBS, trypsinized, and counted using a hemocytometer. Determinations were carried out by triplicate. Adhesion efficiency was expressed as the percentage of the total number of cells seeded.

#### Spreading Assay

Suspensions containing  $4 \times 10^5$  cells were seeded onto 35 mm Petri dishes in MEM supplemented with 10% FCS. Cells were allowed to adhere, and after 60 min nonadherent cells were removed by washing with PBS. The remaining cells were fixed with 4% formaldehyde. The percentage of spread cells was determined by scoring 200–300 cells under a phase-contrast microscope (Nikon, Tokyo, Japan). Data was collected by random observation in triplicate, as described previously by Aguirre Ghiso et al. [33].

#### Wound Migration Assay

Migration was studied as previously described [32]. Briefly, wounds of approximately 400  $\mu\text{m}$  width were made in subconfluent monolayers of the different transfectant cultures. Cells were then allowed to migrate into the cell-free area for a period of 12 h. The same spot was photographed at different times and the migratory area was analyzed using a digital densitometer GS-700 (Bio-Rad). Cell migration was expressed as the percentage of the area occupied by the migratory cells in the original cell-free wounded area.

#### Preparation of Conditioned Media (CM)

Secreted uPA and MMP activities were evaluated in CM. Briefly, semiconfluent cell monolayers growing in 35-mm plastic Petri dishes were extensively washed with PBS. Serum-free medium (1 mL) was then added for 24 h. CM were individually harvested, the remaining monolayers were lysed with 1% Triton-X100-PBS, and cell protein content was determined (Bio-Rad Protein Assay). CM samples were centrifuged (600g, 10 min), aliquoted, and stored at  $-40^\circ\text{C}$ . Samples were used only once after thawing.

#### Quantification of uPA Activity by Radial Caseinolysis

To determine uPA activity, a radial caseinolysis assay was used, as previously described [34]. Briefly, 4 mm wells were punched in the plasminogen-rich casein-agarose gels and 10  $\mu\text{L}$  of CM were seeded. Gels were incubated for 24 h at 37°C in a humidified atmosphere. The diameter of lytic zones was measured, and the areas of degradation were referenced to a standard curve of purified urokinase (0.1–50 IU/mL) and normalized to the original cell culture protein

content [35]. Specificity of uPA activity was determined by blocking its activity with B428 (18  $\mu$ M), a uPA-specific catalytic inhibitor [36].

#### Determination of Plasminogen Activator Inhibitor (PAI) Activity

PAI activity was determined by reverse zymography, as described by Mackay et al. [37]. Briefly, samples of CM were electrophoresed in 10% SDS/PAGE gel co-polymerized with 30 mg/mL of skim dry milk and 2  $\mu$ g/mL plasminogen. Gels were washed with 1% Triton X-100-PBS, rinsed with water and incubated for 2 h in a buffer containing 20 mM Tris-HCl, pH 8.3, 15 mM EDTA, and 1 IU/mL purified urokinase. Gels were stained with Coomassie Brilliant Blue G-250 (Bio-Rad) and PAI activity was visualized as blue bands in a lighter background.

#### Zymography for MMPs

MMPs enzymatic activities were determined on substrate-impregnated gels, as previously described by Aguirre-Ghiso et al. [38] and Simian et al. [39]. Briefly, samples were collected and run on 9% SDS polyacrylamide slab gels containing 1 mg/mL of gelatin (to detect MMP-9 and MMP-2) or  $\alpha$ -casein (to detect MMP-3), under nonreducing conditions. After electrophoresis, gels were washed for 30 min using 2.5% Triton X-100 and subsequently incubated for 48 h at 37°C in a buffer containing 0.25 M Tris-HCl pH 7.4, 1 M NaCl, and 25 mM CaCl<sub>2</sub>. Nonspecific activity was detected with gels incubated in the same buffer solution but supplemented with 40 mM EDTA. After incubation, gels were fixed and stained with 0.5% Coomassie Brilliant Blue G-250 in methanol/acetic acid/H<sub>2</sub>O (30:10:60). Gelatinolytic bands were measured using a digital densitometer GS-700. Data were expressed as arbitrary units and normalized to the values observed in control cells.

## RESULTS

#### Overexpression of PKC $\zeta$ in NMuMG Cells

In order to investigate whether the overexpression of the atypical PKC $\zeta$  isozyme modulated mammary cellular properties associated with malignant transformation and/or metastatic dissemination, we used a stable transfection approach, as previously described [25,40]. Overexpression of PKC $\zeta$  was achieved upon transfection of NMuMG cells with the expression vector pCR3 $\epsilon$ -PKC $\zeta$  [25] followed by selection with G418. To avoid clonal variations, G418-resistant clones were pooled 2 wk after transfection. Western blot analysis with the anti- $\epsilon$ -tag antibody revealed a 75 kDa band corresponding to the  $\epsilon$ -tagged PKC $\zeta$  in PKC $\zeta$ -transfected NMuMG cells but not in cells transfected with the vector alone (Figure 1A). PKC $\zeta$  was expressed in a high percentage of cells (>80%), as determined by immuno-staining assays, and expression levels were moderate but similar for all analyzed

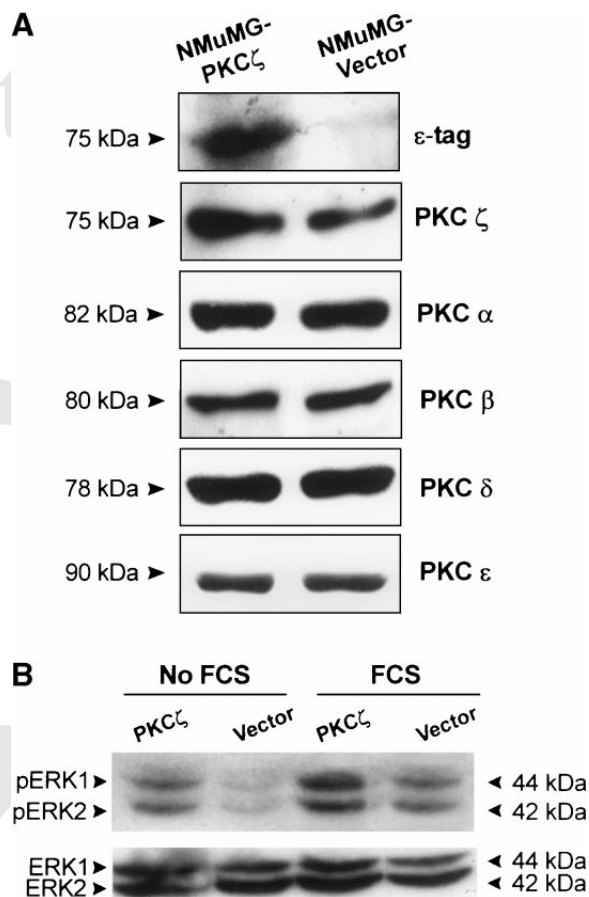


Figure 1. (Panel A) Expression of  $\epsilon$ -tagged protein kinase C- $\zeta$  (PKC $\zeta$ ) in NMuMG cells. Whole cell lysates prepared from NMuMG cells transfected with either pCR3 $\epsilon$ -PKC $\zeta$  or pCR3 $\epsilon$  constructs were resolved on 10% sodium dodecyl sulfate (SDS)-polyacrylamide gel electrophoresis (PAGE) and blotted with an anti- $\epsilon$ -tag antibody or with antibodies against individual PKC isozymes (50  $\mu$ g/lane). (Panel B) Activation of extracellular signal-regulated kinase (ERK)1/2 MAPK by PKC $\zeta$  overexpression in NMuMG cells. Overnight starved cells were treated with 10% FCS (20 min) and subjected to Western blot analysis (40  $\mu$ g/lane) with specific phospho-42/44 ERK or total anti-ERK antibodies. Results are representative of three independent experiments.

cells (data not shown). Using this stable transfection approach, we achieved a 2.8-fold increase in PKC $\zeta$  levels, relative to control transfected cells (Figure 1A, second panel). Overexpression of PKC $\zeta$  did not alter significantly the expression of other PKC isozymes, present in NMuMG cells, as previously described in other cell models [40] (Figure 1A). All pooled cell lines obtained after separated transfection experiments showed similar expression levels and biological properties described below.

#### Evaluation of p42/p44 Mitogen Activated Protein Kinase (MAPK) Pathway Activation

Atypical PKCs activate various intracellular signaling cascades controlling mitogenesis, including the Raf-1/MEK/ERK cascade [41–44]. In order to analyze whether the ERK pathway was indeed activated in

the NMuMG-PKC $\zeta$  stable transfectants, we determined the levels of phosphorylated (active) and total p44/p42 ERK MAPK by Western blot. As shown in Figure 1B, PKC $\zeta$ -expressing cells show increased basal phospho-ERK levels relative to control cells (Figure 1B). Short treatments (20 min) with FCS caused a marked increase in phospho-ERK levels in PKC $\zeta$  cells. The same treatment induced only a moderate activation of ERK in control cells. No changes were observed in total ERK levels (Figure 1B).

#### Effect of PKC $\zeta$ on Proliferation and Survival of NMuMG Cells

Cell counting at early times after plating (5–24 h) revealed a significantly higher cell number in PKC $\zeta$  transfectants compared to control (vector-transfected) cells (Figure 2A). The increase in cell number was blocked by PD98059, a MEK inhibitor (data not shown). Despite the higher number of cells in PKC $\zeta$  transfectants, analysis of growth rate between 48 and 96 h did not show a significant difference in the population doubling time ( $17.8 \pm 3.7$  vs.  $19.1 \pm 2.9$  h, for PKC $\zeta$  and control cells, respectively) (Figure 2B). Then, we explored whether PKC $\zeta$  overexpression conferred a growth advantage under more stringent conditions such as clonal cell growth. As shown in Figure 2C, PKC $\zeta$  transfectants show a higher clonogenic ability than control cells. Again, this capacity was reverted by PD98059 (Figure 2C).

The ability to grow independently of substrate attachment represents a hallmark of malignant cell transformation. Thus, we next assayed whether PKC $\zeta$  overexpression conferred NMuMG cells with anchorage-independent growth capacity in soft agar assay. It was found that neither NMuMG-PKC $\zeta$  nor the vector-transfected control cells were able to form colonies in soft agar.

Transformed cell populations tend to expand in number not only by increasing proliferation rate but also by evading apoptosis [45]. The overexpression of PKC $\zeta$  did not confer any survival advantage when cells were deprived of serum or exposed to the cytotoxic drug doxorubicin. In this regard, NMuMG-PKC $\zeta$  cells presented a similar doxorubicin inhibitory concentration 50 than control transfectants ( $1.4 \pm 0.3$  vs.  $1.8 \pm 0.4$   $\mu$ M, respectively).

#### Role of PKC $\zeta$ in Adhesion, Migration, and Spreading Ability

The increase in cell number in NMuMG-PKC $\zeta$  transfectants observed at early times after plating prompted us to examine whether this could be a consequence of changes in adhesion. Indeed, it was found that PKC $\zeta$  transfectants presented an enhanced adhesive ability relative to control (vector-transfected) cells. A quantitative analysis revealed that 90 min after plating a significant higher number of PKC $\zeta$ -transfected cells adhered to the plastic surface, as compared to control (Figure 3A). The rate

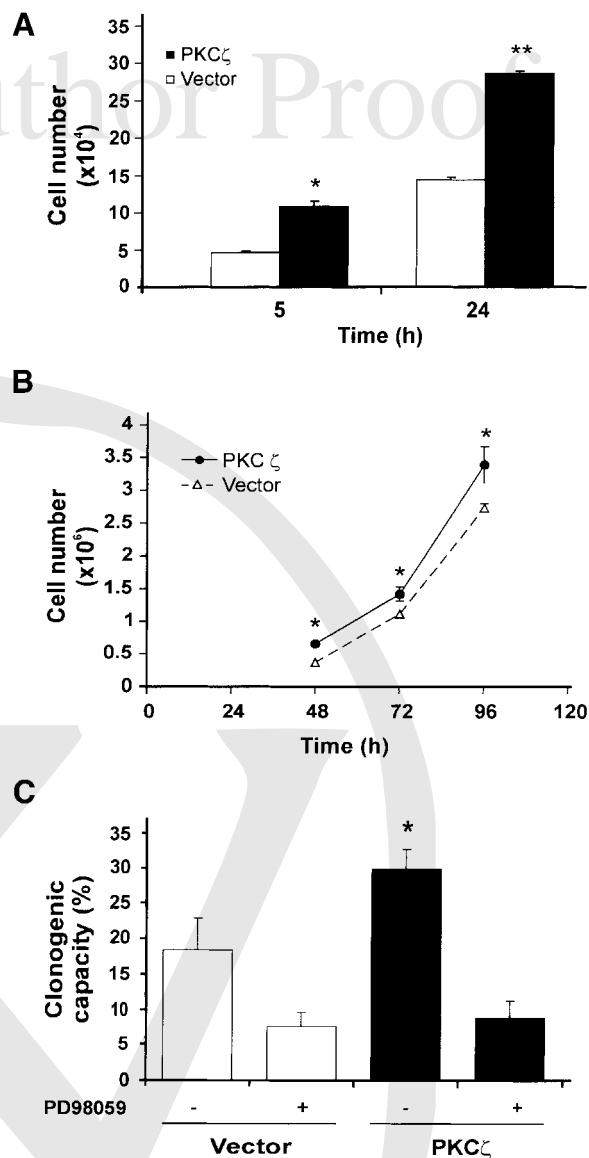
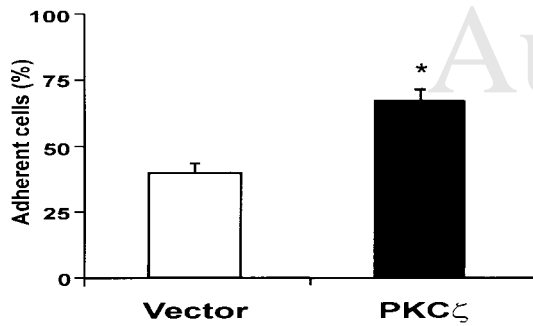


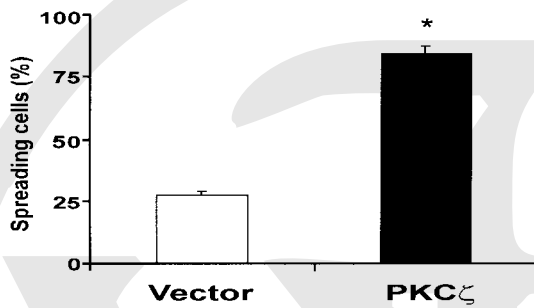
Figure 2. (Panel A) Effect of PKC $\zeta$  on cell growth. Cell number was assessed in unsynchronized NMuMG cultures at different times. At least three independent experiments with similar results were performed. \* $P < 0.05$ , \*\* $P < 0.01$  versus NMuMG-vector cells (Student's  $t$ -test). Each data point represents the mean  $\pm$  SD of triplicate determinations. (Panel B) Effect of PKC $\zeta$  on population doubling time. Cell number was assessed at different times during the exponential growth phase of unsynchronized NMuMG cultures. Results are representative of three independent experiments. \* $P < 0.05$  versus NMuMG-vector cells (Student's  $t$ -test). (Panel C) Clonogenic capacity. Clonal cell growth was studied as previously described in "Materials and Methods" in the presence or absence of PD98059 (50  $\mu$ M). Results are representative of three independent experiments. \* $P < 0.05$  versus NMuMG-vector cells (Student's  $t$ -test).

of adhesion was time-dependent, and 100% of NMuMG-PKC $\zeta$  cells were adhered at 5 h incubation while vector-transfected cells did so only after 10 h incubation (data not shown). The spreading behavior was also markedly enhanced in PKC $\zeta$  overexpressing cells (Figure 3B).

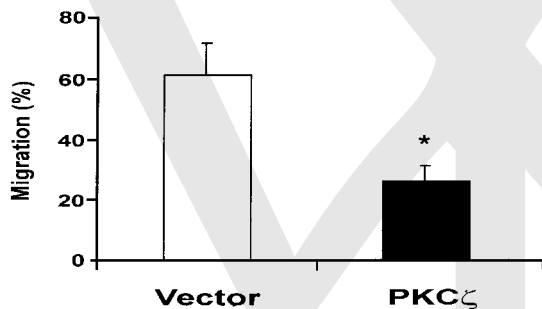
A



B



C

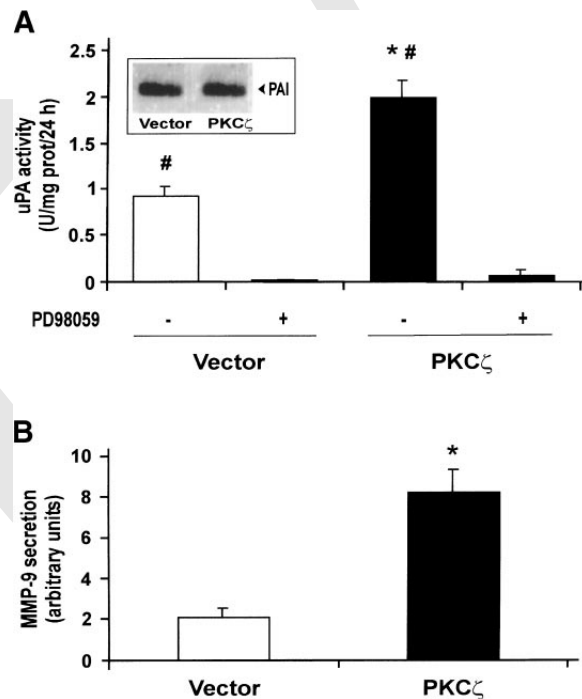


**Figure 3.** Effect of PKC $\zeta$  on adhesion, spreading, and migration. (Panel A) Adhesion studies. Control and PKC $\zeta$  transfectants were assayed for their adhesive capacity as described in "Materials and Methods." The rate of adhesion was expressed as the percentage of the total number of cells seeded. Data expressed as the mean  $\pm$  SD are representative of three independent experiments. \* $P < 0.05$  versus NMuMG-vector cells (Student's *t*-test). (Panel B) Spreading ability. Control and PKC $\zeta$  transfectants were assayed for their spreading ability, as described in "Materials and Methods." The percentage of spreading cells was determined by scoring cells under a phase-contrast microscope. Data were collected by random observation. Data expressed as the mean  $\pm$  SD are representative of three independent experiments \* $P < 0.01$  versus NMuMG-vector cells (Student's *t*-test). (Panel C) Migration assay. Subconfluent monolayers were "wounded" at time 0 and cells were allowed to migrate into the cell free area for 12 h. The same spot was photographed at different times and cell migration was quantified by calculating the percentage of the area occupied by cells that migrated into the original cell-free wounded area. Data expressed as the mean  $\pm$  SD are representative of three independent experiments \* $P < 0.01$  versus NMuMG-vector cells. Student's *t*-test.

In order to analyze whether PKC $\zeta$  overexpression modulates the migratory properties of NMuMG cells, we performed a "wound" assay. Figure 3C shows that, PKC $\zeta$ -transfected cells showed a significantly lower migratory rate than control cells.

#### Role of PKC $\zeta$ in Proteases Secretion

We next analyzed whether PKC $\zeta$  affects the secretion of proteases. Previous reports from our laboratory have shown that phorbol ester-responsive PKC isoforms positively regulate both uPA and MMPs expression in normal and tumor cells [16,29]. Interestingly, we found that overexpression of PKC $\zeta$  in NMuMG mammary cells also induced a significant increase in uPA and MMP-9 secreted activities (Figure 4A and B). In contrast, no changes in MMP-2 and MMP-3 (stromelysin) activities were observed (see Figure 7C below). PAI activity was also determined by reverse zymography assay in the CM of both control and PKC $\zeta$  transfectants. As shown in



**Figure 4.** Effect of PKC $\zeta$  on the secretion of proteolytic enzymes. (Panel A) Quantification of urokinase-type plasminogen activator (uPA) activity by radial caseinolysis. uPA secreted activity was quantified by radial caseinolysis in conditioned media (CM) from PKC $\zeta$  transfected or control cell cultures, either in the presence or absence of PD98059 (50  $\mu$ M). \* $P < 0.01$  versus NMuMG-vector cells, # $P < 0.01$  versus same cell treated with PD98059 (Student's *t*-test). Data are expressed as the mean  $\pm$  SD of triplicate determinations and are representative of at least three independent experiments. Inset: Secreted plasminogen activator inhibitor (PAI) activities, detected by reverse zymography in the same CM. Two additional experiments gave similar results. (Panel B) Quantification of MMP-9 activity by zymography. MMP-9 activity was analyzed as described in "Materials and Methods." Proteolytic activity bands were measured with a digital densitometer, and data were expressed as arbitrary units. \* $P < 0.05$  versus NMuMG-vector cells (Student's *t*-test). Data are expressed as the mean  $\pm$  SD of triplicate determinations and are representative of at least three independent experiments.

Figure 4A (inset), we did not observe measurable differences in PAI activity. These results suggest that PKC $\zeta$  modulation of uPA activity cannot be attributed to a regulation of its inhibitory molecule. Remarkably, the inhibition of MEK with PD98059 impaired both basal uPA secretion as well as the enhanced activity observed in PKC $\zeta$  overexpressors (Figure 4A) indicating that active signaling through PKC $\zeta$  and the MEK/ERK pathway is required for uPA production. Similar results were also observed for MMP-9 (data not shown).

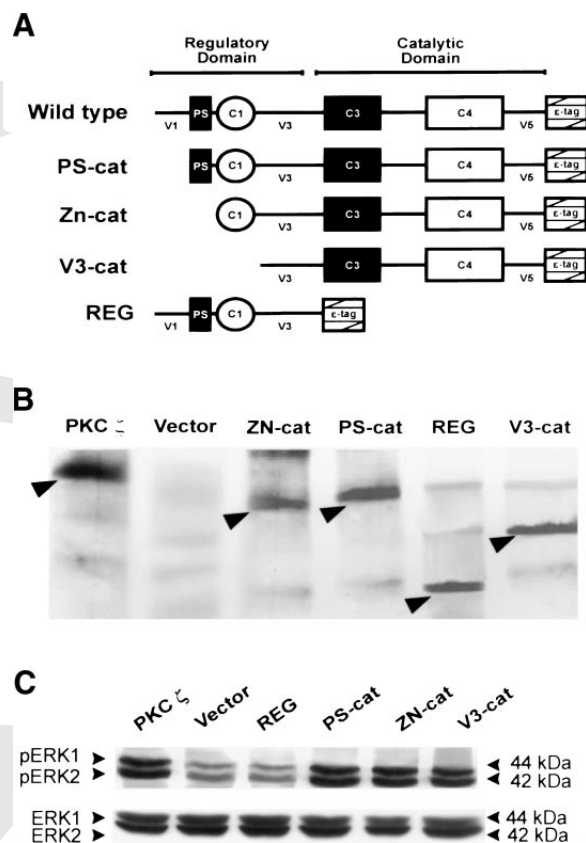
#### Structure-Function Analysis of PKC $\zeta$ in NMuMG Mammary Cells

An important issue in PKC-mediated function is the elucidation of the roles of individual domains in triggering biological effects. This is important because it has been reported that PKC isozymes can signal independently of their catalytic (kinase) activity [25,46–48]. We used a series of deletion mutants comprising different regions of the molecule (Figure 5A). The N-terminal deletion mutants would allow for the assessment of the potential roles of the V1, pseudosubstrate and C1 domains. The constructs, which were epitope-tagged ( $\epsilon$ -tag) to facilitate the comparison of the expression levels, were transfected into NMuMG cells followed by G418 selection. The mobilities of the truncated mutants in SDS-PAGE were consistent with their predicted molecular weights and comparable levels of expression were observed for all mutants (Figure 5B). Only those cell lines expressing an intact PKC $\zeta$  catalytic (kinase) domain showed increased phospho-ERK1/2 levels. Expression of the regulatory domain alone did not affect phospho-ERK1/2 levels, as they were similar to those observed in control (vector-transfected) cells (Figure 5C). Total ERK was similar in all cases (Figure 5C).

In the next set of experiments, we analyzed the effect of the PKC $\zeta$  mutants on adhesion, spreading, and migration. As shown in Figure 6, only those transfectants expressing an intact catalytic (kinase) present similar phenotypes as those observed in wild-type (full-length) PKC $\zeta$  transfectants. Indeed, the adhesion and spreading assays showed that the presence of an intact catalytic domain was required to enhance the adhesive properties of NMuMG cells (Figure 6A and B). Likewise, only those mutants having an intact catalytic domain were capable of inhibiting migration in the wound assay (Figure 6C).

#### Effect of PKC $\zeta$ Mutants on the Production of uPA and MMPs

Finally, we tested whether the PKC $\zeta$  mutants induce changes in the production of proteases. NMuMG cells expressing PKC $\zeta$  mutants with an intact kinase domain secreted more uPA than control cells (twofold to fourfold increase) (Figure 7A). This increase was more evident in ZN-cat expressing cells,

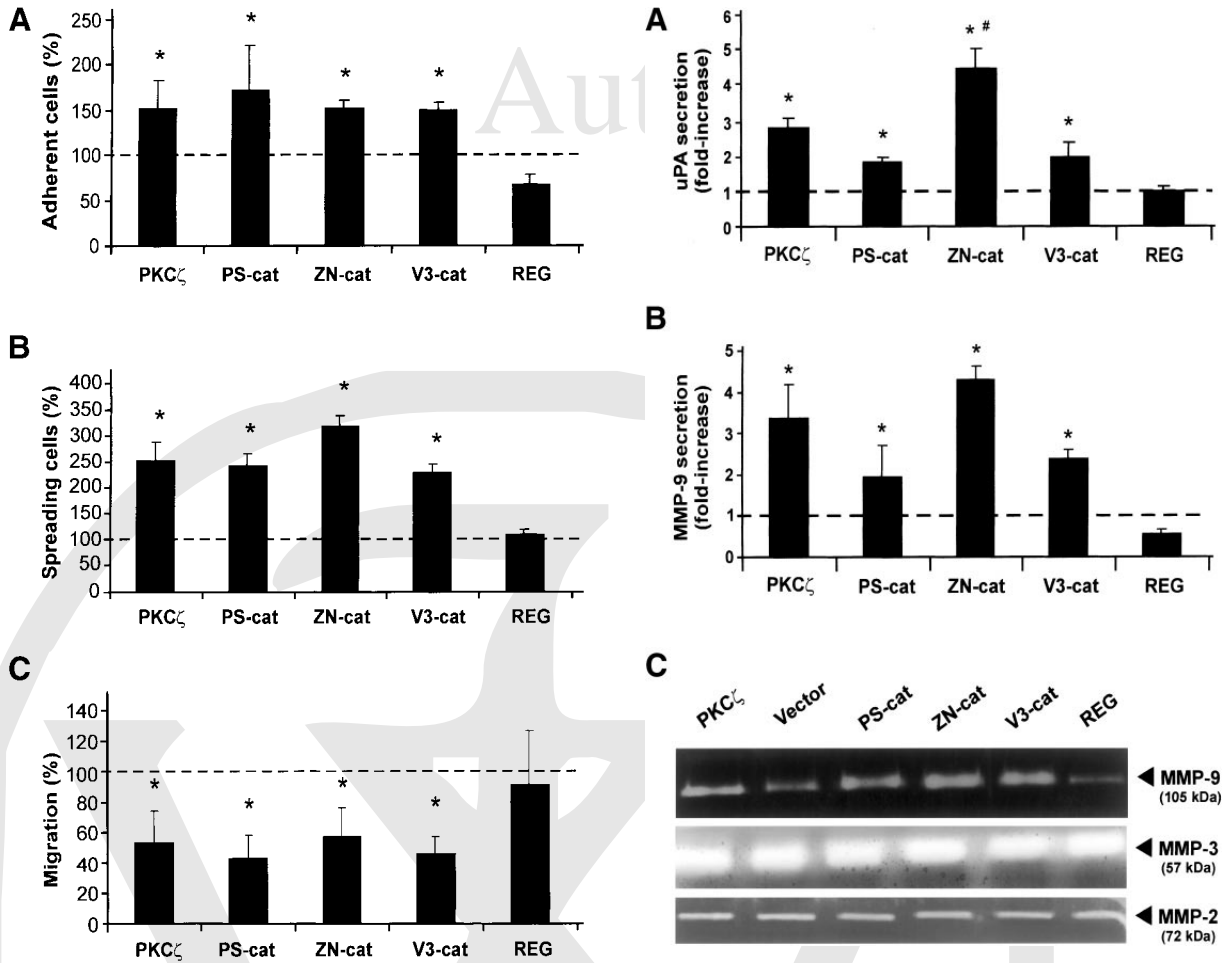


**Figure 5.** Expression of PKC $\zeta$  mutants in NMuMG cells. (Panel A) Schematic representation of PKC $\zeta$  mutants. The pseudosubstrate region (PS), cysteine-rich (Zn-finger) domain (C1), variable regions V1, V3, and V5, and the conserved region C3 are indicated. (Panel B) Expression of  $\epsilon$ -tagged PKC $\zeta$  mutants in NMuMG cells. Western blot analysis was performed with lysates of NMuMG cells stably transfected with each mutant (60  $\mu$ g/lane). The arrows indicate the position of each mutant protein. (Panel C) Modulation of ERK1/2 MAPKs activation by PKC $\zeta$  mutants. Transfectants were lysed and subjected to a Western blot analysis (40  $\mu$ g/lane) with specific antibodies for phospho-p42/44 ERK or total-ERK. Three independent experiments gave similar results.

although the reasons for these differences are not known. Again, the mutant without the catalytic domain (REG), behaved like control cells. In all cases uPA activity could be inhibited by B428 and was not detected in plasminogen-free substrates (data not shown). Gelatin zymograms revealed that MMP-9 was significantly increased in cells overexpressing either wild-type PKC $\zeta$  or the kinase active PKC $\zeta$  mutants (Figure 7B and C). The effect was specific for MMP-9, as no changes were observed in MMP-2 and MMP-3 secreted activity (Figure 7C). No evidence for gelatinolytic or caseinolytic activity was detected in gels incubated in EDTA-containing buffers, confirming that the observed activities corresponded to metalloproteinases (data not shown).

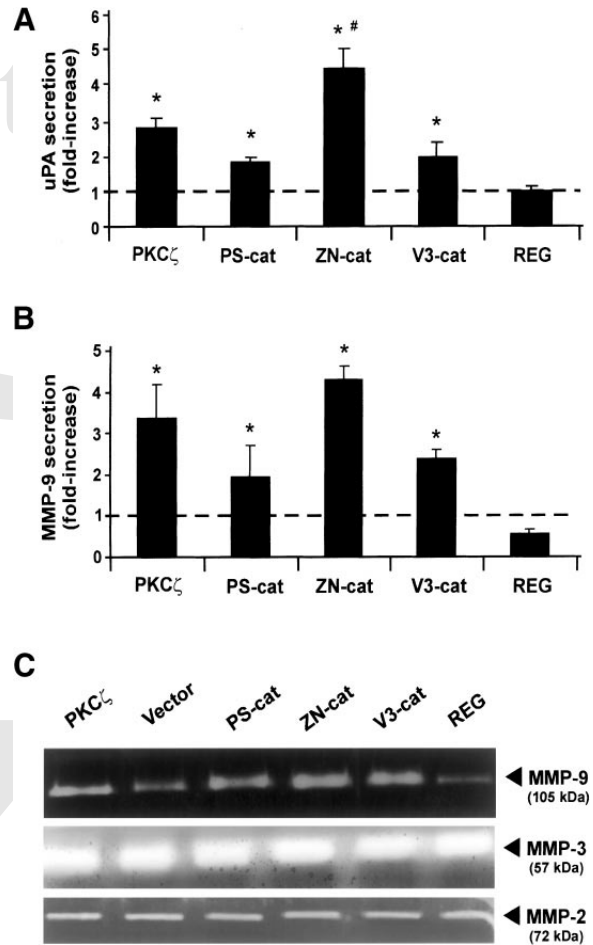
#### DISCUSSION

In the present study, we showed that the overexpression of the atypical PKC $\zeta$  isoform modulates



**Figure 6.** Effect of truncated PKC $\zeta$  mutants on adhesion, spreading, and migration. (Panel A) Adhesion studies. PKC $\zeta$  mutants were assayed for their adhesive capacity as described in "Materials and Methods." Values were compared to those obtained with the vector-transfected cells, which were arbitrarily set as 100% (dotted line). \* $P < 0.05$  versus NMuMG-vector cells (ANOVA test). Data expressed as the mean  $\pm$  SD are representative of three independent experiments. (Panel B) Spreading ability. PKC $\zeta$  mutants were assayed for their spreading ability as described in "Materials and Methods." Values were compared to those obtained with the vector-transfected cells, which were arbitrarily set as 100% (dotted line). \* $P < 0.01$  versus NMuMG-vector cells (ANOVA test). Data expressed as the mean  $\pm$  SD are representative of three independent experiments. (Panel C) Migration assay. Wound migration was studied as previously described in "Materials and Methods." Values were compared to those obtained with the vector-transfected cells, which were arbitrarily set as 100% (dotted line). \* $P < 0.05$  versus NMuMG-vector cells (ANOVA test). Data expressed as the mean  $\pm$  SD are representative three independent experiments.

clonogenicity, cell adhesiveness, motility, and proteases secretion of the immortalized normal murine mammary gland cell line NMuMG through the activation of the ERK/MAPK pathway. In addition, we demonstrated that a functional kinase domain is necessary for these effects. While it has been reported that the overexpression of some classical and novel PKCs confers characteristics associated with malignant transformation to fibroblasts [49] or to mammary epithelial cells [8,29], there have been no



**Figure 7.** Effect of truncated PKC $\zeta$  mutants on the secretion of proteolytic enzymes. (Panel A) Quantification uPA activity by radial caseinolysis. uPA secreted activity was quantified by radial caseinolysis as described in "Materials and Methods." Values were referred to those secreted by vector-transfected cells (dotted line). \* $P < 0.05$  versus NMuMG-vector cells (ANOVA test). # $P < 0.05$  versus PKC $\zeta$  cells (Student's *t*-test). Data are expressed as the mean  $\pm$  SD of three determinations and are representative of at least four independent experiments. (Panel B) Quantification MMP-9 activity by zymography. MMP-9 activity was analyzed as described in "Materials and Methods." Proteolytic activity bands were measured with a digital densitometer. Activities were expressed as arbitrary units and normalized to that of vector-transfected cells (dotted line). \* $P < 0.05$  versus NMuMG-vector cells (ANOVA test). Data are expressed as the mean  $\pm$  SD of three determinations and are representative of at least four independent experiments. (Panel C) Zymography for MMP-2, MMP-3, and MMP-9 detection. MMP-9 and MMP-2 activities were assessed by gelatin zymography. MMP-3 activity was analyzed by casein zymography. Results are representative of at least three independent experiments.

studies addressing the role of the atypical PKC $\zeta$  isoform and the functionality of its structural domains as it pertains to mammary cancer initiation and progression.

PKC enzymes are key elements of the mitogenic signaling pathway that modulate the MEK/ERK phosphorylation cascade, culminating with the activation of transcription factors associated with cell cycle regulation [16,29,43]. Contrarily to the classical and novel PKCs, it has been shown that



PKC $\zeta$  activates this cascade by a mechanism independent of c-Raf1 that probably involves a direct interaction with MEK [43,50]. Although NMuMG cells overexpressing PKC $\zeta$  showed a constitutive activation of the ERK/MAPK pathway, we did not find any significant shortening in the population doubling time. In contrast, other studies have reported enhanced proliferation in response to PKC $\zeta$  overexpression in thyroid gland cells [40], suggesting that the effects of atypical PKCs may be cell-type dependent. Nevertheless, it is important to note that when NMuMG cells were grown under more stringent conditions such as clonal cell growth, overexpression of PKC $\zeta$  conferred a significantly higher clonogenic capacity. In agreement with experiment-sin other cell model [51,52], the overexpression of PKC $\zeta$  was not sufficient to confer anchorage-independent growth capacity.

It is known that proteases expression, ECM remodeling, and adhesive and migratory properties play fundamental roles in the normal physiology as well as in the malignant transformation and tumor progression of the mammary gland [17,53]. Furthermore, an altered expression of proteases, a crucial event for cell invasiveness [54,55], has been associated with poor prognosis in breast cancer [18,56]. In this paper, we report that PKC $\zeta$  overexpression enhanced the secretion of uPA and MMP-9, but not MMP-2 and MMP-3, by NMuMG mammary cells. In the same model, we have previously observed an increase of uPA, MMP-2, and MMP-9 secretion as consequence of a classical PKC overexpression [29]. In support of our conclusions, it has been recently reported that PMA-insensitive (atypical) PKC isozymes are key regulators of MMP-9 expression in rat glioma cells [19]. Conversely, our unpublished results show that PKC $\delta$  overexpression plays the opposite role in NMuMG cells, inhibiting the secretion of both uPA and MMP-9. Taken together, these data are consistent with the idea that individual PKC isozymes have different and even opposing functions within the same cell context [9–11].

To further analyze some of the mechanism implicated in the changes induced by PKC $\zeta$  overexpression in adhesion, spreading, and migration, we have studied the expression of some cell-adhesion molecules and ECM components. Our preliminary results indicate that CD44,  $\beta$ 1 integrin, and fibronectin seem not to be modulated by PKC $\zeta$ , contrarily to our own findings in the same cell model overexpressing a classical PKC [29].

Though, at first sight, the reduced ability to migrate of NMuMG-PKC $\zeta$  cells could be interpreted as a contradictory finding with the increased production of specific proteases observed in the same cells, this effect must be analyzed in the context of the complexity of molecular events and cell-cell and cell-environment interactions necessary for the expression of a fully malignant phenotype *in vivo*, which

are probably not included in the usual, but limited, *in vitro* assays. In fact, preliminary experiments showed that NMuMG-PKC $\zeta$  transfectants presented an increase in their capacity to invade a collagen matrix, assay that integrates both migratory and invasive capacities. Wolf et al. [17,57] have also reported apparent contradictions between protease secretion and migration, as they found that even in the absence of pericellular proteolysis, malignant mammary cells could still invade a collagen matrix due to migratory compensation strategies.

Next, an extensive mutational analysis was performed to address the function of the different PKC $\zeta$  structural domains in proteases secretion. We found that the presence of an intact kinase domain was necessary and sufficient to increase uPA and MMP-9 secretion. The modulation of these two different proteolytic enzymes by PKC $\zeta$  and mutants showed almost identical pattern, and only minor changes could be observed. These subtle differences may be related to unknown functional roles of PKC $\zeta$  domains, as described for other PKCs, even though more biochemical and functional analysis are required to explain them.

Although the PS-cat and V3-cat mutants secreted significantly more uPA and MMP-9 than control cells, their protease secretion level was somehow lower than that of the wild-type PKC $\zeta$  transfectants. This suggested that the V1 region may be relevant in the modulation of protease secretion. When this domain is absent (ZN-cat mutants) proteases secretion level is even higher than that shown by wild-type PKC $\zeta$  cells. Neither MMP-2 nor MMP-3 (stromelysin 1) activities displayed any modulation in wild-type or mutant PKC $\zeta$  transfectants, demonstrating the specificity of this atypical isoform on uPA and MMP-9 regulation. We speculate that these differences may be related to issues of localization due to the important roles that discrete domains in the regulatory region of PKCs play in intracellular targeting.

Our results are coincident with those of Sanz-Navarez [25] in the sense that PKC $\zeta$  overexpression inhibits the migration of murine B16 melanoma cells. However, a more detailed analysis of their results revealed that, although PKC $\zeta$  mutants with a functional catalytic domain were also able to inhibit migration, the highest response was attributed to theregulatory domain devoid of enzymatic activity. This discrepancy with our results, clearly showing that the PKC $\zeta$  regulatory domain did not modulate migration, can be attributed to differences in the tissue origin of the cell model employed, as well as to the fact that NMuMG is a normal, though immortalized, cell line.

Our findings unambiguously revealed that the kinase domain of this atypical PKC $\zeta$  is essential for the analyzed biological effects, probably through the activation of the MEK/ERK cascade. However, the activation and/or inhibition of other pathways by

PKC $\zeta$  signaling is not addressed yet and our results do not rule out this possibility.

In summary, the overexpression of PKC $\zeta$  or its mutants with kinase activity in mammary cells induced an enhanced expression of specific proteases, together with alterations in the adhesive, spreading, and migratory potential, and an increased capability to grow under very stringent conditions. Our results suggest that a PKC $\zeta$ -dependent pathway could be critical for late events leading to an enhanced invasive and metastatic phenotype, thus revealing a novel rationale for targeting this atypical PKC for the control of breast cancer cell dissemination.

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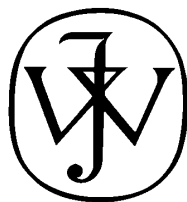
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