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PKC δ is Required for Survival of Cells Expressing Activated p21^{Ras}*

Shuhua Xia, Lora W. Forman, and Douglas V. Faller

Cancer Research Center, Boston University School of Medicine, 715 Albany Street, Boston, MA 02118, USA

Abstract

Inhibition of PKC activity in transformed cells and tumor cells containing activated p21^{Ras} results in apoptosis. To investigate the pro-apoptotic pathway induced by the p21^{Ras} oncoprotein, we first identified the specific PKC isozyme necessary to prevent apoptosis in the presence of activated p21^{Ras}. Dominant-negative mutants of PKC, siRNA vectors, and PKC isozyme-specific chemical inhibitors directed against the PKC δ isozyme demonstrated that PKC δ plays a critical role in p21^{Ras}-mediated apoptosis. An activating p21^{Ras} mutation, or activation of the PI₃K Ras effector pathway, increased the levels of PKC δ protein and activity in cells, whereas inhibition of p21^{Ras} activity decreased the expression of PKC δ protein. Activation of the Akt survival pathway by oncogenic Ras required PKC δ activity. Akt activity was dramatically decreased after PKC δ suppression in cells containing activated p21^{Ras}. Conversely, constitutively-activated Akt rescued cells from apoptosis induced by PKC δ inhibition. Collectively, these findings demonstrate that p21^{Ras} through its downstream effector PI₃K, induces PKC δ expression and that this increase in PKC δ activity, acting through Akt, is required for cell survival. The p21^{Ras} effector molecule responsible for the initiation of the apoptotic signal after suppression of PKC δ activity was also determined to be PI₃K. PI₃K (p110-CAAX), was sufficient for induction of apoptosis after PKC δ inhibition. Thus, the same p21^{Ras} effector, PI₃K, is responsible for delivering both a pro-apoptotic signal, and a survival signal, the latter being mediated by PKC δ and Akt. Selective suppression of PKC δ activity and consequent induction of apoptosis is a potential strategy for targeting of tumor cells containing an activated p21^{Ras}.

The Ras oncogene family is among the most commonly mutated group of genes in human cancer. Its protein products code for three closely-related, p21^{Ras} proteins including H-ras, K-ras and N-ras. p21^{Ras} proteins are localized in the inner plasma membrane, bind GDP and GTP and possess an intrinsic GTPase activity. p21^{Ras} proteins function as plasma membrane-bound guanine nucleotide binding proteins and act as molecular switches, thereby regulating signal transduction pathways for hormones, growth factors and cytokine receptors (1). Several downstream effector proteins of p21^{Ras} have been identified which bind preferentially to p21^{Ras} in the GTP-bound state, including Raf, phosphoinositide 3-OH kinase (PI₃K), and a family of GDP-GTP exchange factors for the Ral small GTPases (Ral-GDS). Raf proteins, which are proto-oncogene-encoded serine/threonine kinases, activate the MEK-ERK signaling pathway. PI₃K activation results in the activation of the anti-apoptotic serine/threonine kinase Akt, among other molecules. Other p21^{Ras} targets include

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Address correspondence to: Douglas V. Faller, Cancer Research Center, K-701, 715 Albany St. Boston, MA 02118, Tel. 617 638-4173; Fax, 617 638-4176; dfaller@bu.edu.

the GTPase-activating proteins, p120GAP and neurofibromin (2). p21^{Ras} proteins were shown to influence proliferation, differentiation, transformation, and apoptosis by relaying mitogenic and growth signals from the membrane into the cytoplasm and the nucleus.

Specific point mutations localized in codons 12, 13, 59, 61, 63, 116, 117, and 146 can lock the p21^{Ras} protein in the active GTP-bound state and permit stimulation of downstream signaling cascades in the absence of extrinsic p21^{Ras} activation. Ras mutations can be found in human malignancies with an overall frequency of 20%. A particularly high incidence of Ras gene mutations has been reported in malignant tumors of the pancreas (80–90%, K-ras), in colorectal carcinomas (30–60%, K-ras), in non-melanoma skin cancer (30–50%, H-ras), and in hematopoietic neoplasia of myeloid origin (18–30%, K-and N-ras) (3).

In addition to its central involvement in cell proliferation, recent studies indicate that the presence of an activated p21^{Ras} protein sensitizes transformed or malignant cells to apoptotic stimuli (4–9). Various signaling pathways have been proposed for this pro-apoptotic activity. Chou et al (10) reported activated p21^{Ras} can cause apoptosis in transformed murine fibroblast cells through activation of the transcription factor NFκB. Another study suggested that the p21^{Ras}/MAP kinase pathway is involved in Ras-specific apoptosis (11). The latter study also found that activating p21^{Ras} mutations increased colon cancer cell sensitivity to 5-FU-induced apoptosis through the negative regulation of gelsolin expression (12). Our previous studies demonstrated suppression of protein kinase C (PKC) activity in cells expressing activated p21^{Ras} rapidly induces apoptosis via FADD/caspase-8 signaling (9). We also found that reactive oxygen species are necessary as downstream effectors of the Ras-mediated apoptotic response to PKC inhibition (7).

There are at least 12 PKC isoforms that are classified into three subfamilies according to the structure of the N-terminal regulatory domain, which determines their sensitivity to the second messengers Ca²⁺ and diacylglycerol (DAG) (13). Despite the high degree of homology, however, there is a surprising degree of non-redundancy. Thus, individual PKC isoforms mediate different and unique cellular functions in different cell types and different tissues (14). PKCδ belongs to the subfamily of novel isoforms (PKCδ, PKCε, PKCθ and PKCη), which are insensitive to Ca²⁺. PKCδ is widely regarded as having pro-apoptotic properties (15–17). Caspase activation mediates cleavage of PKCδ which results in release of the active catalytic domain (18, 19). In addition, PKCδ activity is known to initiate a number of pro-apoptotic signals, such as increased expression and stability of p53 (20, 21), mitochondrial cytochrome C release (22, 23) and c-Ab1 activation (24). But recent studies have also shown that PKCδ can protect cells against apoptotic stimuli under certain conditions (25). PKCδ has been reported to regulate B-lymphocyte survival (26). Knock-out experiments have shown that PKCδ-deficient mice have a severely deregulated immune system and develop autoimmune disease (27, 28). Thus, PKCδ activation can serve as pro-apoptotic signal, or as survival signal, to determine cell fate.

The present study examines the mechanism of apoptotic signaling induced by the p21^{Ras} oncoprotein. We found that PKCδ plays a critical role in suppressing p21^{Ras}-mediated apoptosis, and selective inhibition of this isozyme initiates apoptosis in cells containing activated p21^{Ras}. Our data further demonstrate that unregulated Ras activity, through activation of the downstream effector PI₃K, up-regulates PKCδ expression and subsequently activates Akt, generating an anti-apoptotic effect and protecting against Ras-mediated apoptosis.

EXPERIMENTAL PROCEDURES

Plasmids and Reagents

The activated Ras effectors, Raf-1 (Raf-22W), PI₃K (p110-CAAX), Rlf (Rlf-CAAX) and Ral-GDS (RalA-28N, dominant-negative) were cloned into the pBabe puro vector (29) and the HRas effector loop mutants, S35, G37 and C40 (double mutations) and H-ras V12 single mutant (V12) were cloned into the pSG5 vector (30). These vectors were kindly provided by C. Counter (Duke University) and J. Downward (MRC, UK). The pEGFP-PKC δ -KR (dominant-negative PKC δ) vector (31) was kindly provided by Dr. D. Kufe (Dana-Farber Cancer Institute). pEF1 α -vAkt and pEF1 α -cAkt were kindly provided by Dr. Geoffrey Cooper (Boston University). The GST-NORE CT and FLAG-MST1 CT vectors were generously provided by Dr. J. Avruch (Massachusetts General Hospital). Chemical inhibitors used in this study specific to PKC isozymes, PI₃K, p21^{Ras} and MAPK are listed in Table 1. All inhibitors were dissolved in dimethyl sulfoxide for use, and their effects were measured relative to dimethyl sulfoxide (vehicle)-treated controls. The concentration of all inhibitors was optimized to produce greater than 90% inhibition of target molecule activity.

Cell culture and treatment

NIH/3T3 and Balb cell lines were obtained from ATCC (American Type Culture Collection, Rockville, MD). NIH/3T3-Ras cells were produced by stable transfection of v-Harvey ras, and selected and maintained in 0.5 mg/ml of Geneticin (Gibco BRL, Gaithersburg, MD). Ki-v-ras-Balb (KBalb) cells were produced by stable infection of Balb with retroviral vector stocks containing v-Kirsten ras, and were selected and maintained with Geneticin. The human pancreatic tumor lines Hs 766s, BxPC-3 and MIA PaCa-2, were obtained from ATCC. All cell lines were maintained in Dulbecco Modified Eagle medium (DMEM) (Gibco BRL) supplemented with 2 mM l-glutamine, 100 U/ml of penicillin, and 100 μ g/ml of streptomycin (Gibco BRL). Media was additionally supplemented with 10% DCS (NIH/3T3, NIH/3T3-Ha-v-Ras, Balb/3T3, Ki-v-ras-Balb, and Balb-myc), or 10% FBS (BxPC-3, MIA PaCa-2). Cells were cultured at 37°C and 5% CO₂.

Cell proliferation assay

Cell proliferation was assessed using an MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assay (Roche, Mannheim, Germany). The number of viable cells growing in a single well on a 96-well microtiter plate was estimated by adding 10 μ l of MTT solution (5 mg/ml in phosphatebuffered saline [PBS]). After 4 h of incubation at 37°C, the stain was diluted with 100 μ l of dimethyl sulfoxide. The optical densities were quantified at a test wavelength of 550 nm and a reference wavelength of 630 nm on a multiwell spectrophotometer.

siRNA knockdown of PKC δ and PKC α

siRNA duplexes for PKC δ (siRNAs) were obtained from Qiagen (Valencia, CA). The siRNA sequences for targeting PKC δ were PKC δ -siRNA-1 (5'-GAUGAAGGAGGCGCUCAGTT-3') and PKC δ -siRNA-2 (5'-GGCUGAGUUCUGGCUGGA-CTT-3') (32). The corresponding scrambled siRNAs were used as negative control. These siRNA sequences were also cloned into the pRNA6.1-Neo vector with a GFP tag according to the manufacturer's instructions (GenScript, Piscataway, NJ). siRNA for PKC α (PKC-PKC α -V6) was purchased from Upstate (Lake Placid, NY). Transfection of siRNA (oligo) was performed using 50 nM PKC δ siRNA, or the same amount of scrambled siRNA and Lipofectamine 2000 (Invitrogen, Carlsbad, CA), according to the manufacturer's instructions. Transfection of plasmid-based siRNA vectors was carried out using the same method. PKC δ protein levels were determined by immunoblot analysis.

Assay of PKC α and PKC δ kinase activities

PKC α and PKC δ activities were measured with an assay kit (Upstate Cell Signaling). After two days of treatment with rottlerin, cells were lysed in 50 mM Tris-HCl, pH 8.0, 150 mM NaCl, 1% Triton X-100, 20 mM MgCl₂, 5 mM EGTA, 1 mM orthovanadate, 50 μ g/ml PSMF and 3 μ g/ml aprotinin. PKC α and PKC δ were immunoprecipitated from 200 μ g of protein extracts as described above. Immunocomplexes were washed three times with the kinase buffer (20 mM Tris-HCl, 10 mM MgCl₂, pH 7.5) and then incubated with a PKC-specific peptide substrate, [³²P]ATP, and inhibitors of cAMP-dependent kinase and calmodulin kinase for 10 min at 30°C. ³²P incorporated into the substrate was separated from residual ³²P using p81 filters and subsequently quantified by scintillation counting. β -actin antibody was used as negative control in immunoprecipitations.

p21^{Ras} activity assays

Cells were cultured in DMEM containing 0.5% serum for 48 h. Then cells were lysed in a buffer containing 25 mM HEPES, pH 7.5, 150 mM NaCl, 1% Igepal CA-630, 0.25% sodium deoxycholate, 10% glycerol, and 25 mM NaF. Protein was normalized to 1 μ g/ μ l, and activated Ras was affinity-precipitated by mixing 1 mg of cell lysate with 10 μ g of Raf-RBD-agarose bead conjugate (Upstate Biotechnology, Inc.) for 60 min at 4°C. The conjugates were washed three times in lysate buffer and then separated on a 10% SDS-polyacrylamide gel. Proteins were transferred onto a polyvinylidene difluoride (PVDF) membrane and immunoblotted with a monoclonal p21^{Ras} antibody (BD Transduction Laboratories).

DNA profile analysis

1 \times 10⁵ cells were plated in a 60 mm dish and grown until confluent. Cells were harvested and resuspended with 1 ml of a 35% ethanol/DMEM solution for 5 min at room temperature. Cells were collected and stained with solution containing 50 μ g propidium iodide /ml and 25 units RNase/ml in PBS, and incubated in the dark for 30 min at room temperature for flow cytometric analysis.

Immunoblotting analysis

Harvested cells were disrupted in a buffer containing 20 mM Tris (pH 7.4), 0.5% NP-40, and 250 mM NaCl. Total protein (40 μ g) was separated on 10% SDS-polyacrylamide gels and transferred to nitrocellulose membranes or PVDF membranes. Membranes were blocked overnight and probed with affinity-purified antibodies against p21^{Ras} (BD Transduction Labs), PKC α , β , η , θ , δ (BD Transduction Lab), caspase-3, caspase-9 (Cell Signaling), β -actin (Sigma) Akt, phospho-Akt specific for serine473, Erk1/2, or p-Erk1/2 (Santa Cruz). After washing, the blots were incubated with horseradish peroxidase-conjugated secondary antibodies and visualized using the Amersham enhanced chemiluminescence ECL system.

Cell apoptosis assay

Terminal deoxynucleotidyl transferase-mediated deoxyuridine triphosphate nick end-labeling (TUNEL) assay was used for apoptosis assay. Briefly, cells were fixed with 4% paraformaldehyde in PBS overnight at 4°C. The samples were washed three times in PBS and permeabilized with 0.2% Triton X-100 in PBS for 15 min on ice. After rinsing twice with PBS, the samples were incubated with the TMR red TUNEL reagent (Roche) at 37°C in the dark, according to the manufacturer's instructions. Apoptotic cells were identified by fluorescent microscopy.

Statistical analyses

Results are expressed as mean \pm SD. Statistical analysis was performed using Student's *t*-test, and *P*-values $<$ 0.05 were considered significant.

RESULTS

Selective down-regulation of PKC δ induces apoptosis in cells with activated p21^{Ras}

To begin determination of whether a specific PKC isozyme is responsible for suppression of Ras-mediated apoptosis, six isozyme-specific or non-specific PKC inhibitors were used to suppress PKC α , PKC α/β , PKC β 1/2, PKC δ or general PKC activity (Table 1). Optimal maximal effective concentrations of each inhibitor ($>$ 90% inhibition of relevant enzyme activity) were determined in pilot experiments (Fig. 1A or data not shown). The elevated activity of p21^{Ras} in the KBalb, the MIA PaCa-2, and NIH/3T3-Ras cell lines, compared their counterpart cell lines (Balb, BxPc-3, and NIH/3T3, respectively), was confirmed by an activated p21^{Ras} pull-down assay (Fig. 1B). Cells were treated for 48 h and cell proliferation was quantitated by an MTT assay. Among the isozymespecific inhibitors, rottlerin dramatically and specifically decreased the proliferation of both MIA PaCa-2 and NIH/3T3-Ras cells (both of which express a mutant, activated p21^{Ras} protein), compared to the corresponding Hs766T cells and NIH/3T3 cells (which contain wild-type p21^{Ras}) (Figs. 1C; 1D). The pan-PKC inhibitor Bis-1 and the DAG antagonist HMG also strongly suppressed the growth of the cells containing activated p21^{Ras} consistent with our previous studies, while having no significant effect on most of the cell lines which contained a wild-type p21^{Ras} (7, 8). In contrast, rottlerin modestly but consistently stimulated proliferation of those cell lines expressing wild-type p21^{Ras} (Balb and Hs766T cells). Cytofluorometric analysis of PI-stained nuclei showed that rottlerin caused 24–35% apoptosis (manifested as hypodiploid nuclei) in activated-Ras-containing cell lines after 60 h of treatment, whereas the matched cells containing wild-type p21^{Ras} displayed only 12–14% apoptosis (Fig. 1E). PKC δ expression and activity was analyzed in the rottlerin-treated cells to confirm suppression of activity.

In order to confirm inhibition of PKC δ activity by rottlerin, both *in vivo* and *in vitro* kinase assays were performed. For the *in vivo* assays, cells were treated with rottlerin at 20 μ M for 48 h, then lysed and PKC δ was immunoprecipitated by a specific anti-PKC δ antibody. The incorporation of [γ -³²P] ATP into a PKC substrate peptide (QKRPSQRSKYL) by the immunoprecipitates was quantitated. Exposure to rottlerin, at the concentrations used in the apoptosis studies described above, blocked greater than 85% of the PKC δ activity in all of the cell lines tested, except Balb (Fig. 2A). In contrast, rottlerin produced only a slight, statistically-insignificant decrease in PKC α activity (Fig. 2B). For the *in vitro* kinase assay, rottlerin was added to immuno-purified PKC δ protein and incubated for 4 h, and PKC δ kinase activities were assayed using the artificial substrate. Rottlerin directly inhibited 55–90% of PKC δ activity (Fig. 2C),

Cell lines expressing activated p21^{Ras} consistently demonstrated elevated levels of PKC δ activity relative to the corresponding lines containing wild-type p21^{Ras} (Figs. 2A&B). Similarly, total levels of PKC δ protein were elevated in the cell lines containing activated p21^{Ras} relative to the corresponding lines containing wild-type p21^{Ras} (Fig. 2D). The data obtained with the chemical inhibitors of PKC isozymes is consistent with PKC δ being the relevant PKC target for Ras-mediated apoptosis.

Unexpectedly, exposure to rottlerin for 48 h suppressed PKC δ protein levels in all cell lines tested (Fig. 2D), whereas the levels of other PKC isozymes, including PKC α , $-\beta$, $-\eta$, and $-\theta$, were not changed by rottlerin (Fig. 2E). The magnitude of suppression of PKC δ levels by rottlerin in the cells containing activated p21^{Ras} (3–5 fold) approached the magnitude of

the suppression of PKC δ activity in these cells by rottlerin (6–10 fold). Thus, although we can demonstrate direct inhibition of PKC δ activity by rottlerin in *in vitro* assays, the marked suppression in PKC δ activity observed after 48 h *in vivo* may be due to suppression of isozyme protein levels as well as direct inhibition of enzyme activity.

As the specificity of chemical kinase inhibitors is never absolute, we employed two specific genetic techniques to suppress PKC δ activity. At 48 h after transfection of each of two PKC δ -specific hairpin vectors targeted at different PKC δ sequences into matched pairs of cell lines, BxPc-3 / MIA PaCa-2 and NIH/3T3 / NIH/3T3-Ras, immunoblot analysis demonstrated that expression of PKC δ protein was significantly diminished by transfection with the PKC δ -siRNA-2 vector. In contrast, the PKC δ -siRNA-1 vector did not produce significant knockdown of PKC δ protein (Fig. 3A). Because the efficiency of transient transfection in these cells was less than 50%, these analyses likely underestimate the activity of the siRNAs in an individual cell. Control experiments showed no changes of PKC δ by vehicle alone or scrambled siRNA. PKC δ siRNA also had no effect on PKC α expression. Seventy-two hours after transfection with pRNA-U6.1-GFP-control siRNA (scrambled hairpin sequence) or pRNA-U6.1-GFP-PKC δ -siRNA-2, all cell lines were harvested to determine the apoptotic fraction by TUNEL assay. Cells which took up the vector DNA were identified by green fluorescence. TUNEL-positive cells stained red. Superimposition displayed transfected, apoptotic cells as yellow (Figs. 3C&D). For cells transfected with the PKC δ hairpin vector, 40–50% cells were undergoing apoptosis at the 48 h time point, whereas cells transfected with the control scrambled hairpin vector displayed a frequency of apoptosis of less than 10% (Fig. 3H).

Similar results were obtained when a dominant-negative, kinase-dead PKC δ mutant protein was used as an alternative method of blocking specifically PKC δ activity in cells expressing an activated p21^{Ras} or wild-type p21^{Ras} (Figs. 3E&F). Transfection of NIH/3T3-Ras cells with the dn-PKC δ vector produced a 30–40% fraction of cells with a hypodiploid (apoptotic) DNA content, compared to a less than 5–10% apoptotic fraction in cells expressing a wild-type p21^{Ras} (Fig. 3H). Transfection of the empty vector as a control generated no significant apoptosis above background levels. The induction of apoptosis by the competitive expression of dominant-negative PKC δ with a single-base mutation, rendering it catalytically inactive, also demonstrates that it is the kinase activity of PKC δ that is required for the survival of cells expressing activated p21^{Ras} rather than a non-catalytic function of the molecule. We studied the effects of knockdown of PKC α (*via* expression of a PKC α -siRNA) as a specificity control. Forty-eight hours after transfection of PKD-PKC α V6, the levels of PKC α protein were significantly decreased in both NIH/3T3 and NIH/3T3-Ras cells (Fig. 3B). Analysis of apoptosis demonstrated that PKC α inhibition by PKC α -siRNA induced apoptosis in approximately 20% of both NIH/3T3 and NIH/3T3-Ras cells, with no selective toxicity for cells containing an activated p21^{Ras} (Figs. 3G&H). This finding was consistent with results of the MTT assay (Fig. 1C & D). Collectively, the data demonstrate that the PKC δ isozyme plays a critical survival role in p21^{Ras}-induced apoptosis.

PKC δ inhibition induces mitochondrial apoptotic pathways in cells expressing an activated p21^{Ras}

To further characterize the molecular mechanisms of p21^{Ras}-mediated apoptotic pathways, we investigated the influence of mitochondrial apoptotic signaling when PKC δ activity is suppressed by assay of caspase-3 and caspase-9 activation. Immunoblot analysis demonstrated that exposure to rottlerin activated both procaspase-3 and procaspase-9 exclusively in the cells expressing activated p21^{Ras} (Figs. 4A&B). For caspase-3, the full-length protein (35 kDa) and the large cleavage fragment (17 kDa) were detected (Fig. 4A);

three activation fragments from caspase-9 zymogen (35, 17 & 10 kDa) were detected (Fig. 4B).

The PI₃K Ras effector pathway is sufficient to sensitize cells to apoptosis by PKC δ inhibition

Although previous studies have demonstrated that either constitutive expression of activated p21^{Ras} or acute increases in endogenous p21^{Ras} activity stimulate apoptosis following inhibition of PKC activity in multiple types and lineages of cells, the roles of specific p21^{Ras} downstream effectors in the process have never been determined. In general, three major effector pathways activated by Ras have been defined: Raf1/MAPK, Ral-GDS, and PI₃K. To begin to address this question, p21^{Ras} effector loop mutants, consisting of the activating Ras mutation (V12) and a second mutation (S35, G37, or C40) were employed. The three RasV12 mutants (S35, G37, and C40) differ in their ability to bind to p21^{Ras} effectors (Raf, Ral-GEFs, and the p110 subunit of PI₃K, respectively) (29). Cells were treated with 20 μ M rottlerin for 60 h, and subjected to flow cytometric analysis. All three p21^{Ras} downstream effector-loop mutants stimulated apoptosis to some extent after inhibition of PKC δ , although expression of the C40 mutant consistently generated the greatest amount of apoptosis (data not shown).

The Ras-effector loop mutants are not completely specific in their activation of a single Ras effector pathway, and each activates all three pathways to some degree. In order to more clearly identify the down-stream effector of p21^{Ras} relevant to Ras-mediated apoptosis, we activated single effector pathways using expression vectors for activated PI₃K (p110-CAAX), Raf (Raf-22W), and Ral-GEF (RIF-CAAX) into NIH/3T3 cells. The dominant-negative RalA-22N vector was used as control. Constitutive activation of the PI₃K pathway (transfection of p110-CAAX) was capable of inducing apoptosis after PKC δ inhibition (apoptotic frequency: 38.22%). In contrast, the other Ras effectors Raf and Ral-GEF induced little apoptosis in response to PKC δ suppression (Fig. 5A).

To independently confirm that the downstream p21^{Ras} effector PI₃K plays a critical role in Ras-mediated apoptosis, a PI₃K-specific inhibitor (LY294002) and a MAPK inhibitor (PD98095) were employed. In dose-ranging studies, we evaluated concentrations of LY and PD from 1.0 to 60 μ M, assaying for efficiency of target enzyme inhibition and for toxicity. Even at 60 μ M concentrations of LY294002 or PD98095, only modest and non-statistically-significant levels of apoptosis were observed. At 10 μ M concentrations, there were no signs of toxicity or apoptosis (data not shown). Their effectiveness in suppressing the relevant target pathways at 10 μ M concentrations was confirmed by immunoblot (Figs. 5B&C). A NIH/3T3 cell line stably-expressing p110-CAAX was pre-treated with Ly294002 (10 μ M) for 30 min, and then rottlerin (20 μ M) was added for 60 h. In the presence of the PI₃K inhibitor, the apoptosis induced by rottlerin decreased nearly 50% (Fig. 5D). Similarly, in KBalb cells, LY294002 blocked 50% of the apoptosis induced by PKC δ inhibition (Fig. 5E). In cells which contained an activated p21^{Ras} but not those with activation of the single PI₃K-effector pathway, the MAPK inhibitor PD98095 also suppressed p21^{Ras}-dependent apoptosis to some extent, but to a substantially lesser degree than did PI₃K-inhibition. Collectively, these data demonstrate that the pro-apoptotic signal manifested during Ras-mediated apoptosis following PKC δ inhibition is mediated mainly through the downstream effector PI₃K, although the MAPK effector pathway may contribute to some extent in this process.

The anti-apoptotic Akt Ras-effector pathway is regulated by PKC δ

One direct downstream target of the PI₃K pathway are the members of the Akt kinase family, which have well-characterized pro-survival and anti-apoptotic activities, mediated

through regulation of the mitochondrial apoptotic machinery and the expression of genes involved in this process. To determine the role of Akt in p21^{Ras}-mediated apoptosis, we first examined Akt levels and activity in three paired cell lines with wild-type or mutant Ras alleles. These cells were treated with rottlerin for 48 h or left untreated. Akt activity, as assessed by quantitation of serine⁴⁷³-phosphorylated Akt, was down-regulated approximately 50% by rottlerin in all three cell lines expressing activated Ras (Fig. 6A). In contrast, no significant changes in levels of p-Akt were induced in the cell lines expressing wild-type Ras. Exposure to rottlerin also suppressed total Akt levels to a modest extent in some cell lines, but this effect was independent of the presence of activated p21^{Ras}. If the suppression of Akt activation by PKC δ suppression in activated Ras-containing cells were responsible for the resulting apoptosis, enforced expression of activated Akt should be able to effect rescue from apoptosis. MIA Paca-2 and NIH/3T3-Ras were co-transfected with a PKC δ -hairpin vector (which efficiently suppressed PKC δ levels in these experiments by at least 85%, see Fig. 3A for an example) and a constitutively-activated Akt (vAkt) expression vector, or a cAkt expression vector as a control. Akt activity in these cells was confirmed by immunoblotting for phosphorylated (activated) Akt (Fig. 6B). In both cell lines containing activated Ras, expression of vAkt reversed approximately 70–80% of the apoptosis induced by knock-down of PKC δ . Expression of a non-activated, wild-type Akt, however, did not similarly rescue the cells (Figs. 6C, D&E). To further support a role for Akt in Ras-mediated apoptosis induced by PKC δ inhibition, Akt activity was analyzed after PKC δ knock-down. Balb and KBalb cells were treated with a PKC δ siRNA or a scrambled siRNA with a fluorescein tag (serving as control and to quantitate transfection efficiency). After 48 h, cells were harvested and p-Akt expression was analyzed. p-Akt expression was dramatically decreased when PKC δ expression was knocked-down by PKC δ siRNA in KBalb cells (Fig. 6F). For reasons not yet elucidated, we found PKC δ -siRNA was consistently not as effective in downregulating PKC δ in Balb cells as it was in KBalb cells containing a mutated, activated p21^{Ras} (KBalb cells) (*e.g.* 62% suppression in Balb compared to 83% suppression in KBalb). Accordingly, the effects of PKC δ downregulation on p-AKT levels cannot be directly compared between Balb and K-Balb cells in these studies. Collectively, these studies indicate that activation of Akt is required for survival of cells expressing activated Ras, that PKC δ is required for the activation of Akt by Ras, and that induction of Ras-mediated apoptosis by suppression of PKC δ is effected through interference with this anti-apoptotic Ras effector pathway.

A pro-apoptotic p21^{Ras} effector, NORE1 (33), which binds to Ras-GTP and is a member of the RASSF family of tumor suppressors, has been described by Eckfeldt (34). To determine whether activation of this Ras-effector pathway plays any role in Ras-mediated apoptosis induced by suppression of PKC δ , we employed vectors expressing the fragment of NORE1 which binds to its apoptotic effector MST1 (aa 358–413), or the carboxy-terminal non-catalytic segment of MST1 (aa 307–487) as GST fusion peptides, each of which has been demonstrated to block formation of the NOR-MST1 apoptotic complex (33). Neither peptide attenuated the apoptosis induced by suppression of PKC δ (data not shown).

p21^{Ras} up-regulates PKC δ protein levels post-transcriptionally

We observed that cell lines expressing an activated p21^{Ras} invariably expressed substantially more PKC δ protein than did their wild-type p21^{Ras}-expressing counterparts (Fig. 7A), suggesting that p21^{Ras} activity may upregulate PKC δ protein expression as well as activity (Fig. 2D&2E). To test this hypothesis, NIH/3T3 cells were transfected with pSG5-H-(V12)Ras and PKC δ protein levels were assayed over time. PKC δ protein levels increased rapidly after transfection, reaching peak levels at 18 h (Fig. 7B). In contrast, transfection with the empty pSG5 vector had no effect on PKC δ protein levels. Conversely, when p21^{Ras} activity, or PI₃K activity, in KBalb and NIH/3T3-Ras cells was inhibited by the Ras

inhibitor FPT III or the PI₃K inhibitor LY294002, respectively, PKC δ protein levels fell (Fig. 7C). This regulation of PKC δ by p21^{Ras} is not at the level of transcription, as PKC δ transcript levels, assessed by quantitative RT-PCR, did not vary as a function of p21^{Ras} activity (data not shown).

To determine which p21^{Ras} effector pathway mediated the upregulation of PKC δ protein, p110-CAAX, Raf-22w, RIF-CAAX, and RalA-28N expression vectors were each stably transfected into two different cell lines, and PKC δ protein levels were quantitated (Fig. 8A). The p110-CAAX expression vector was consistently the most potent inducer of PKC δ expression in both Balb and NIH/3T3 cells, although activation each of the other two effector pathways could also induce PKC δ protein expression to a variable extent. In agreement, the C40 p21^{Ras}-effector loop mutant, which activates predominantly the PI₃K pathway, was the most potent of the effector mutants for induction of PKC δ protein levels (Fig. 8B). The expression of other isozymes of PKC ($-\alpha$ and $-\theta$), β -actin and tubulin were not changed by expression of p21^{Ras} or PI₃K (Fig. 8C&D). Similarly, levels of p53, cyclin D1 and p16 were not changed by expression of the p21^{Ras} or PI₃K (data not shown).

DISCUSSION

The p21^{Ras} family members comprise critical molecular switches transducing signals to diverse downstream pathways, ultimately controlling such processes as proliferation, cytoskeletal integrity, apoptosis, cell adhesion, and cell migration (35–38). p21^{Ras} and Ras-related proteins are frequently deregulated in cancers, leading to invasion and metastasis and enhancement of survival by activation of anti-apoptotic pathways. Paradoxically, several studies have demonstrated that enforced, high-level expression of oncogenic p21^{Ras} can induce a permanent growth arrest in normal cells, mimicking natural senescence (39, 40). Activation of the Raf-1/MEK/p38MAPK pathway is thought to be essential for oncogenic p21^{Ras}-induced senescence (40, 41). Additionally, inactivation of the ARF/p53 tumor suppressor pathway in mouse fibroblasts and skin keratinocytes, or inactivation of the p16/Rb tumor suppressor pathway in human fibroblasts, can bypass p21^{Ras}-induced senescence, suggesting that cellular fate resulting from oncogenic p21^{Ras} signaling is dependent upon the cellular context and the integration of tumor suppressor signals (39, 42). Our laboratory and others have established p21^{Ras} as a modulator of apoptosis in transformed cells and malignant cells, and also in normal cells. Through FADD, caspase 8, and downstream effector reactive oxygen species (ROS), p21^{Ras} sensitizes cells to apoptosis induced by PKC inhibition (7)(43, 44). However, the specific molecular mechanism by which oncogenic p21^{Ras} mediates apoptosis in the setting of inhibition of PKC activity remains incompletely understood. As different PKC isozymes may have opposing functions with respect to cellular proliferation, differentiation, and apoptosis, a more complete understanding of this process required identification of the specific PKC isozyme required for survival of cells expressing activated p21^{Ras} along with elucidation of the specific p21^{Ras} downstream effectors evoking the apoptotic outcome.

In this study, we demonstrate that PKC δ activity is required to prevent the induction of apoptosis in cells expressing activated p21^{Ras}. It is noteworthy that p21^{Ras} activity, and in particular activation of the PI₃K pathway, up-regulates PKC δ protein levels, thus positively reinforcing an anti-apoptotic, protective response to p21^{Ras} dysregulation in the cell. Conversely, when this induction is prevented by siRNA knockdown of PKC δ , programmed cell death is initiated. Pro-apoptotic activity engendered by activated Ras has been described in a number of other systems (45), and isoform differences in the proapoptotic effects of Ras have been reported, with K-ras sensitizing cells to gamma irradiation-induced apoptosis, but H-Ras exerting a protective effect (46). Activated K-Ras has been reported to sensitize cells to the pro-apoptotic effects of PKC agonists, though phosphorylation of serine residue 181

of p21^{Ras} and its translocation to intracellular membranes (47). K-ras, but not H-ras was reported to bind to, and potentially inactivate or sequester, the anti-apoptotic proteins Bcl-2 and Bcl-XL (48). In contrast to these observations of isotype-specific differences in p21^{Ras} functions, we have previously reported, and confirm herein, that both activated K-ras and H-ras sensitize cells to apoptosis induced by inhibition of PKC δ activity, which prevents their obligate activation of the Akt survival pathway.

PKC δ has been reported to both positively and negatively regulate apoptotic programs (49–52). These findings have generated conflicting hypotheses as to the role of PKC δ in the control of cell proliferation and survival. The normal phenotype of PKC δ -null mice demonstrates that PKC δ is not required for appropriate control of cell proliferation during normal development (28, 53). In contrast, PKC δ may be suborned during cellular transformation and become necessary for one or more components of the malignant phenotype. Inhibition of PKC δ was reported to suppress the metastatic potential of breast cancer cells (54) and to reduced their survival (55). Similar results were reported using non-small cell lung cancer cells (56). Our findings support this hypothesis. We find that PKC δ functions as a survival signal in a variety of cells with dysregulated activation of p21^{Ras}. PKC δ expression is upregulated in response to p21^{Ras} activity, primarily through PI₃K activation, and is required for the survival of these cell lines. However, PKC δ is not required for the survival or proliferation of the counterparts of these cell lines containing wild-type p21^{Ras} whether or not they are transformed, and indeed suppression of PKC δ actually leads to a small but reproducible increase in the proliferation of normal cells.

Activated Akt is a well-established survival factor, exerting anti-apoptotic activity by suppressing the release of cytochrome c from the mitochondria (57). Activated Akt also inactivates the pro-apoptotic factors BAD and procaspase-9 by direct phosphorylation (58). A number of reports have suggested relationships between PKC activity and Akt activation. One recent study showed that PKC δ contributes to the phosphorylation of Ebp1 in PC12 cells (59). Phosphorylation of Ebp1 is required for its association/interaction with active nuclear Akt, and the Akt/Ebp1 complex mediates an anti-apoptotic effect in intact cells. A very recent study suggested that both PKC δ and PKC ϵ can negatively regulate Akt activity in mouse keratinocytes (60), although this study utilized only chemical inhibitors lacking absolute specificity. Gonzalez-While PKC α has been shown to promote the dephosphorylation and inactivation of Akt in prostate cancer cells (61), another study demonstrated that phosphorylation of PKC δ protects glioma cells from TRAIL-induced apoptosis by activation of Akt (62). In our study, we prove that Akt activity can be regulated by PKC δ (Fig. 6F), and that cells in which PKC δ has been selectively suppressed by treatment with PKC δ -siRNA can be rescued from apoptosis by enforced expression of a constitutively-activated Akt (Fig. 6C). These data suggest that oncogenic p21^{Ras} protein, through PI₃K and PKC δ , induces Akt activity, initiating an anti-apoptotic signaling cascade which is required for their survival. However, whether PKC δ alone is the major regulator of Akt activity under all conditions remains to be elucidated, as the levels of p-Akt were only slightly changes after knocking-down PKC δ in cells containing wild-type (normal) Ras (Fig. 6F). There are two possible explanations for the differences observed on PKC δ actions of Akt between normal cells and those containing activated p21^{Ras}: the first is the relative differences in efficiency of transient transfection, while the second is that the robust regulation of Akt activity by PKC δ requires the involvement of an activated p21^{Ras} protein.

It is noteworthy that the PI₃K effector pathway, in addition to generating the anti-apoptotic signal mediated through PKC δ and activation of Akt (and upregulation of PKC δ levels and activity), is also responsible for the pro-apoptotic signal delivered in Ras-transformed cells, which is uncovered and made manifest by inhibition of PKC δ . We show that isolated activation of the PI₃K pathway is sufficient to render cells susceptible Ras-mediated

apoptosis, and conversely, that inhibition of the PI₃K pathway is sufficient to protect cells from apoptosis initiated by suppression of PKC δ .

In conclusion, this study significantly extends our understanding of the mechanism underlying p21^{Ras}-mediated apoptosis by identifying the molecules required to redirect p21^{Ras} signaling from a proliferative/transforming outcome towards instead an apoptotic fate. Furthermore, elucidation of the particular PKC isozyme necessary for survival of cells transformed by p21^{Ras} suggests that selective suppression of PKC δ activity, and the consequent induction of apoptosis, is a potential strategy for targeting of tumor cells containing an activated p21^{Ras} GTPase.

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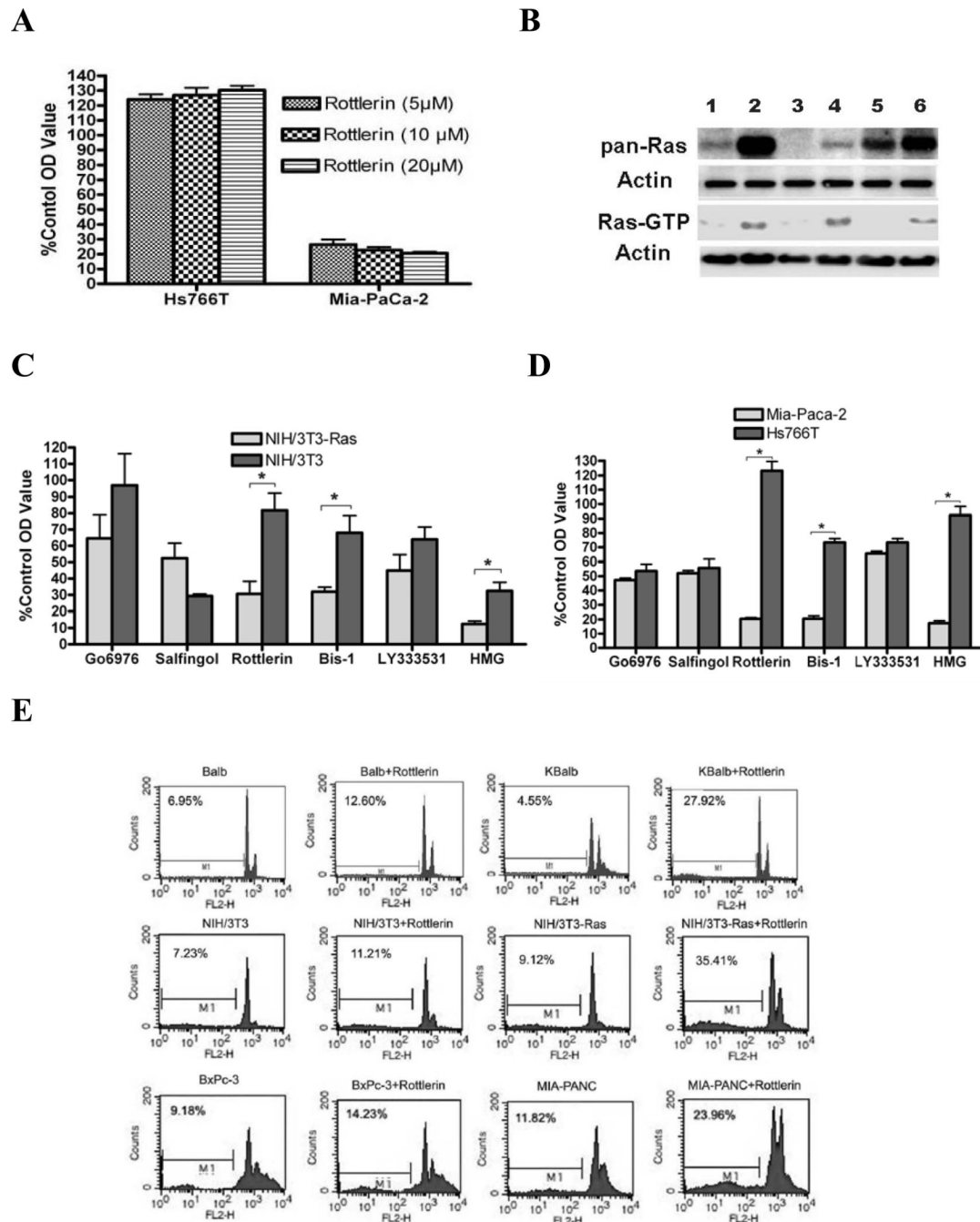
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The abbreviations used are

MAPK	mitogen-activated protein kinase
BIM-1	bisindolylmaleimide I
HMG	1-O-hexadecyl-2-O-methyl-rac-glycerol
FPT III	(E,E)-[2-Oxo-2-[(3,7,11-trimethyl-2,6,10-dodecatrienyl)oxy]amino]ethyl]phosphonic Acid, (2,2-Dimethyl-1-oxopropoxy) methyl Ester, Na
PKC	Protein kinase C

**Figure 1.**

Effects of isozyme-specific and non-specific PKC inhibitors on proliferation or apoptosis of mouse fibroblast cells and human pancreatic carcinoma cells with wild-type or activated p21^{Ras}. Cells were grown to 80% confluence in 96-well plates, then treated with the inhibitors at the concentrations indicated in Table 1. The corresponding solvents at equivalent volumes were used as vehicle controls. After 48 h treatment, cell growth was evaluated by MTT assay or FACS. Results are presented as the means \pm SD. Statistical significance was determined using a paired Student's *t*-test, and *p*-values <0.05 were considered significant (*, *p*<0.01). (A) p21^{Ras}-specific suppression of growth by PKC δ inhibition. Human pancreatic carcinoma cell cultures (Hs766T [wild-type p21^{Ras}] and

MiaPaCa-2 [activated p21^{Ras}] were treated with different doses of rottlerin, and cell growth assayed by MTT assay. **(B)** Assay of cellular p21^{Ras} activity. Nuclear-free lysates containing a total of 400 µg protein from each indicated cell type were used for analysis of Ras activity by Raf-RBD pull-down. Equal loading was demonstrated by re-probing the blot with anti-β-actin Ab. Pan-Ras protein expression levels were also analyzed. Lanes 1 to 6 represent: Balb, KBalb, NIH/3T3, NIH/3T3-Ras, BxPc-3 and MIA PaCa-2 cells respectively. **(C)** Growth of NIH/3T3 and NIH/3T3-Ras cells treated with isozyme-specific and non-specific PKC inhibitors. **(D)** Growth of Hs766T and MIA PaCa-2 cells treated with isozyme-specific and non-specific PKC inhibitors. **(E)** Nuclear DNA content of cells after inhibition of PKCδ. After 60 h treatment with rottlerin, cells were fixed and stained with propidium iodide, and the apoptotic (hypodiploid) fractions (M1 fractions) were evaluated by flow cytometry. Counts refers to cell numbers; FLH2 is a log scale of fluorescence channels. Similar results were observed in three independent experiments.

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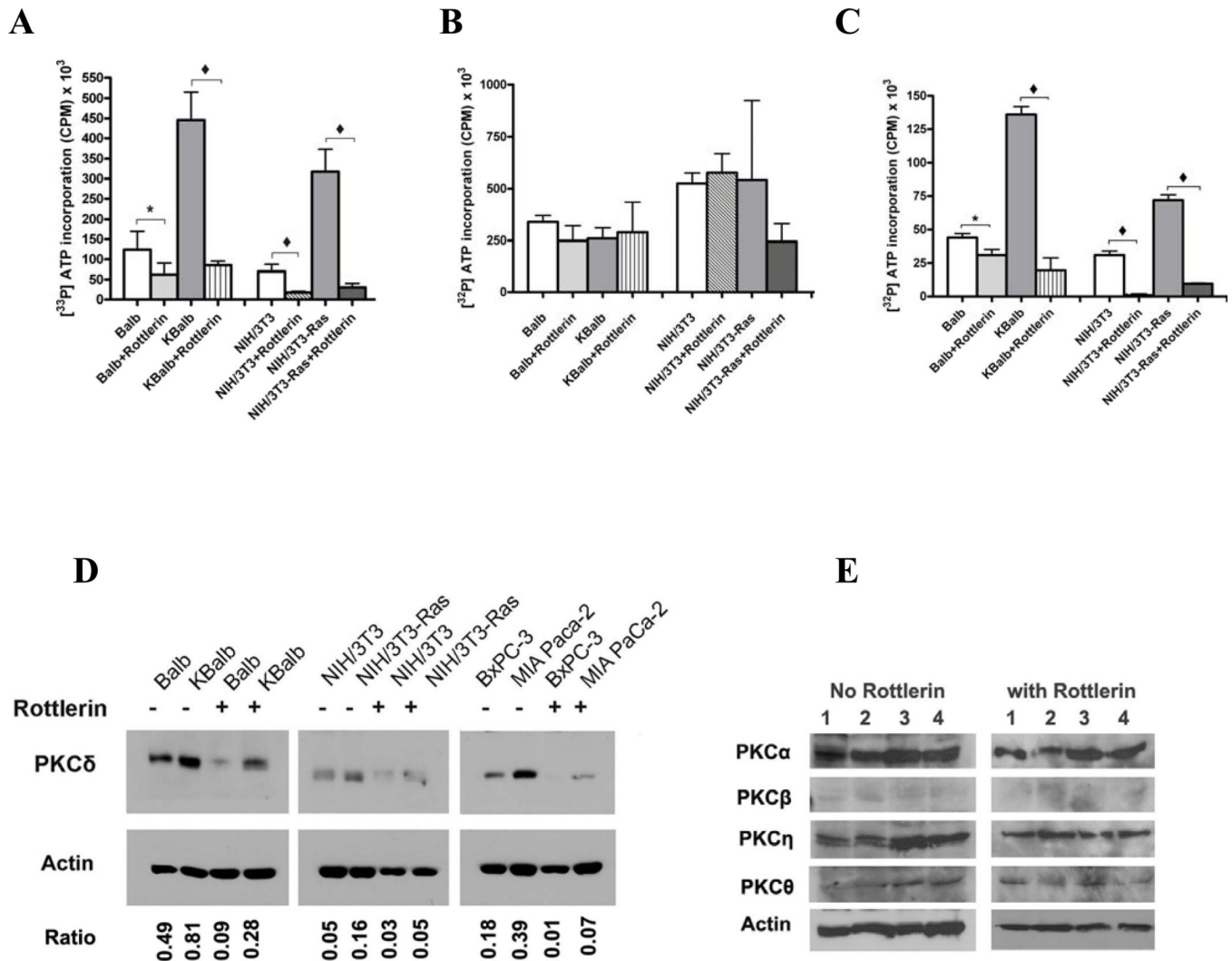


Figure 2.

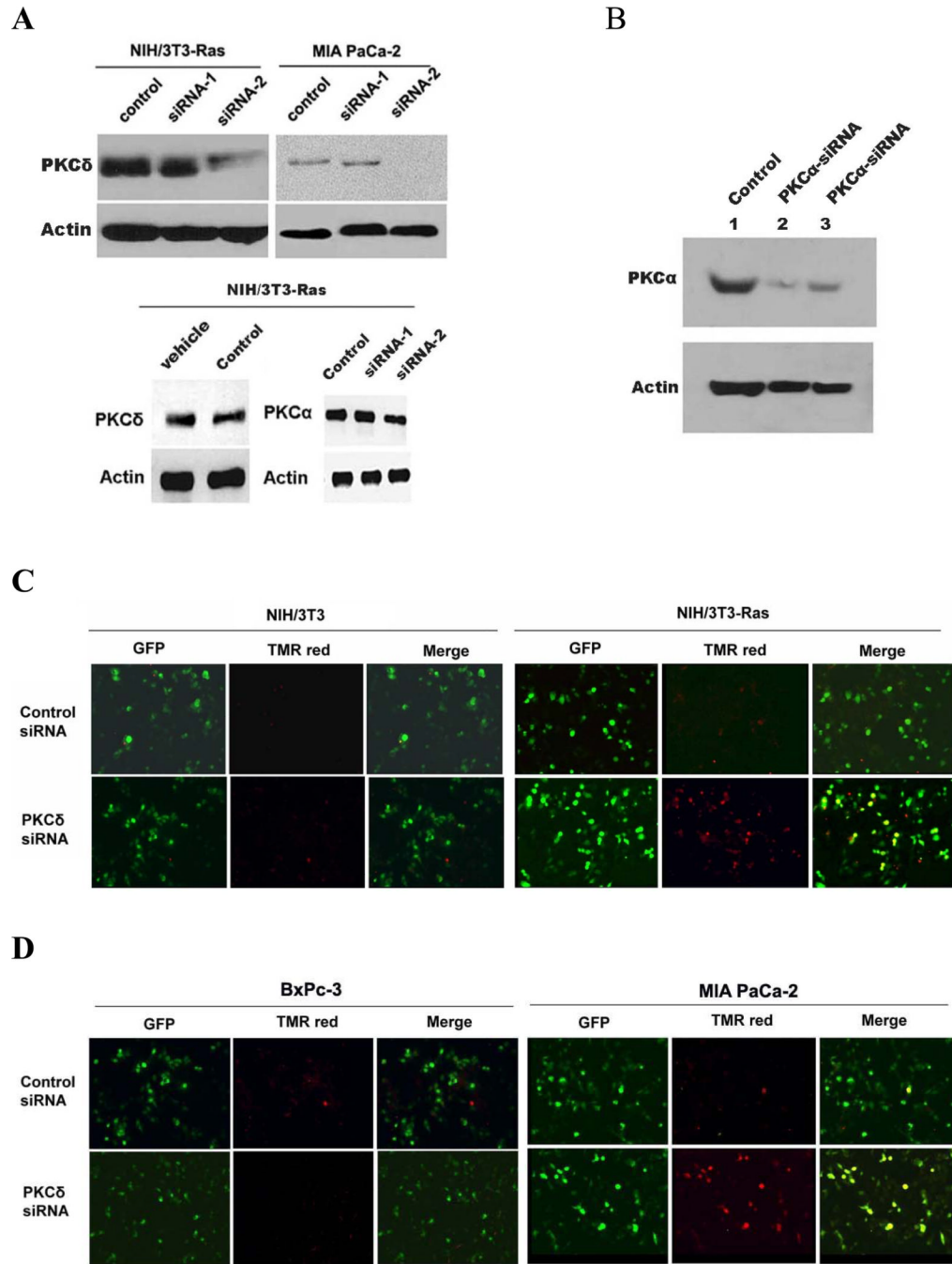
Effects of rottlerin on PKC δ and PKC α enzyme activity and protein levels. (A) PKC δ and (B) PKC α kinase activities. The indicated cell lines were treated with rottlerin (20 μ M) for 48 h, then harvested for analysis. 400 μ g of protein lysates were used for immunoprecipitation by isozyme-specific antibodies, and the immunopurified proteins were assayed using an artificial substrate. (C) PKC δ activity measured by *in vitro* assay. PKC δ was pulled down from cell lysates by IP and treated with rottlerin (20 μ M) for 4 h, and then subjected to an *in vitro* kinase assay. (D) PKC δ protein levels in cells after rottlerin treatment. 40 μ g aliquots of the same protein lysates used in the studies in panels A and B were separated and immunoblotted using an anti-PKC δ monoclonal antibody. Ratio: refers to the value of PKC δ expression/ β -actin expression measured by Software BandLead 3.0. (E) PKC isozyme expression levels after treatment with rottlerin. 1×10^5 cells were plated in each well of a 6-well plate. Cells were treated with rottlerin (20 μ M) or with the same volume of vehicle after they reached 80% confluence. After 48 h, cells were harvested, lysed in 50 μ l lysis buffer and the lysates separated and subjected to immunoblot analysis using anti-PKC isozyme-specific antibodies. Lanes 1 to 4 represent: Balb, KBalb, NIH/3T3 and NIH/3T3-Ras cells respectively. The immunoblots shown in the left and right panels were separated on the same gel and immunoblotted at the same time. Because the samples in the left panel came from untreated cells, whereas the samples in the right panel came from

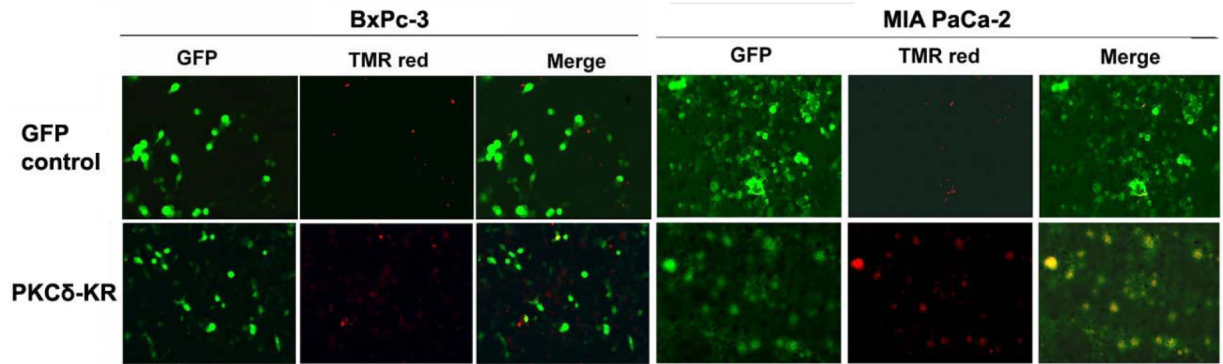
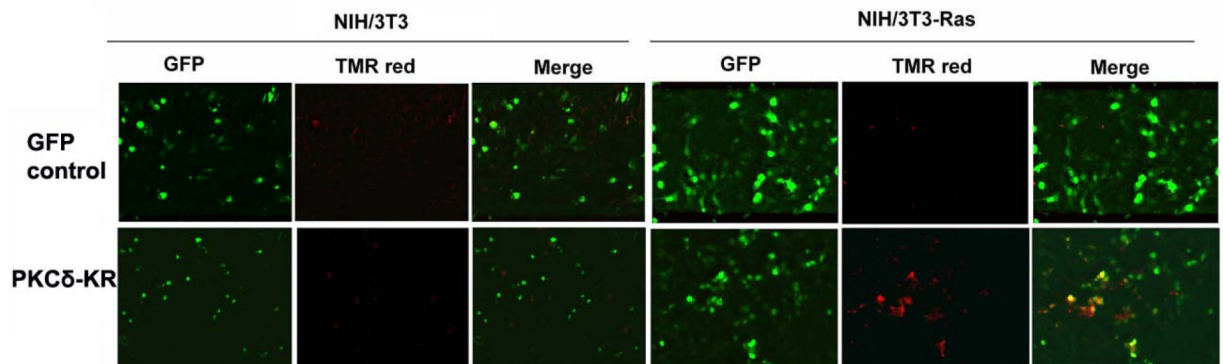
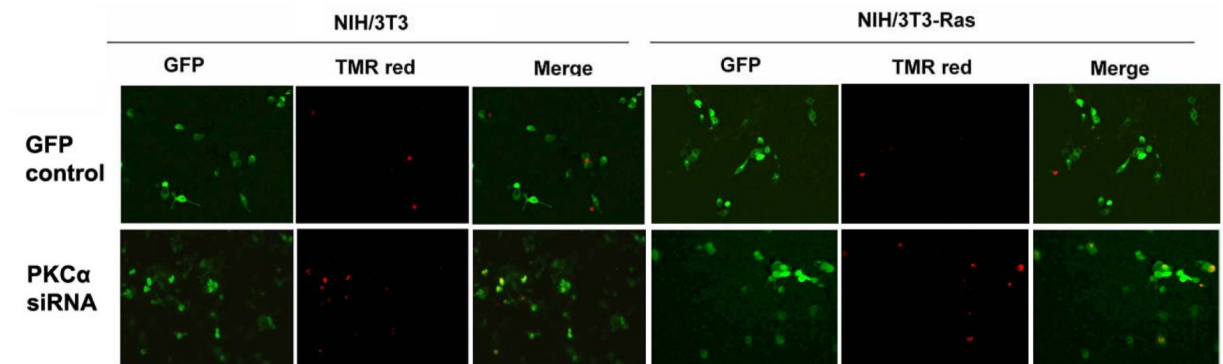
rottlerin-treated cells, and because rottlerin-treatment inhibited cell growth/proliferation, fewer numbers of cells were obtained in these groups, and therefore less protein/lysate was available for assay from the treated group compared to the vehicle control group. The mean activities \pm SD were obtained from at least three independent experiments, and assays from immunoprecipitations using an anti- β -actin antibody were used to determine the background activity, which was subtracted. Student's t-test was used for statistical analysis. (*, $p < 0.01$; \blacklozenge , $p < 0.001$).

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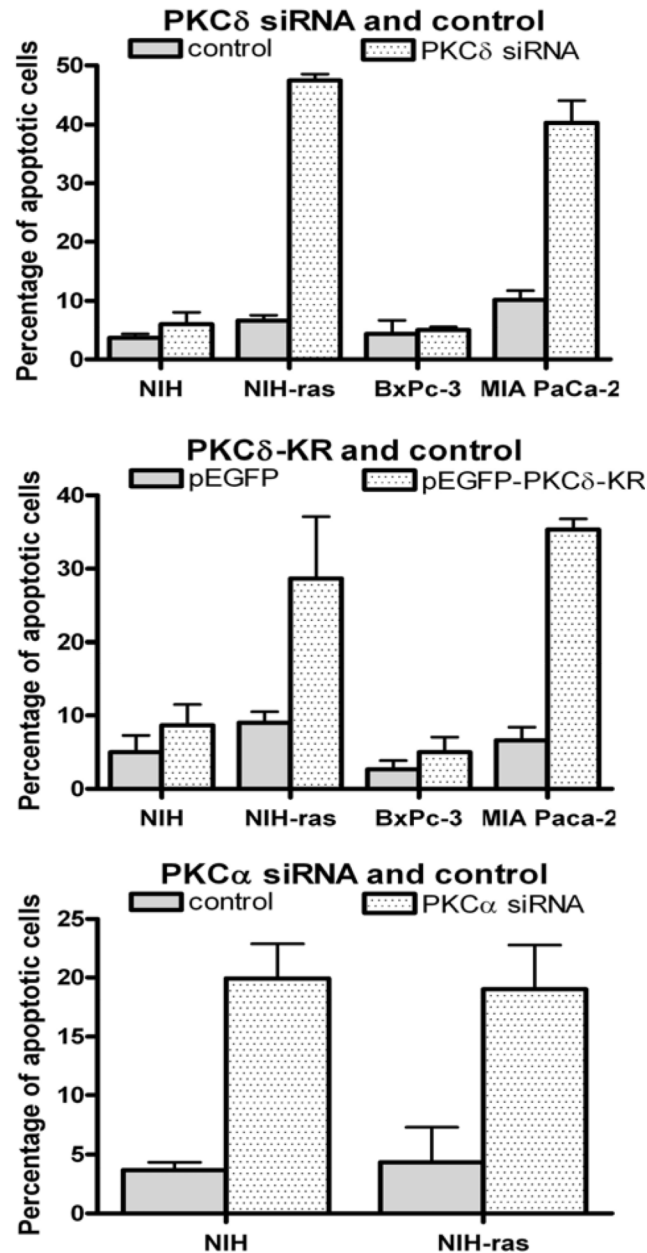
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**Figure 3.**

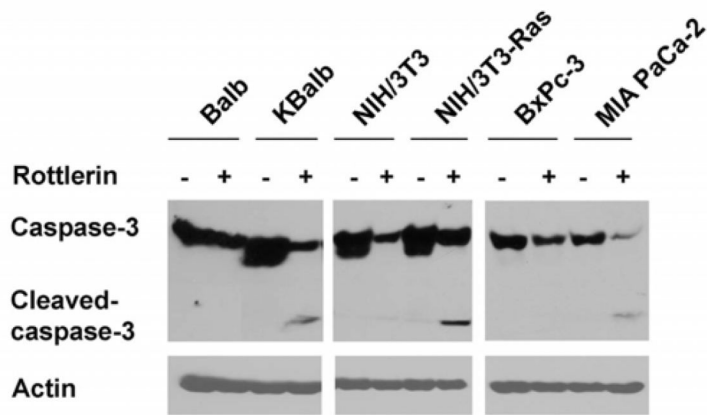
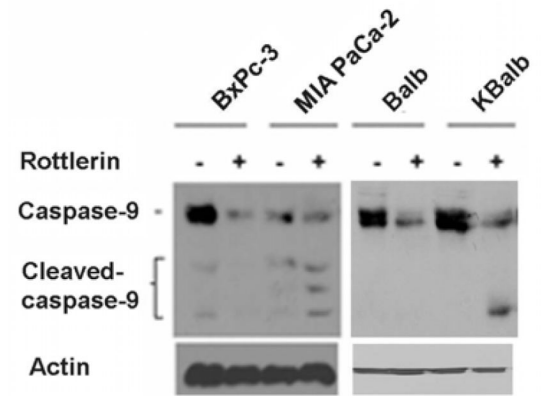
Effect of PKC δ inhibition on the viability of cells expressing activated p21^{Ras}. (A) Immunoblot analysis of PKC δ , PKC α or β -actin expression in NIH/3T3-Ras and MIA PaCa-2 cells transfected with PKC δ siRNA-1 or -2. A total of 50 nM of PKC δ siRNAs were used. Transfections with a scrambled siRNA or with vehicle alone were used as negative controls. In all experiments, transfections with scrambled siRNA or with vehicle alone yielded identical results, and in subsequent experiments only the scrambled siRNA transfection control is shown. (B) Knock-down activity of PKC α siRNA or scrambled siRNA (control) on NIH/3T3 (lanes 1 and 2) and NIH/3T3-Ras cells (lane 3) evaluated by

immunoblotting cell lysates for PKC α or β -actin. NIH/3T3-Ras (**C**) and MIA PaCa-2 (**D**) were transfected with pRNA-U6.1-GFP scrambled siRNA hairpin vector (control) or pRNA-U6.1-GFP-PKC δ siRNA-2 hairpin vector. After 72 h, apoptosis was detected by TUNEL assay. NIH/3T3 and BxPc-3 were used as control cell lines, respectively. Shown are 200x magnifications. MIA PaCa-2 cells (**E**) and NIH/3T3-Ras cells (**F**) were transfected with a dominant-negative, kinase-dead PKC δ vector. After 72 h, apoptosis was detected by TUNEL assay. BxPc-3 and NIH/3T3 were used as control cell lines (respectively). Shown are 200x magnifications. (**G**) NIH/3T3 and NIH/3T3-Ras cells were co-transfected with PKD-PKC α siRNA and pEGFP or scrambled siRNA and pEGFP. After 72 h, apoptotic cells were assayed by TUNEL reagent. (**H**) Quantitation of apoptotic cells in the transfected populations, with error bars indicating SD. Top panel: PKC δ siRNA; middle panel: PKC δ -KR; bottom panel: PKC α siRNA. Data shown are representative of at least three independent experiments.

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A**B****Figure 4.**

Inhibition of PKC δ induces the cleavage of caspase-3 and caspase-9 in cells expressed activated p21^{Ras}. All the cells were treated with 20 μ M rottlerin for 60 h. Thereafter, cells were harvested, and cell lysates were prepared and subjected to immunoblot analysis to detect the levels of caspase-3/cleaved caspase-3 (**A**) and caspase-9/cleaved caspase-9 (**B**) and β -actin. A representative blot from duplicate experiments, producing similar results, is shown.

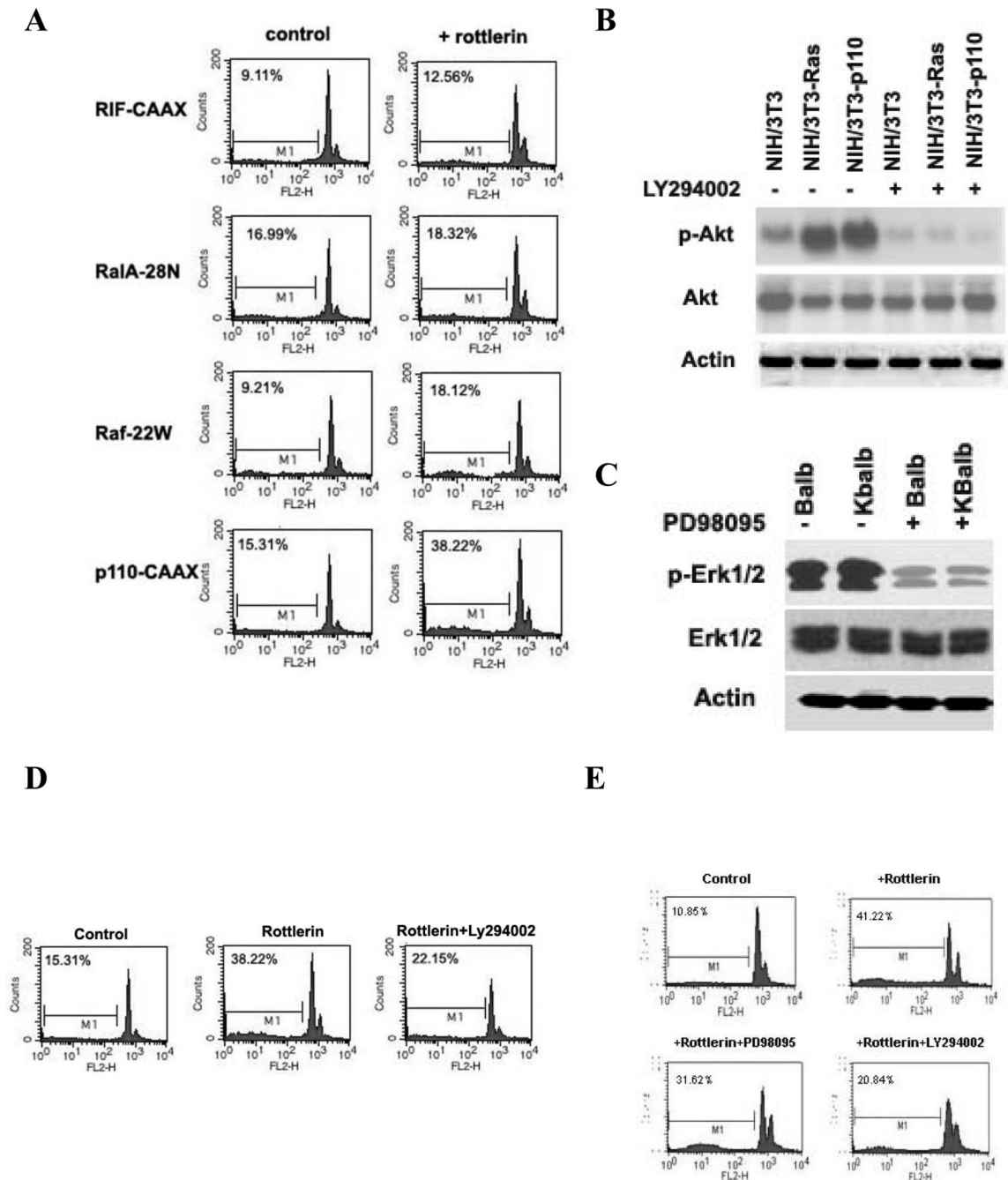


Figure 5.

Activation of apoptosis by individual p21^{Ras}-effector pathways. (A) NIH/3T3 cell lines stably expressing the p21^{Ras} downstream effectors Raf-22W (Raf-1), P110-CAAXC (PI₃K), RIF-CAAX (RIF), or RalA-28N (as a negative control) were treated with rottlerin (20 μ M) for 60 h and analyzed for the apoptotic fraction. (B) immunoblot analysis of Akt and p-Akt expression in NIH/3T3, NIH/3T3-Ras and NIH/3T3-p110CAAX cells after 48 h treatment with 10 μ M LY294002. (C) Erk1/2 and p-Erk1/2 expression in Balb and KBalb cells after 48 h treatment with 10 μ M PD98095. The left and right sides of immunoblot 5B and of immunoblot 5C were separated on the same gels, transferred to the same filters and

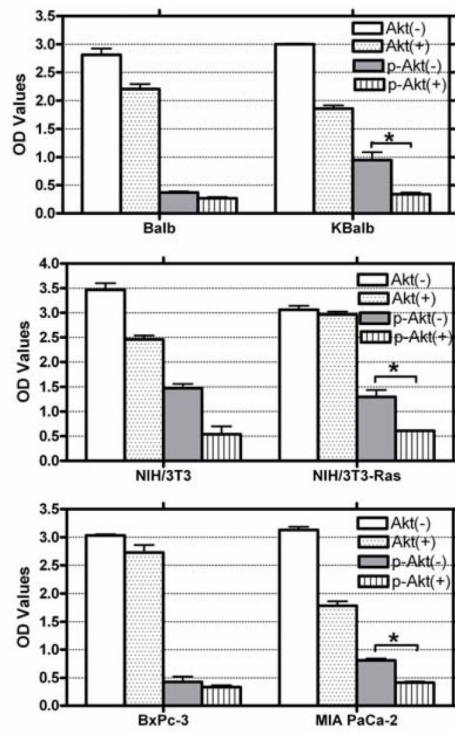
immunoblotted together. **(D)** Suppression of apoptosis by inhibition of PI₃K activity. NIH/3T3 cells stably expressed p110-CAAX were pre-treated for 30 minutes with LY294002 (10 μM) before rottlerin (20 μM) was added. The apoptotic (hypodiploid) fractions were evaluated 60 h later. **(E)** KBalB cells were pre-treated with vehicle solvent, LY294002 (10 μM), or PD98095 (10 μM) and then treated with rottlerin (20 μM). The apoptotic (hypodiploid) fractions were evaluated 60 h later. Cells were fixed and stained with propidium iodide. DNA fragmentation was analyzed by flow cytometry using the FL2-H channel. Data shown are representative of at least three independent experiments. Yes, (In this revision, gels 5C has been rerun, as requested by Reviewer.)

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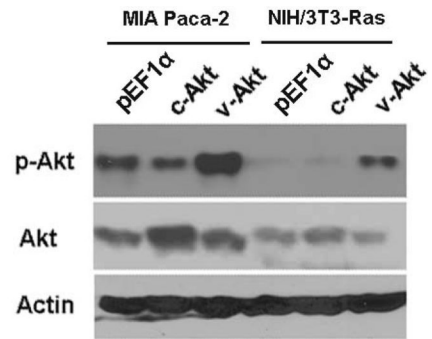
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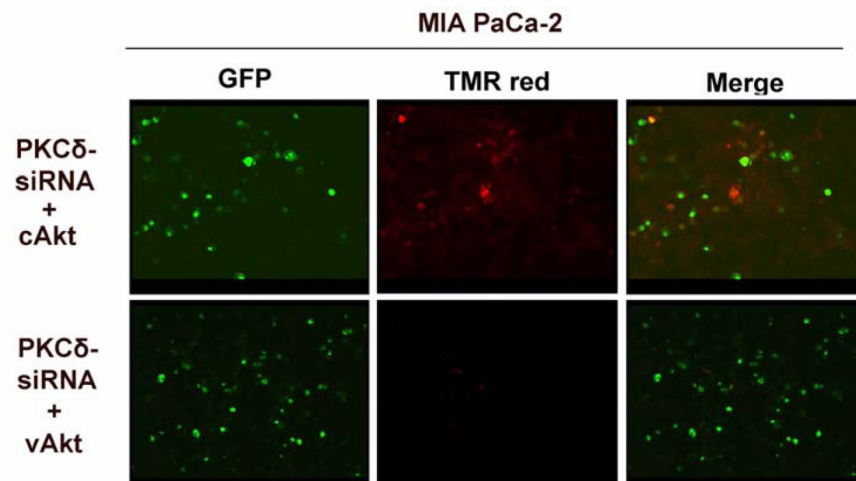
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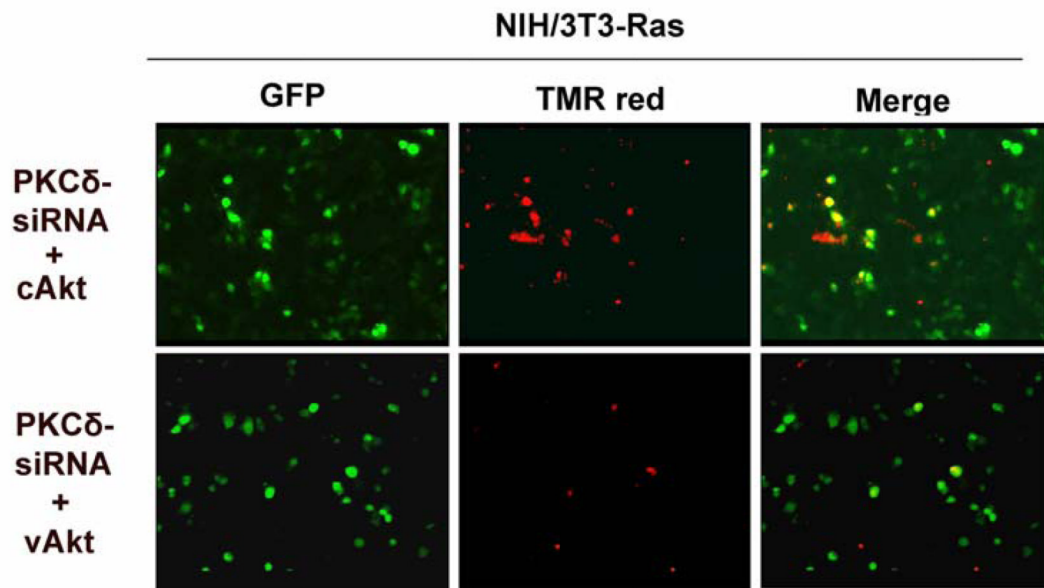
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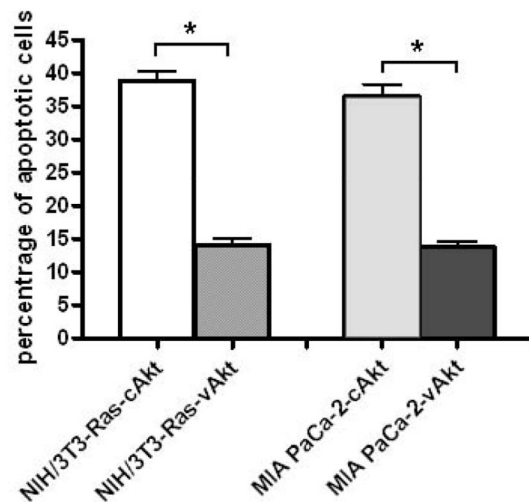
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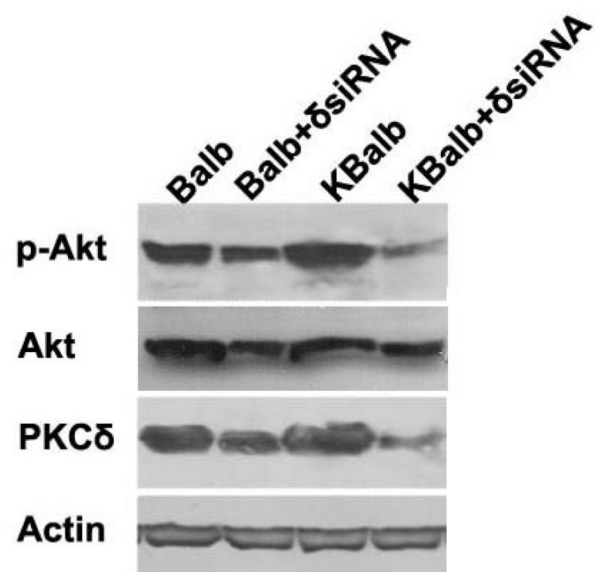


Figure 6.

Regulation of AKT activity by PKC δ . (A) Treatment with rottlerin reduces Akt activity in cells expressing activated Ras. All cells were treated with 20 μ M rottlerin for 60 h in 96-well plates. Total Akt and p-Akt levels were assayed using a SuperArray Case ELISA kit (*, $p < 0.005$). (+): treated with 20 μ M rottlerin; (-): treated with vehicle (control); (B) pAkt protein levels in MIA PaCa-2 and NIH/3T3-Ras cells, assayed by immunoblotting with an antibody specific for phosphorylated serine⁴⁷³ Akt, or for total Akt, or for β -actin. Both cell lines were transfected with pEGFP and with pEF1 α empty vector, or pEF1 α -cAkt, or pEF1 α -vAkt. GFP expression was used to normalize for transfection efficiency. (C & D) Activated Akt rescues cells from apoptosis induced by PKC δ knock-down. MIA PaCa-2 and NIH/3T3-Ras cells were cultured on 4-well chamber slides and were co-transfected with

pRNA-U6.1-GFP-PKC δ hairpin vector and with pEF1 α -vAkt or pEF1 α -cAkt vectors. The PKC δ -hairpin vector efficiently suppressed PKC δ levels in these experiments by at least 85% (e.g., see Fig. 3A). After 72 h, cells were fixed with 4% paraformaldehyde for 1 h at room temperature, and then subjected to TUNEL assay. Shown are 200x magnifications. **(E)** Quantitation of apoptotic cells in the transfected populations (*, $p < 0.001$). **(F)** Immunoblot of lysates from Balb or KBalB cells treated with PKC δ -specific siRNA or scrambled, control siRNA. Antibodies against PKC δ , total Akt, phospho-serine⁴⁷³ Akt, and β -actin were used. The results are representative of at least two independent experiments.

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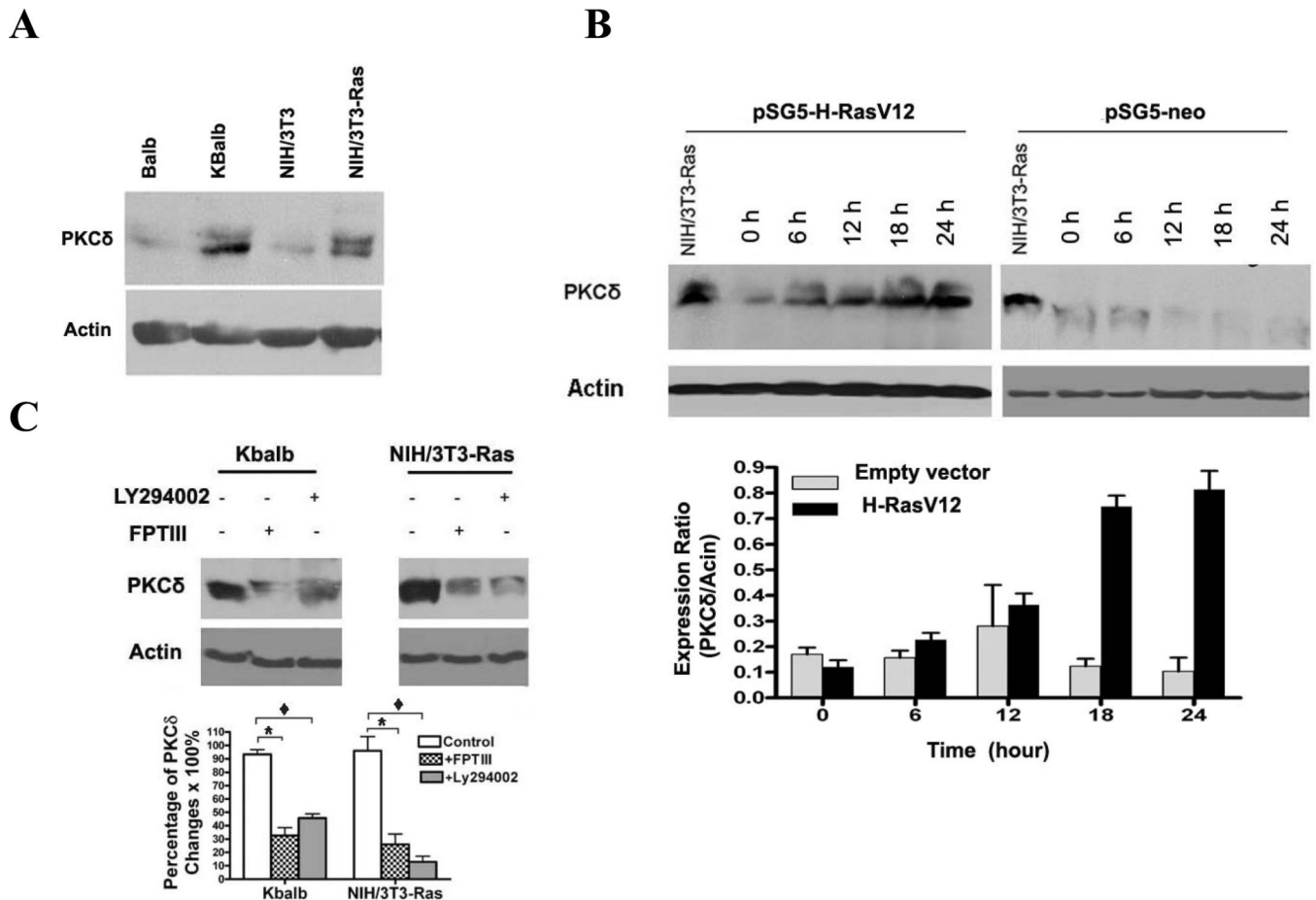
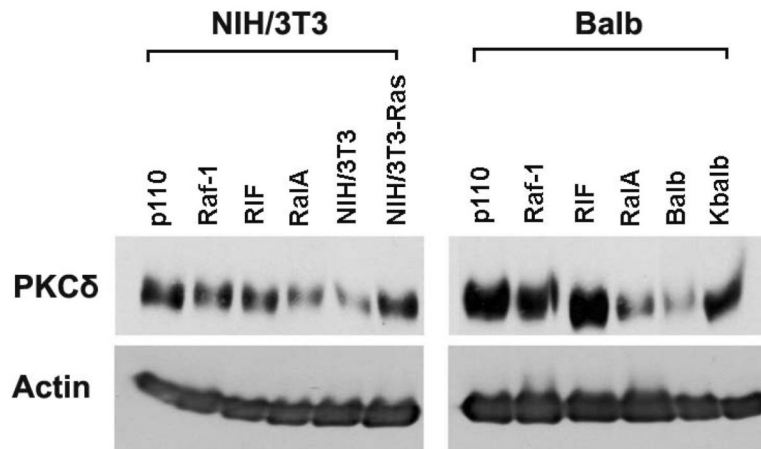
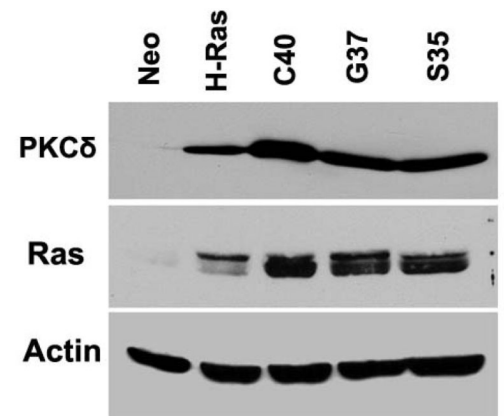


Figure 7. Regulation of PKC δ levels by p21^{Ras}. (A) Immunoblot analysis of PKC δ expression in matched cell line pairs NIH/3T3 and NIH/3T3-Ras; and Balb and Kbalb. Blots were stripped and reprobed for β -actin. (B) (Upper panel) NIH/3T3 cells were transfected with the pSG5-H-ras vector and harvested at the time points indicated. The NIH/3T3-Ras cell line was used as positive control. (Lower panel) The relative levels of PKC δ expression were quantitated by Software BandLead 3.0. (C) Effects of a p21^{Ras} inhibitor and a PI₃K inhibitor on PKC δ expression. (Upper panel) NIH/3T3-Ras cells were treated with FPT inhibitor III (100 μ M) or with Ly294002 (10 μ M) for 48 h, then lysed and subjected to immunoblot assay for PKC δ . Blots were then stripped and reprobed for p21^{Ras} and β -actin. (Lower panel) Quantitative measurement of changes in PKC δ Levels. Data shown are representative of at least three independent experiments (*, p < 0.01, and ♦, p < 0.05).

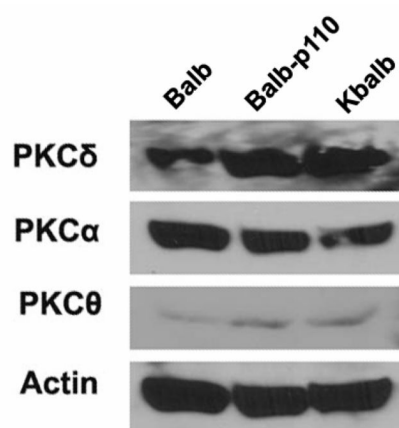
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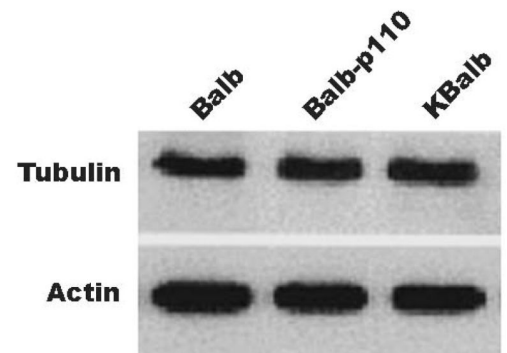


Figure 8. Effect of isolated p21^{Ras} effector pathways on PKC expression. Immunoblots of PKC δ , β -actin, Ras, PKC α , PKC θ , or tubulin in: (A) NIH/3T3-Ras cells or NIH-3T3 cells stably-expressing the indicated Ras downstream effectors or empty vector as control (left panel) and KBalb cells or Balb cells stably-expressing the indicated Ras downstream effectors or empty vector as control (right panel); (B) Balb cells stably-expressing Ras-effector loop mutants; (C & D) Balb cells stably-expressing PI₃K catalytic domain p110CAAX or KBalb Cells. A total of 50 μ g of protein was separated on a 10% SDS-PAGE for each sample. The results are representative of at least three independent experiments.

Table 1

Chemical kinase inhibitors used in this study

Name	Concentration (μM)	Target	Vendor
FPT III	100	p21 ^{Ras}	Calbiochem
Rottlerin	20	PKC δ	Calbiochem
Ly294002	10	PI ₃ K	Calbiochem
PD98095	10	MAPK	Calbiochem
Safingol	5	PKC α	Calbiochem
Go6976	0.3	PKC α/β	Calbiochem
Bisindolylmaleimide I	10	pan-PKC	Calbiochem
Ly333531	0.01	PKC β 1 and β 2	A.G.Scientific
HMG	120	pan-PKC	Calbiochem