

# Caspase-3-Dependent Proteolytic Cleavage of Protein Kinase C $\delta$ Is Essential for Oxidative Stress-Mediated Dopaminergic Cell Death after Exposure to Methylcyclopentadienyl Manganese Tricarbonyl

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In the present study, we characterized oxidative stress-dependent cellular events in dopaminergic cells after exposure to an organic form of manganese compound, methylcyclopentadienyl manganese tricarbonyl (MMT). In pheochromocytoma cells, MMT exposure resulted in rapid increase in generation of reactive oxygen species (ROS) within 5–15 min, followed by release of mitochondrial cytochrome C into cytoplasm and subsequent activation of cysteine proteases, caspase-9 (twofold to threefold) and caspase-3 (15- to 25-fold), but not caspase-8, in a time- and dose-dependent manner. Interestingly, we also found that MMT exposure induces a time- and dose-dependent proteolytic cleavage of native protein kinase C $\delta$  (PKC $\delta$ , 72–74 kDa) to yield 41 kDa catalytically active and 38 kDa regulatory fragments. Pretreatment with caspase inhibitors (Z-DEVD-FMK or Z-VAD-FMK) blocked MMT-induced proteolytic cleavage of PKC $\delta$ , indicating that cleavage is mediated by caspase-3. Furthermore, inhibition of PKC $\delta$  activity with a specific inhibitor, rottlerin, significantly inhibited

caspase-3 activation in a dose-dependent manner along with a reduction in PKC $\delta$  cleavage products, indicating a possible positive feedback activation of caspase-3 activity by PKC $\delta$ . The presence of such a positive feedback loop was also confirmed by delivering the catalytically active PKC $\delta$  fragment. Attenuation of ROS generation, caspase-3 activation, and PKC $\delta$  activity before MMT treatment almost completely suppressed DNA fragmentation. Additionally, overexpression of catalytically inactive PKC $\delta$ <sup>K376R</sup> (dominant-negative mutant) prevented MMT-induced apoptosis in immortalized mesencephalic dopaminergic cells. For the first time, these data demonstrate that caspase-3-dependent proteolytic activation of PKC $\delta$  plays a key role in oxidative stress-mediated apoptosis in dopaminergic cells after exposure to an environmental neurotoxic agent.

*Key words: apoptosis; oxidative stress; Parkinson's disease; environmental factors; manganese; dopaminergic degeneration*

Parkinson's disease (PD) is an idiopathic neurodegenerative disorder characterized by profound loss of dopaminergic neurons in the nigrostriatal tract. Although debated, most studies have concluded that aging, environmental neurotoxicant exposures, and genetic alterations are potential risk factors in the development of PD (Oertel and Kupsch, 1993; Langsten and Hill, 1998; Aschner, 2000; Simon et al., 2000). Recently, a study conducted on thousands of twins concluded that genetic factors do not play a role in the pathogenesis of geriatric onset of PD, which further supports the view that environmental factors are dominant risk factors in the etiology of PD (Tanner et al., 1999). Results of several epidemiological studies conducted in rural areas have also suggested that certain pesticides and other environmental factors, including transition metals such as manganese, have a positive association with increased incidences of PD (Seidler et al., 1996; Liou et al., 1997; Gorell et al., 1999). Occupational exposure to manganese during mining was shown to cause a Parkinson's-like

syndrome known as Manganism (Mena et al., 1967; Barbeau, 1984; Donaldson, 1987; Gorell et al., 1999). Furthermore, exposure to manganese-containing compounds such as manganese ethylene-bis-dithiocarbamate (a fungicide) and Bazooka (a cocaine-based drug) among farm workers and abusers, respectively, has been shown to result in adverse neurological defects (Roels et al., 1987; Ferraz et al., 1988; Wang et al., 1989; Thiruchelvam et al., 2000).

Methylcyclopentadienyl manganese tricarbonyl (MMT) has been used in Canada as an anti-knock gasoline agent and has been recently legalized for use in the United States as a replacement for tetraethyl lead [(CH<sub>3</sub>CH<sub>2</sub>)<sub>4</sub>Pb] in gasoline (Lynam et al., 1999; Zayed et al., 1999). Because MMT is a manganese-containing compound, its use has raised great a concern regarding increased exposure to the public and its possible adverse health effects (Frumkin and Solomon, 1997; Davis, 1998; Lynam et al., 1999; Zayed et al., 1999). Exposure to MMT produces a prolonged and more pronounced accumulation of manganese in rat brain as compared with manganese derived from an inorganic source, for example, MnCl<sub>2</sub> (Zheng et al., 2000). Administration of MMT produces seizures in mice (Fishman et al., 1987) and also results in depletion of dopamine in the mouse striatum (Gianutsos and Murray, 1982). Furthermore, MMT administration has been shown to be an effective inhibitor of complex I in mitochondrial electron transport chain (Autissier et al., 1977), an action similar to the pyridinium metabolite of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), a Parkinsonian toxin. Recently, we demonstrated that MMT exposure induces reactive oxygen species (ROS) generation, dopamine depletion, and cell death in dopamine-producing rat pheochromocytoma (PC12)

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cells, which can be protected by pretreatment with antioxidants (Wagner et al., 2000). To further understand the cellular mechanism of MMT-mediated apoptosis, we investigated whether oxidative stress induced by MMT can activate a series of cellular factors associated with apoptotic pathways, which could subsequently lead to programmed cell death in dopaminergic cells. Herein, we report that MMT exposure activates a novel apoptotic pathway in dopaminergic cells through caspase-3-dependent proteolytic cleavage of PKC $\delta$ .

## MATERIALS AND METHODS

**Reagents.** MMT was obtained from Sigma-Aldrich (St. Louis, MO); rottlerin was purchased from Calbiochem (San Diego, CA); acetyl-Asp-Glu-Val-Asp-aldehyde (Ac-DEVD-CHO), acetyl-Iso-Glu-Thr-Asp-7-amino-4-methylcoumarin (Ac-IETD-AMC), acetyl-Leu-Glu-His-Asp-7-amino-4-methylcoumarin (Ac-LEHD-AMC), and Z-Asp-Glu-Val-Asp-fluoromethyl ketone (Z-DEVD-FMK) were obtained from Alexis Biochemicals (San Diego, CA); Z-Val-Ala-Asp-fluoromethyl ketone (Z-VAD-FMK) was obtained from Enzyme Systems (Livermore, CA). Acetyl-Asp-Glu-Val-Asp-7-amino-4-methylcoumarin (Ac-DEVD-AMC) was obtained from Bachem (King of Prussia, PA); fluorescein isothiocyanate conjugated to VAD-FMK (FITC-VAD-FMK) was purchased from Promega (Madison, WI); antibodies to PKC $\delta$ , PKC $\alpha$ , PKC $\beta$ I, and PKC $\beta$ II were purchased from Santa Cruz Biotechnology (Santa Cruz, CA), cytochrome C (mouse monoclonal) from PharMingen (San Diego, CA), green fluorescent protein (GFP) (mouse monoclonal) from Clontech (Palo Alto, CA), and  $\beta$ -actin (mouse monoclonal) from Sigma (St. Louis, MO). ECL chemiluminescence kit was purchased from Amersham Pharmacia Biotech (Piscataway, NJ). PC12 cells were purchased from American Type Culture Collection (ATCC) (Rockville, MD), and immortalized rat mesencephalic dopaminergic neuronal cell line (1RB<sub>3</sub>AN<sub>27</sub>) was a kind gift of Dr. Kedar N. Prasad (University of Colorado Health Sciences Center, Denver, CO). Hydroethidine and Hoechst 33342 were purchased from Molecular Probes (Eugene, OR). Cell Death Detection ELISA Plus assay kit was purchased from Roche Molecular Biochemicals (Indianapolis, IN). PKC $\delta$  catalytic fragment, acridine orange, histone H1,  $\beta$ -glycerophosphate, superoxide dismutase (SOD), ATP, Protein-A-Sepharose, phosphatidylserine, and dioleoylglycerol were purchased from Sigma. Mn(III)tetrakis(4-Benzoic acid)porphyrin chloride (MnTBAP) was purchased from Oxis Health Products (Portland, OR). [ $\gamma$ -<sup>32</sup>P]ATP was purchased from NEN (Boston, MA). Bradford protein assay kit was purchased from Bio-Rad (Hercules, CA). Lipofectamine Plus reagent, Roswell Park Memorial Institute (RPMI)-1640 medium, horse serum, fetal bovine serum, L-glutamine, penicillin, streptomycin, and PCEP4 plasmid were purchased from Invitrogen (Gaithersburg, MD). BioPORTER, protein delivery reagent was purchased from Gene Therapy Systems (San Diego, CA), and plasmids PKC $\delta$ <sup>K376R</sup>-GFP fusion protein and pEGFP-N1 were kind gifts of Dr. Stuart Yuspa (National Cancer Institute, Bethesda, MD).

**Cell culture.** PC12 (ATCC CRL1721) cells were grown in RPMI medium supplemented with 10% horse serum, 5% fetal bovine serum, 1% L-glutamine, penicillin (100 U/ml), and streptomycin (100 U/ml) and maintained at 37°C in a humidified atmosphere of 5% CO<sub>2</sub>. Immortalized rat mesencephalic cells (1RB<sub>3</sub>AN<sub>27</sub>) were grown in RPMI medium supplemented with 10% fetal bovine serum, 1% L-glutamine, penicillin (100 U/ml), and streptomycin (100 U/ml), maintained at 37°C in a humidified atmosphere of 5% CO<sub>2</sub> (Prasad et al., 1998).

**Stable transfection.** Plasmid pPKC $\delta$ <sup>K376R</sup>-GFP encodes protein kinase C $\delta$ -GFP fusion protein, the number K376R refers to the mutation of lysine residue at position 376 to arginine in the catalytic site of PKC $\delta$  rendering it inactive (Li et al., 1999). Plasmid pEGFP-N1 encodes the green fluorescent protein alone and used as vector control. pEGFP-N1 and pPKC $\delta$ <sup>K376R</sup> were transfected into 1RB<sub>3</sub>AN<sub>27</sub> cells using Lipofectamine Plus reagent according to the procedure recommended by the manufacturer. In brief, 8  $\mu$ g of DNA, 24  $\mu$ l of lipid, and 24  $\mu$ l of Plus reagent were used to transfect 1RB<sub>3</sub>AN<sub>27</sub> cells in 100 mm tissue culture dishes at 50% confluency in 4 ml of culture medium without serum. Fresh medium containing serum was added 3 hr later. For stable cell lines, the 1RB<sub>3</sub>AN<sub>27</sub> cells were selected in 400  $\mu$ g/ml hygromycin, 48 hr after cotransfection with PCEP4 plasmid, which confers hygromycin resistance. Colonies were isolated with trypsin and glass cloning cylinders, and they were then replated and grown to confluence in T75 flasks. Subsequently, the stable cell lines were maintained in 200  $\mu$ g/ml hygromycin.

**Treatment paradigm.** After 2–4 d in culture, PC12 cells and 1RB<sub>3</sub>AN<sub>27</sub> were harvested and resuspended in serum-free growth medium at a cell density of 1–3  $\times$  10<sup>6</sup>/ml. Cell suspensions were treated with varying concentrations of MMT (30–500  $\mu$ M) over a period of 0.5–5 hr at 37°C. In inhibitor studies SOD (ROS inhibitor, 100 U/ml), MnTBAP (ROS inhibitor, 10  $\mu$ M), rottlerin (PKC $\delta$  inhibitor, 5–20  $\mu$ M), Ac-DEVD-CHO (caspase-3-specific inhibitor, 100–300  $\mu$ M), Z-DEVD-FMK (caspase-3-specific inhibitor, 10–50  $\mu$ M), or Z-VAD-FMK (a broad spectrum caspase inhibitor, 30–100  $\mu$ M) were added 30–90 min before the addition of MMT. The reaction samples were removed at 0.25, 0.5, 1, 2, 3, and 5 hr, then spun at 200  $\times$  g, and after 5 min, the cell pellets were used for assessing cytochrome C release, caspase-3, caspase-8, and caspase-9 enzymatic activities, extent of PKC $\delta$  cleavage, and DNA fragmentation. Dimethylsulfoxide (DMSO) (0.5–1%) was used as a vehicle in control experiments.

**Lactate dehydrogenase assay.** Lactate dehydrogenase (LDH) activity in the cell-free extracellular supernatant was quantified as an index of cell death (Vassault, 1983). We modified the original method to a 96-well format (Kitazawa et al., 2001). Briefly, PC12 cells were plated in 96-well plate, and after treatment 10  $\mu$ l of the extracellular supernatant was added to 200  $\mu$ l of 0.08 M Tris buffer, pH 7.2, containing 0.2 M NaCl, 0.2 mM NADH, and 1.6 mM sodium pyruvate. LDH activity was measured continuously by monitoring the decrease in the rate of absorbance at 339 nm using a microplate reader (Molecular Devices, Sunnyvale, CA), and the temperature was maintained at 37°C during reading. Changes in absorbance per minute ( $\Delta A/\Delta T$ ) were used to calculate LDH activity ( $U/I$ ), using the following equation:  $U/I = (\Delta A/\Delta T) \times 9682 \times 0.66$ , where 9682 was a coefficient factor, and 0.66 was a correction factor at 37°C.

**Detection of reactive oxygen species and lipid peroxidation by flow cytometry.** Flow cytometry analysis was performed on a Becton Dickinson (San Francisco, CA) FACScan flow cytometer. Hydroethidine, a sodium borohydride-reduced derivative of ethidium bromide, is used to detect ROS produced specifically inside the cell (Narayanan et al., 1997). When hydroethidine is loaded in the cells, it binds to cellular macromolecules. Once O<sub>2</sub><sup>-</sup> is generated, it converts hydroethidine to ethidium bromide and increases red fluorescence (620 nm). A 15 mW air-cooled argon-ion laser was used as an excitation source for hydroethidine at 488 nm, and the optical filter was 585/42 nm bandpass. Cells were detected and distinguished from the background by forward-angle light scattering and orthogonal light scattering characteristics. All the flow cytometric data were analyzed by Cellquest data analysis software to determine the significant increase or decrease of fluorescence intensity.

PC12 cells and engineered 1RB<sub>3</sub>AN<sub>27</sub> cells expressing kinase inactive PKC $\delta$  protein were resuspended with HBSS with 2 mM calcium at a density of 0.5  $\times$  10<sup>6</sup> cells/ml. Cells were then incubated with 10  $\mu$ M hydroethidine for 15 min at 37°C in the dark to allow dye loading into the cells. After incubation with dye, excess dye was removed, and the cells were resuspended with HBSS. After addition of MMT (30–500  $\mu$ M) ROS generation was measured at 0, 5, 15, 30, and 45 min after the exposure. In inhibitor studies, cells were incubated with SOD (100 U/ml) and MnTBAP (10  $\mu$ M) 10–30 min before MMT exposure.

**Quantification of cytochrome C release.** Cytochrome C release was quantified using a recently developed ELISA kit developed by MBL (Watertown, MA). This is a fast, highly sensitive and reliable assay for the detection of early changes in cytochrome C levels. Briefly, after 2–4 d in culture, PC12 cells were harvested and resuspended in serum-free growth medium at a cell density of 5  $\times$  10<sup>6</sup>/ml. Cell suspensions were exposed to 200 and 500  $\mu$ M MMT for 15–30 min at 37°C. After treatment the cells were spun at 200  $\times$  g, and after 5 min, washed once with 1 $\times$  ice-cold PBS and resuspended in 1 ml of ice-cold homogenization buffer (10 mM Tris HCl, pH 7.5, 0.3 M sucrose, 1 mM phenylmethylsulfonyl fluoride, 25  $\mu$ g/ml aprotinin, and 10  $\mu$ g/ml leupeptin) and homogenized on ice. Cells were then centrifuged for 10,000  $\times$  g for 60 min at 4°C. The resulting supernatants were collected as cytoplasmic fraction and used for cytochrome C release measurements. The MBL ELISA kit measures cytochrome C by one-step sandwich ELISA. The assay uses affinity-purified two polyclonal antibodies against cytochrome C. The cytoplasmic fractions were incubated with peroxidase conjugated anti-cytochrome C polyclonal antibody in the 96-well microtiter for 60 min at room temperature (RT). After washing with buffer (provided with the kit), the peroxidase substrate is mixed with the chromogen and allowed to incubate for an additional 15 min. An acid solution provided with the kit is then added to each well to terminate the enzyme reaction and to stabilize the developed color. The optical density of each well is then measured at 450 nm using a microplate reader. The concentration of

cytochrome C is calibrated from a standard curve based on reference standards.

**Confocal analysis of *in situ* caspase activity.** For this study, we used CaspACE kit (Promega) to label PC12 cells. The kit uses FITC-VAD-FMK, an FITC conjugate of the cell-permeable caspase inhibitor Z-VAD-FMK, which binds to activated caspase and serves as an *in situ* marker for apoptosis. The experiment was performed as per the manufacturer's protocol with slight modifications. Briefly, PC12 cells were grown on laminin (5  $\mu$ g/ml)-coated slides for 2–3 d in a 37°C, 5% CO<sub>2</sub> incubator. Cells were then exposed to 200  $\mu$ M MMT for 1 hr in the dark. After exposure, the cells were treated with 10  $\mu$ M FITC-VAD-FMK for 20 min at 37°C. Cells were then rinsed with 1 $\times$  PBS and fixed in 10% buffered formalin for 30 min at RT in the dark. After fixing, the cells were washed three times with PBS to remove formalin and then mounted with medium and coverslips, and observed under a Leica TCS-NT confocal microscope (Leica Microsystems Inc., Exton, PA).

**Enzymatic assay for caspases.** Caspase-3, caspase-8, and caspase-9 activities were performed as previously described by Yoshimura et al. (1998). Briefly, after treatment cells were spun and the cell pellets were lysed with Tris buffer, pH 7.4 (50 mM Tris HCl, 1 mM EDTA, and 10 mM EGTA) containing 10  $\mu$ M digitonin for 20 min at 37°C. Lysates were centrifuged at 900  $\times$  g for 3 min, and the resulting supernatants were incubated with specific fluorogenic caspase substrates at 37°C for 1 hr. Ac-DEVD-AMC (50  $\mu$ M), Ac-IETD-AMC (50  $\mu$ M), and Ac-LEHD-AMC (50  $\mu$ M) were used as substrates for determining caspase-3-, caspase-8-, and caspase-9-like protease activities, respectively. Levels of cleaved (active) caspase substrate were monitored at excitation  $\lambda$  380 nm and emission  $\lambda$  460 nm using a fluorescence plate reader (model: Fluoroskan-11; Titertek). Caspase activities were expressed as fluorescence units per milligram of protein per hour. The protein concentrations were determined using the Bio-Rad protein assay kit.

**Isolation of cytoplasmic fractions.** After incubation, the PC12 cells were spun at 200  $\times$  g for 5 min. Cell pellets were then washed once with ice-cold Ca<sup>2+</sup>-free PBS saline and resuspended in 2 ml of homogenization buffer (20 mM Tris-HCl, pH 8.0, 10 mM EGTA, 2 mM EDTA, 2 mM dithiothreitol, 1 mM phenylmethylsulfonyl fluoride, 25  $\mu$ g/ml aprotinin, and 10  $\mu$ g/ml leupeptin). The suspensions were sonicated for 10 sec, and centrifuged at 100,000  $\times$  g for 1 hr at 4°C. The supernatants were collected as cytosolic fractions. Protein concentration of each sample was determined, and the SDS-gel electrophoresis was performed as described below.

**Western blotting.** Cytoplasmic fractions containing equal amounts of protein were loaded in each lane and separated on a 10–12% SDS-polyacrylamide gel. Proteins were then transferred to nitrocellulose membrane by electroblotting for 75–90 min at 100 V. Nonspecific binding sites were blocked by treating the nitrocellulose membranes with 5% nonfat dry milk powder for 2 hr before treatment with primary antibodies. The nitrocellulose membranes containing the proteins were incubated with primary antibodies for 1 hr at RT with antibody directed against PKC $\delta$  (1:2000 dilution), PKC $\alpha$  (1:2000 dilution), cytochrome C (1:2000 dilution), or GFP (1:1000 dilution). The primary antibody treatments were followed by treatment with secondary HRP-conjugated anti-rabbit or anti-mouse IgG (1:2000 dilution) for 1 hr at RT. Secondary antibody-bound proteins were detected using an ECL chemiluminescence kit (Amersham). To confirm equal protein loading, blots were reprobbed with a  $\beta$ -actin antibody (1:5000 dilution). Gel photographs were taken with a gel imaging system and quantification of bands was performed using the imaging software from Scion Corp. (Frederick, MD).

**Immunoprecipitation kinase assays.** PKC $\delta$  enzymatic activity was assayed using an immunoprecipitation kinase assay as described by Reylan et al. (1999) and Vancurova et al. (2001). Briefly, after treatment with MMT, PC12 cells were washed once with PBS and resuspended in 1 ml of PKC lysis buffer (25 mM HEPES, pH 7.5, 20 mM  $\beta$ -glycerophosphate, 0.1 mM sodium orthovanadate, 0.1% Triton X-100, 0.3 M NaCl, 1.5 mM MgCl<sub>2</sub>, 0.2 mM EDTA, 0.5 mM DTT, 10 mM NaF, and 4  $\mu$ g/ml each aprotinin and leupeptin). In inhibition experiments, cells were pretreated with 10  $\mu$ M rottlerin before the addition of 200  $\mu$ M MMT. The cell lysates were allowed to sit on ice for 30 min and centrifuged at 13,000  $\times$  g for 5 min, and the supernatants were collected as cytosolic fraction. Protein concentration was determined using a Bradford assay. Cytosolic protein (0.25–0.5 mg) was immunoprecipitated overnight at 4°C using 2  $\mu$ g of anti-PKC $\delta$ , anti-PKC $\alpha$ , anti-PKC $\beta$ I, or anti-PKC $\beta$ II antibodies. The immunoprecipitates were then incubated with Protein-A Sepharose (Sigma) for 1 hr at 4°C. The protein A bound antigen-antibody complexes were then washed three times with PKC

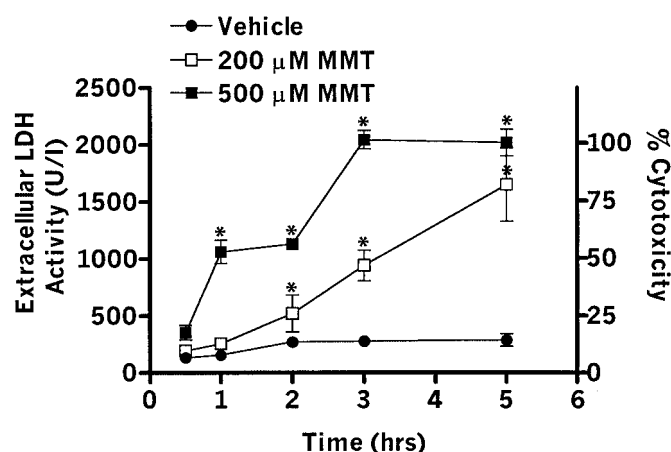
lysis buffer, three times with 2 $\times$  kinase buffer (40 mM Tris, pH 7.4, 20 mM MgCl<sub>2</sub>, 20  $\mu$ M ATP, and 2.5 mM CaCl<sub>2</sub>), and resuspended in 20  $\mu$ l of 2 $\times$  kinase buffer. Reaction was started by adding 20  $\mu$ l of reaction buffer containing 0.4 mg Histone H1, 50  $\mu$ g/ml phosphatidylserine, 4.1  $\mu$ M dioleoylglycerol, and 5  $\mu$ Ci of [ $\gamma$ -<sup>32</sup>P] ATP (3000 Ci/mM) to the immunoprecipitated samples and incubated for 10 min at 30°C. SDS gel-loading buffer (2 $\times$ ) was added to terminate the reaction, the samples were boiled for 5 min, and the products were separated on a 12.5% SDS-PAGE gel. For *in vitro* inhibition of PKC $\delta$  kinase activity, 5–20  $\mu$ M rottlerin was added to 200  $\mu$ M MMT-treated immunoprecipitated sample 15 min before the addition of 2 $\times$  reaction buffer containing 5  $\mu$ Ci of [ $\gamma$ -<sup>32</sup>P] ATP (3000 Ci/mmol). The H1 phosphorylated bands were detected using a Personal Molecular Imager (FX model; Bio-Rad), and quantification was done using Quantity One 4.2.0 software.

**Intracellular delivery of PKC $\delta$  catalytic fragment.** Intracellular delivery of PKC $\delta$  fragment was performed using a recently developed lipid-mediated delivery system (BioPORTER; Gene Therapy Systems, San Diego, CA). This is a fast and reliable procedure that delivers proteins in a functionally active form into the cytoplasm of cells (Zelphati et al., 2001). The protein delivery system is composed of a new trifluoroacetylated lipopolyamine (TFA-DODAPL) and dioleoyl phosphatidylethanolamine. This cationic formulation has recently been used for delivery of various bioactive molecules, including antibodies, enzymes (caspase-3, caspase-8,  $\beta$ -galactosidase, and granzyme B), cytochrome C, dextran sulfates, phycobiliproteins, and albumins into the cytoplasm of numerous adherent and suspension cells (Zelphati et al., 2001). Active PKC $\delta$  catalytic fragments were delivered into cells using the protein delivery reagent by following the manufacturer's protocol. PC12 cells ( $\sim$ 1–2  $\times$  10<sup>5</sup> cells/ml) were subcultured in 24-well tissue culture plate for 24 hr. PKC $\delta$  catalytic fragment (5 ng) was mixed with 3  $\mu$ l of protein delivery reagent and 300  $\mu$ l of serum-free DMEM media and added to each well. The cells were incubated at 37°C for 4 hr. Cells were then lysed, and caspase-3 activity measured as described above. Heat-inactivated PKC $\delta$  catalytic fragment was used as negative control and inactivation was performed by incubating the active PKC $\delta$  fragment at 95°C for 15 min. The delivery efficiency was  $\sim$ 70% in PC12 cells as determined using a FITC-tagged antibody control (supplied with the assay kit). Also, the protein delivery system produced no significant cytotoxic response as measured by Trypan blue dye exclusion method.

**In situ assessment of apoptosis.** To assess nuclear morphology and DNA damage, we stained the cells with fluorescent DNA-binding dyes acridine orange and Hoechst 33342. Acridine orange, a useful probe for detecting apoptotic cells, exhibits metachromatic fluorescence that is sensitive to DNA conformation. Apoptotic cells stained with acridine orange show reduced green and enhanced red fluorescence in comparison with normal cells (Pulliam et al., 1998). Briefly, PC12 cells were grown on laminin (5  $\mu$ g/ml)-coated slides for 2–3 d in a 37°C, 5% CO<sub>2</sub> incubator. Cells were washed twice with PBS and after 1 hr treatment with 200  $\mu$ M MMT, the cells were incubated with 10  $\mu$ M acridine orange for 15 min at RT in the dark. The cells were again washed with PBS, mounted with coverslips, and observed under a Nikon DiaPhot microscope, and pictures were captured with a SPOT digital camera (Diagnostic Instruments, Sterling Heights, MI).

Morphological changes associated with apoptosis were also assessed by staining with Hoechst 33342. Cells stained with Hoechst 33342 dye fluoresce bright blue after binding to DNA in the nucleus. The nucleus of apoptotic cells exhibit strong blue staining and staining pattern is heterogeneous and occurs in patches, indicative of chromatin condensation, whereas the nucleus of nonapoptotic cells exhibit more diffused, weak and homogenous staining (Shimizu et al., 1996; Du et al., 1997). Briefly, PC12 cells were plated on collagen (6  $\mu$ g/cm<sup>2</sup>)-coated cover slides and treated with 200  $\mu$ M MMT. After 1 hr of exposure, the cells were fixed with 10% buffered formaldehyde for 30 min at room temperature and stained with Hoechst 33342 (10  $\mu$ g/ml) for 3 min in dark. The cells were again washed three times with PBS, mounted with coverslips, and observed under a Nikon DiaPhot microscope under UV illumination, and pictures were captured with a SPOT digital camera (Diagnostic Instruments).

**Quantification assay for DNA fragmentation.** DNA fragmentation assay was performed using a recently developed Cell Death Detection ELISA Plus assay kit. This is a fast, highly sensitive and reliable assay for the detection of early changes in apoptotic cell death and measures the appearance and amount of histone-associated low molecular weight DNA in the cytoplasm of cells. This assay has been recently used in quantitation of apoptosis because of its reliability and high sensitivity



**Figure 1.** MMT exposure induces cell death. PC12 cells were exposed to 200 and 500  $\mu\text{M}$  of MMT for 0.5–5 hr at 37°C. After the exposure, cell-free extracellular supernatants were collected, and LDH activity was measured by spectrophotometer. Values represent mean  $\pm$  SEM for three to five separate experiments in triplicate. Significance was determined by ANOVA followed by Dunnett's post-test between the vehicle-treated group and each treatment group (\* $p < 0.05$ ).

(Reyland et al., 1999). Briefly, PC12 and engineered 1RB<sub>3</sub>AN<sub>27</sub> cells were exposed to 200–500  $\mu\text{M}$  MMT for 1–3 hr. In inhibitor studies, SOD (100 U/ml), MnTBAP (10  $\mu\text{M}$ ), rottlerin (10  $\mu\text{M}$ ), Z-DEVD-FMK (50  $\mu\text{M}$ ), or Z-VAD-FMK (100  $\mu\text{M}$ ) were treated for 30 min at 37°C before the addition of MMT. After MMT treatment, cells were spun down at 200  $\times g$  for 5 min and washed once with 1 $\times$  PBS. Cells were then incubated with a lysis buffer (supplied with the kit) at RT. After 30 min, samples were centrifuged, and 20  $\mu\text{l}$  aliquots of the supernatant were then dispensed into streptavidin-coated 96-well microtiter plates followed by addition of 80  $\mu\text{l}$  of antibody cocktail and incubated for 2 hr at RT with mild shaking. The antibody cocktail consisted of a mixture of anti-histone biotin and anti-DNA-HRP directed against various histones and antibodies to both single-stranded DNA and dsDNA, which are major constituents of the nucleosomes. After incubation, unbound components were removed by washing with the incubation buffer supplied with the kit. Quantitative determination of the amount of nucleosomes retained by anti-DNA-HRP in the immunocomplex was determined spectrophotometrically with 2,2'-azino-di-(3-ethylbenzthiazoline sulfonate (6)) diammonium salt (ABTS) as an HRP substrate (supplied with the kit). Measurements were made at 405 nm against an ABTS solution as a blank (reference wavelength  $\sim$ 490 nm) using a Molecular Devices Spectramax Microplate Reader.

**Data analysis.** Data analysis was performed using Prism 3.0 software (GraphPad Software, San Diego, CA). Data from caspase enzymatic activities and DNA fragmentation assays were first analyzed using one-way ANOVA. Neuman-Keuls or Dunnett's post-tests were then performed to compare control with MMT-treated groups, and differences with  $p < 0.05$  were considered significant. For individual comparisons, student  $t$  test or Welch's corrected  $t$  test where appropriate was used.

## RESULTS

### MMT-induced cytotoxicity

PC12 cells were exposed to 200 and 500  $\mu\text{M}$  MMT for varying amounts of time. The amount of LDH released into the extracellular media was measured as an index of cytotoxicity (Vassault, 1983; Kanthasamy et al., 1995). Extracellular LDH activity showed a dose- and time-dependent increase with MMT treatment ranging from 2- to 20-fold over the control group (Fig. 1). For example, exposure to 200  $\mu\text{M}$  MMT resulted in 3-, 10-, and 17-fold increase in LDH release over untreated cells at 1, 3, and 5 hr, respectively. To determine the percent cell death, total LDH content of untreated cells ( $\sim$ 2000 U/I) was normalized to 100%. PC12 cells exposed to 200  $\mu\text{M}$  MMT for 3 hr produced  $\sim$ 50% cell death. Similarly, exposure to 500  $\mu\text{M}$  MMT resulted in a signifi-

cant ( $p < 0.05$ ) toxicity in MMT-treated groups as compared with vehicle-treated cells at all time points. There were no significant differences in LDH release between vehicle-treated and untreated PC12 cells during the 5 hr exposure. These data suggest that MMT induces a dose- and time-dependent cell death in dopamine-producing cells.

### Generation of ROS after MMT treatment

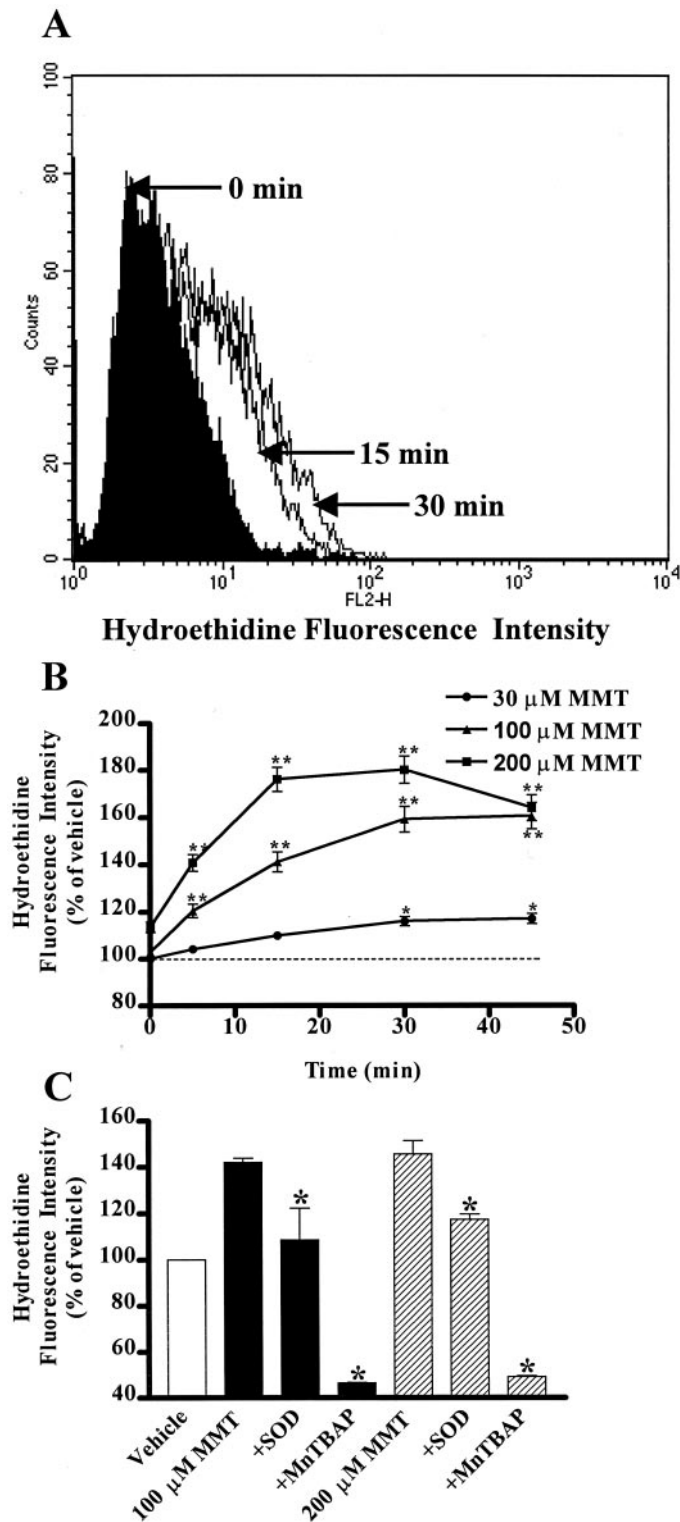
Flow cytometric analysis using the ROS-sensitive fluorescence probe hydroethidine revealed that MMT treatment induces ROS generation. Figure 2*A* depicts a representative flow cytometric histogram of 200  $\mu\text{M}$  MMT-treated PC12 cells exhibiting time-dependent increases in red fluorescence. MMT treatment increased ROS production in a dose- and time-dependent manner (Fig. 2*B*). For example, a 15 min exposure to 30, 100, and 200  $\mu\text{M}$  MMT resulted in a 10, 41, and 76% increase in ROS production, respectively. Exposure to 200  $\mu\text{M}$  MMT resulted in 40, 76, and 80% increase in ROS production over vehicle treatment at 5, 15, and 30 min, respectively. The time course study also revealed that 500  $\mu\text{M}$  MMT treatment induced a rapid and dramatic increase in ROS generation ( $>200\%$  of control) within 5 min and then the response rapidly declined over time (data not shown). Pretreatment with SOD (100 U/ml) or MnTBAP (SOD mimetic, 10  $\mu\text{M}$ ) significantly ( $p < 0.05$ ) reduced MMT-induced ROS production, indicating that MMT predominantly generates superoxide species (Fig. 2*C*).

### Accumulation of cytochrome C in the cytosol after MMT treatment

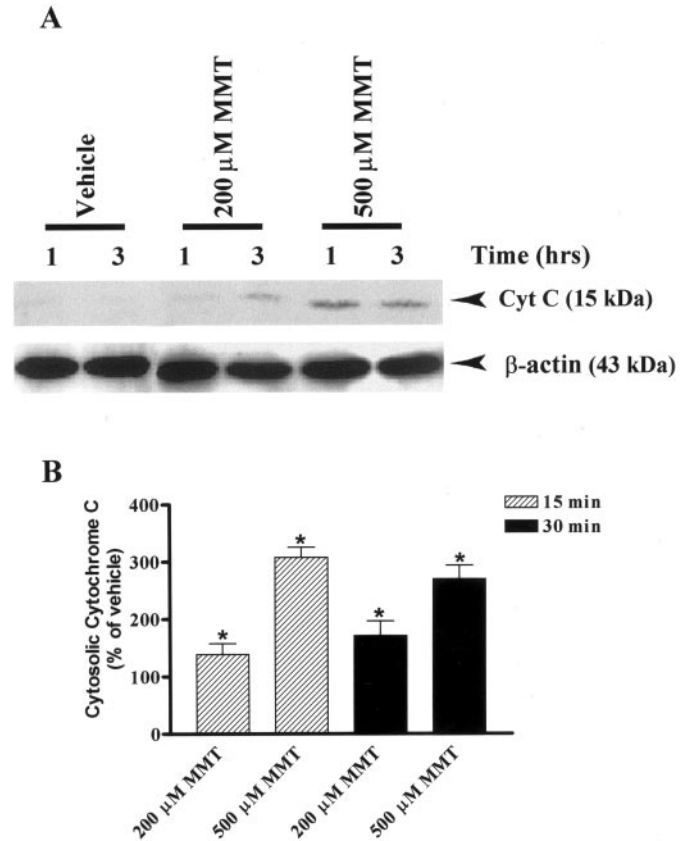
ROS production in the cells is known to activate many cellular factors including cytochrome C, which subsequently triggers apoptotic cell death (Tan et al., 1998; Cassarino et al., 1999). Release of cytochrome C from the mitochondria into the cytoplasm is an early event that occurs during programmed cell death (Muller-Hocker, 1992; Crompton, 1999), and therefore we determined whether MMT induces release of cytochrome C in PC12 cells. Figure 3*A* shows a time-dependent increase of cytochrome C in the cytoplasmic fractions of PC12 cells treated with MMT. No detectable levels of cytochrome C were detected in the cytosol of vehicle (DMSO)-treated cells up to 3 hr, whereas a profound release of cytochrome C was observed as early as 1 hr in MMT-treated cells. Nitrocellulose membranes were reprobed with  $\beta$ -actin antibody, and the density of 43 kDa  $\beta$ -actin bands was identical in all lanes confirming equal protein loading. To further accurately quantify how soon cytochrome C is released, we used a highly sensitive cytochrome C ELISA sandwich assay. MMT exposure resulted in a dose-dependent increase in cytosolic cytochrome C as early as 15 min (Fig. 3*B*). A 15 min exposure to 200 and 500  $\mu\text{M}$  MMT resulted in an increase in cytosolic cytochrome C by 40 and 200% over the vehicle-treated group, respectively, and a 30 min exposure resulted in an increase in cytosolic cytochrome C by 70 and 170%, respectively. The reason for the decrease in amount of cytochrome C released at 30 min after 500  $\mu\text{M}$  MMT exposure might be attributed to loss of cellular integrity caused by the observed necrotic cell death (Fig. 1).

### Activation of caspase-3 and caspase-9 but not caspase-8 after MMT treatment

Because the release of cytochrome C is known to activate a group of cysteine proteases, namely caspases (Cohen, 1997; Earnshaw et al., 1999; Schultz and Andreasen, 1999; Jellinger, 2000), we examined whether caspase-8, caspase-9, and caspase-3 are activated during MMT exposure. PC12 cells exposed to MMT showed a



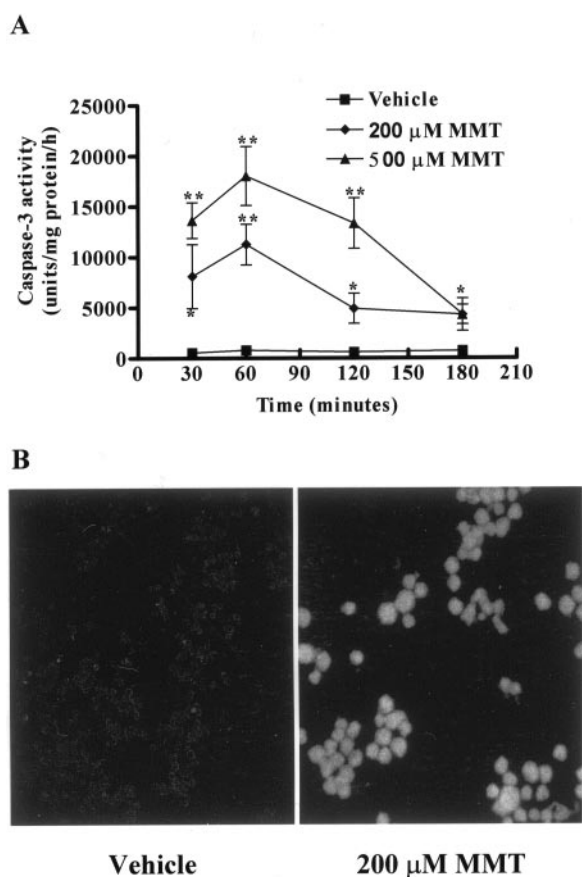
**Figure 2.** MMT treatment generates ROS in PC12 cells. PC12 cells were suspended in HBSS supplemented with 2 mM Ca<sup>2+</sup> at a density of 0.5–0.75  $\times$  10<sup>6</sup> cells/ml. A concentration of 10  $\mu$ M hydroethidine was added to the cells and incubated for 15 min at 37°C in the dark. *A*, Time-dependent change in hydroethidine fluorescent intensity in PC12 cells treated with MMT. A concentration of 200  $\mu$ M MMT was added, and fluorescent intensity was measured at 0, 15, and 30 min by flow cytometry as described in Materials and Methods. The data are a representative flow cytometric histogram of MMT-treated PC12 cells exhibiting a time-dependent increase in red fluorescence. *B*, Dose- and time-dependent increase in ROS production. Various doses of MMT were added, and



**Figure 3.** Dose- and time-dependent accumulation of cytosolic cytochrome C in MMT-treated PC12 cells. *A*, Western blot. *B*, Cytochrome C ELISA assay. *A*, Subconfluent cultures of undifferentiated PC12 cells were harvested at 1 and 3 hr after treatment with 200 or 500  $\mu$ M MMT. The cytosolic fractions were obtained as described in Materials and Methods. Cytosolic fractions were separated by 12% SDS-PAGE, transferred to a nitrocellulose membrane, and cytochrome C (*Cyt C*) was detected using polyclonal antibody raised against cytochrome C. For  $\beta$ -actin measurements, the membrane used for cytochrome C was reprobed with  $\beta$ -actin antibody to confirm equal protein loading in each lane. The immunoblots were visualized using ECL detection agents from Amersham. *B*, Subconfluent cultures of undifferentiated PC12 cells were harvested at 15 and 30 min after treatment with 200 or 500  $\mu$ M MMT. The cytosolic fractions were obtained as described in Materials and Methods. The value of each treatment group is the mean  $\pm$  SEM from two separate experiments in triplicate. Asterisks ( $*p < 0.05$ ) indicate significant differences compared with vehicle-treated cells.

significant increase in caspase-9 activity, however, no significant increase in caspase-8 enzyme activity was observed (data not shown). A 30 min exposure to 200 and 500  $\mu$ M MMT produced a threefold and twofold increase in caspase-9 activity, respectively. The lack of dose–response in caspase-9 enzymatic activity at 500

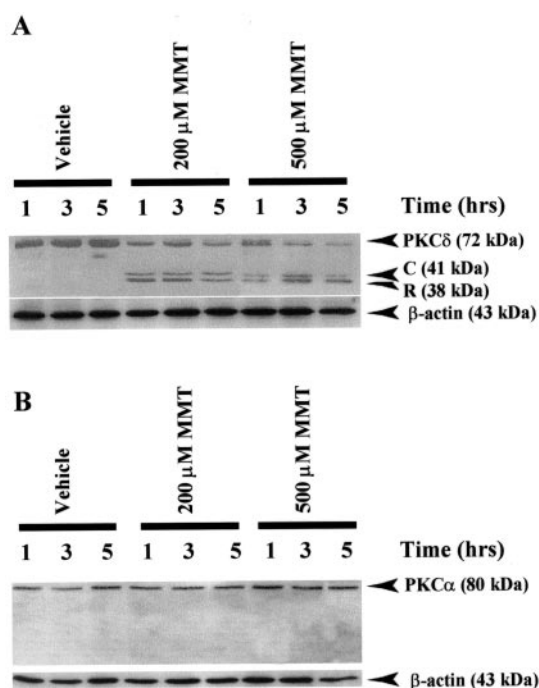
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fluorescent intensity was measured at 0, 5, 15, and 30 min. Data represent the mean  $\pm$  SEM of two to five separate experiments in triplicate. Asterisks ( $*p < 0.5$  and  $**p < 0.01$ ) indicate significant differences compared with the time-matched vehicle-treated cells. *C*, Effect of SOD and MnTBAP on ROS production. Cells were pretreated with ROS inhibitors, SOD (100 U/ml) and MnTBAP (10  $\mu$ M), and then exposed to 100 or 200  $\mu$ M MMT for 15 min. The value of each treatment group is the mean  $\pm$  SEM from two to three separate experiments performed in triplicate. Asterisks ( $*p < 0.05$ ) indicate significant differences compared with MMT-treated cells.



**Figure 4.** MMT treatment increases caspase-3 activity. *A*, Caspase-3 enzymatic activity. *B*, *In situ* caspase-3 activity. *A*, Subconfluent cultures of undifferentiated PC12 cells were harvested at 30 min, 1, 2, and 3 hr after MMT treatment. Caspase-3 activity was assayed using specific fluorogenic substrate, Ac-DEVD-AMC (50  $\mu$ M), as described in Materials and Methods. The data represent mean  $\pm$  SEM of nine individual measurements from three separate experiments. Asterisks (\*\* $p < 0.01$ ; \* $p < 0.05$ ) indicate significant differences compared with temporally matched vehicle (DMSO)-treated cells. *B*, PC12 cells were grown on laminin-coated slides for 2–3 d and then exposed to 0.5% DMSO (vehicle) and 200  $\mu$ M MMT for 1 hr in the dark. After exposure, cells were treated with 10  $\mu$ M FITC-VAD-FMK (Promega caspACE, *in situ* marker for caspase-3 activity) and processed as described in Materials and Methods. Confocal images were obtained using a Leica TCS-NT microscope.

$\mu$ M MMT concentration is probably caused by acute cytotoxic effects of the toxic compound at higher doses.

MMT treatment in PC12 cells resulted in dramatic increase in caspase-3 enzymatic activity (Fig. 4). After exposure to 200  $\mu$ M MMT, caspase-3-specific activity was increased 12-, 17-, 7-, and 6-fold over the vehicle-treated groups at 0.5, 1, 2, and 3 hr after treatment, respectively (Fig. 4*A*). Similarly, exposure to 500  $\mu$ M MMT resulted in an increase in caspase-3 specific activity by 20-, 27-, 20-, and 6-fold over the vehicle-treated groups after 0.5, 1, 2, and 3 hr exposure, respectively. The overall pattern of the time course study of caspase-3 activity revealed a clear pattern of dramatic increase in enzyme activity peaking at 1 hr and then progressively decreasing over time, returning nearly to that of vehicle-treated cells at 3 hr. To further confirm the activation of caspase-3, *in situ* fluorometric analysis was performed using FITC-VAD-FMK in live cells. In these experiments, we found majority of PC12 cells were labeled within 1 hr of exposure to 200  $\mu$ M MMT, indicating a profound increase in caspase-3 activity *in situ*, whereas no labeling was seen in vehicle-treated cells (Fig. 4*B*).

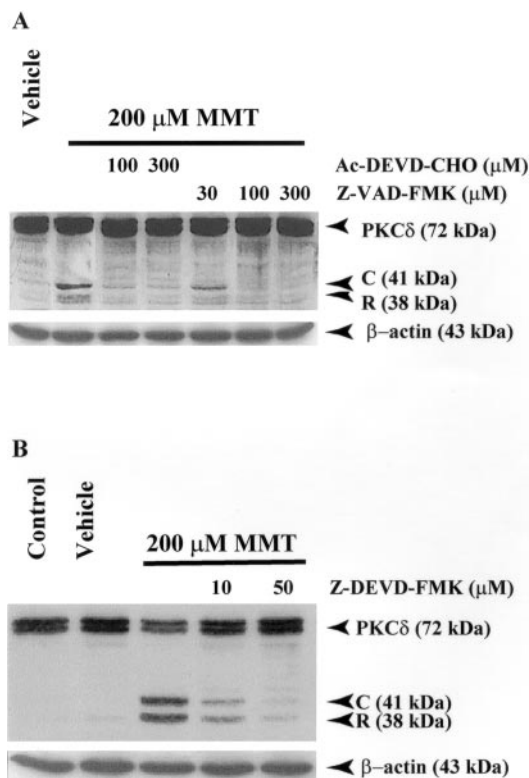


**Figure 5.** Proteolytic cleavage of PKC $\delta$  but not of PKC $\alpha$  in MMT-treated PC12 cells. *A*, PKC $\delta$ ; *B*, PKC $\alpha$ . Subconfluent undifferentiated PC12 cells were harvested at 1, 3, and 5 hr after treatment of 200 or 500  $\mu$ M MMT. Cytosolic fractions were obtained as described in Materials and Methods, and were separated by 10% SDS-PAGE, transferred to nitrocellulose membrane, and PKC $\alpha$  and PKC $\delta$  were detected using antibodies directed against their catalytic subunits. To confirm equal protein loading in each lane, the membranes were reprobbed with  $\beta$ -actin antibody. The immunoblots were visualized using ECL detection agents from Amersham. C, PKC $\delta$  catalytic subunit; R, PKC $\delta$  regulatory subunit.

### Proteolytic cleavage of protein kinase C $\delta$ but not PKC $\alpha$ by MMT

Recent studies have indicated PKC $\delta$  to be one of the endogenous substrates for caspase-3, which cleaves the kinase to yield a 41 kDa catalytically active and a 38 kDa regulatory PKC $\delta$  fragments in non-neuronal cell lines, salivary gland acinar cells, (Reyland et al., 1999), rat fibroblasts (Dal Pra et al., 1999), and neutrophils (Pongracz et al., 1999). Because MMT exposure resulted in a profound activation of caspase-3, we decided to examine the proteolytic cleavage of PKC $\delta$  in MMT-treated PC12 cells. After treatment of PC12 cells with MMT at 37°C, we observed over a 5 hr period a significant proportion of native PKC $\delta$  (72–74 kDa) protein was proteolytically cleaved to yield 38 kDa regulatory and 41 kDa catalytically active fragments when immunoblotted with an antibody raised against PKC $\delta$  (Fig. 5*A*). The time course study revealed that almost all of the native PKC $\delta$  (72–74 kDa) protein was cleaved within 3 hr of incubation with MMT, evidenced by a reduction in the intensity of the native 72–74 kDa band and a concomitant increase in the catalytically active 41 kDa cleaved fragment. PC12 cells exposed to increasing concentrations of MMT (200 and 500  $\mu$ M) showed a dose-dependent cleavage of PKC $\delta$ . However, no cleavage of PKC $\delta$  was observed in vehicle-treated cells during the entire 5 hr experimental time period at the doses tested. Nitrocellulose membranes were reprobbed with  $\beta$ -actin antibody, and the density of 43 kDa  $\beta$ -actin band was identical in all lanes confirming equal protein loading.

MMT-induced proteolytic cleavage of PKC was also isoform-specific, because exposure of PC12 cells to 200 or 500  $\mu$ M MMT



**Figure 6.** Caspase-3 mediates the proteolytic cleavage of PKC $\delta$  in MMT-treated PC12 cells. *A*, Effect of Ac-DEVD-CHO and Z-VAD-FMK on PKC $\delta$  cleavage. *B*, Effect of Z-DEVD-FMK on PKC $\delta$  cleavage. Subconfluent undifferentiated PC12 cells were treated with 200  $\mu$ M MMT, with or without the inclusion of caspase inhibitors Ac-DEVD-CHO, Z-VAD-FMK, or Z-DEVD-FMK. Inhibitors were added 30 min before the addition of MMT. Cells were harvested 3 hr after the addition of MMT. The cytosolic fractions were obtained as described in Materials and Methods, and were analyzed by 10% SDS-PAGE and Western blot. To confirm equal protein loading in each lane, the membranes were reprobbed with  $\beta$ -actin antibody. C, PKC $\delta$  catalytic subunit; R, PKC $\delta$  regulatory subunit.

for up to 5 hr failed to induce proteolytic cleavage of PKC $\alpha$  (Fig. 5*B*). Membranes were reprobbed with  $\beta$ -actin antibody, and the density of 43 kDa  $\beta$ -actin band was identical in all lanes confirming equal protein loading. Additionally, MMT exposure did not result in the translocation of either PKC $\alpha$  or  $\delta$  from the cytoplasm to the membrane for their activation (data not shown).

### MMT-induced proteolytic cleavage of PKC $\delta$ is caspase-3-dependent

To further confirm that PKC $\delta$  cleavage is mediated by caspase-3, we used caspase-3-specific inhibitors Ac-DEVD-CHO (Fig. 6*A*), Z-DEVD-FMK (Fig. 6*B*) or a broad-spectrum caspase inhibitor, Z-VAD-FMK (Fig. 6*A*), to block the cleavage. Pretreatment of PC12 cells for 30 min with any of the three inhibitors used here before 3 hr exposure of cells to 200  $\mu$ M MMT prevented the appearance of the 41 kDa catalytically active PKC $\delta$  fragment, and effects of all three inhibitors were dose-dependent. Furthermore, Z-DEVD-FMK and Z-VAD-FMK appeared to be more potent in blocking PKC $\delta$  cleavage than Ac-DEVD-CHO. Membranes were reprobbed with  $\beta$ -actin antibody (Fig. 6*A,B*), and the density of 43 kDa  $\beta$ -actin bands was identical in all lanes confirming equal protein loading.

### Rottlerin blocks MMT-induced caspase-3 enzymatic activity: possible feedback activation of caspase-3 by PKC $\delta$

As reported above, we observed a dramatic increase in caspase-3 (6- to 27-fold) (Fig. 4*A*) enzymatic activity in MMT-treated PC12 cells at 1 hr after treatment. These results prompted us to determine the cause for the dramatic increase in MMT-induced caspase-3 activity, and so we investigated whether PKC $\delta$  is capable of activating caspase-3 by a positive feedback mechanism. To address this hypothesis, we tested the ability of PKC $\delta$  specific inhibitor rottlerin to modulate caspase-3 activity in MMT-treated PC12 cells by pre- and post-treatment. Pretreatment with rottlerin 30 min before the addition of 200  $\mu$ M MMT suppressed caspase-3 activity in a dose-dependent manner (Fig. 7*A*). Rottlerin at 5, 10, and 20  $\mu$ M suppressed MMT-induced caspase-3 activity by 38, 64, and 70%, respectively, whereas the basal caspase-3 activity was not altered by treatment with rottlerin alone. Post-treatment with rottlerin 30 min after the addition of 200  $\mu$ M MMT also suppressed caspase-3 activity in a dose-dependent manner (Fig. 7*B*). Rottlerin at 5 and 20  $\mu$ M suppressed MMT-induced caspase-3 activity by 60 and 98%, respectively, whereas the basal caspase-3 activity was unaltered by post-treatment with rottlerin alone. The extent of MMT-induced caspase-3 inhibition by 5  $\mu$ M rottlerin was not statistically significant ( $p > 0.05$ ) between pre- and post-treatments, whereas inhibition with 20  $\mu$ M rottlerin post-treatment was very significant ( $p < 0.01$ ) as compared with pre-treatment, reducing the caspase-3 activity to almost the basal level. Overall, these results indicate that there may be a positive feedback activation of caspase-3 by PKC $\delta$ , and this activation can be blocked by rottlerin.

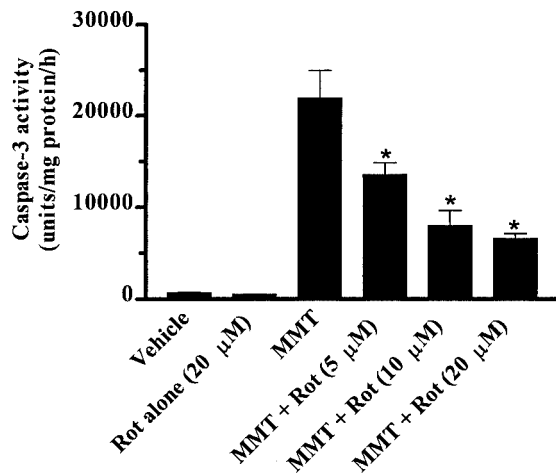
### Rottlerin blocks caspase-3 mediated proteolytic cleavage of PKC $\delta$

Because the pretreatment study with rottlerin blocked caspase-3 enzymatic activity, we further tested whether rottlerin pretreatment attenuates caspase-3-dependent proteolytic cleavage of PKC $\delta$ . Pretreatment with rottlerin before the addition of 200  $\mu$ M MMT prevented the accumulation of PKC $\delta$  cleavage product in a dose-dependent manner (Fig. 8). Rottlerin at 20  $\mu$ M markedly reduced the appearance of PKC $\delta$  cleavage product in PC12 cells exposed to 200  $\mu$ M MMT, indicating that the activation of PKC $\delta$  is essential to its cleavage by caspase-3. Membranes were reprobbed with  $\beta$ -actin antibody (Fig. 8), and the density of 43 kDa  $\beta$ -actin band was identical in all lanes, confirming equal protein loading.

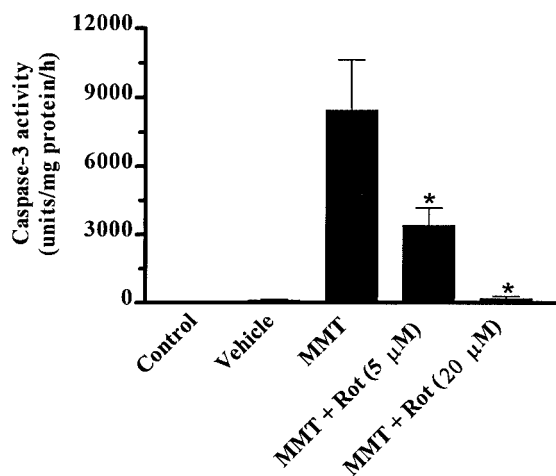
### Rottlerin inhibits MMT-induced increases in PKC $\delta$ kinase activity in PC12 cells

To determine whether the MMT induced caspase-3- and PKC $\delta$ -dependent accumulation of PKC $\delta$  cleaved product is attributed to an increase in PKC $\delta$  enzyme activity, we performed kinase assays in immunoprecipitated samples from cytosolic fractions using PKC $\delta$  specific polyclonal antibody and by examining the ability of PKC $\delta$  to phosphorylate histone H1. The enzymatic activity of PKC $\delta$  increased after 1 hr exposure to MMT in dose-dependent manner (Fig. 9*A*). Densitometric analysis of phosphorylated histone H1 bands revealed a three-fold and five-fold increase in protein kinase activity in cells exposed to 200 and 500  $\mu$ M MMT for 1 hr, respectively, and was coincident with generation of PKC $\delta$  cleavage. We attribute this increased kinase activity to the persistently active PKC $\delta$  catalytic fragment, because activation of

## A. Pre-treatment



## B. Post-treatment

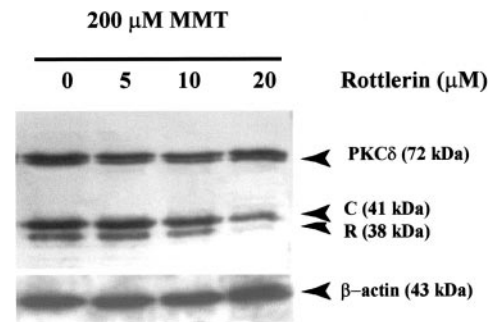


**Figure 7.** Suppression of caspase-3 activity by rottlerin in MMT-treated PC12 cells. *A*, Pre-treatment; *B*, post-treatment. Subconfluent undifferentiated PC12 cells were treated with 200  $\mu$ M MMT with or without the inclusion of rottlerin (Rot; 5–20  $\mu$ M) for 1 hr. Rottlerin was added 30 min before or 30 min after the addition of MMT. Caspase-3 activity was assayed using Ac-DEVD-AMC (50  $\mu$ M) as substrate, as described in Materials and Methods. The data represent an average of four to nine individual measurements from two or three separate experiments  $\pm$  SEM. Asterisk (\* $p$  < 0.05) indicates significant difference compared with cells exposed to 200  $\mu$ M MMT.

intact PKC $\delta$  by translocation to the membrane does not occur during MMT treatment (data not shown). There was no increase in kinase activity of PKC $\alpha$ , PKC $\beta$ I, and PKC $\beta$ II in immunoprecipitated samples of treated cells (data not shown) suggesting that the MMT-induced increase in kinase activity is isoform specific for PKC $\delta$ . Pretreatment with 10  $\mu$ M rottlerin resulted in 80% reduction in the kinase activity (Fig. 9*B*), suggesting that the activation of PKC $\delta$  is essential for MMT-induced increases in kinase activity, and this may be facilitated via the positive-feedback activation of caspase-3 by the catalytically active PKC $\delta$  fragment.

#### Rottlerin directly inhibits PKC $\delta$ kinase activity *in vitro*

Rottlerin was originally reported to inhibit PKC $\delta$  kinase activity by competing for the ATP-binding site (Gschwendt et al., 1994).



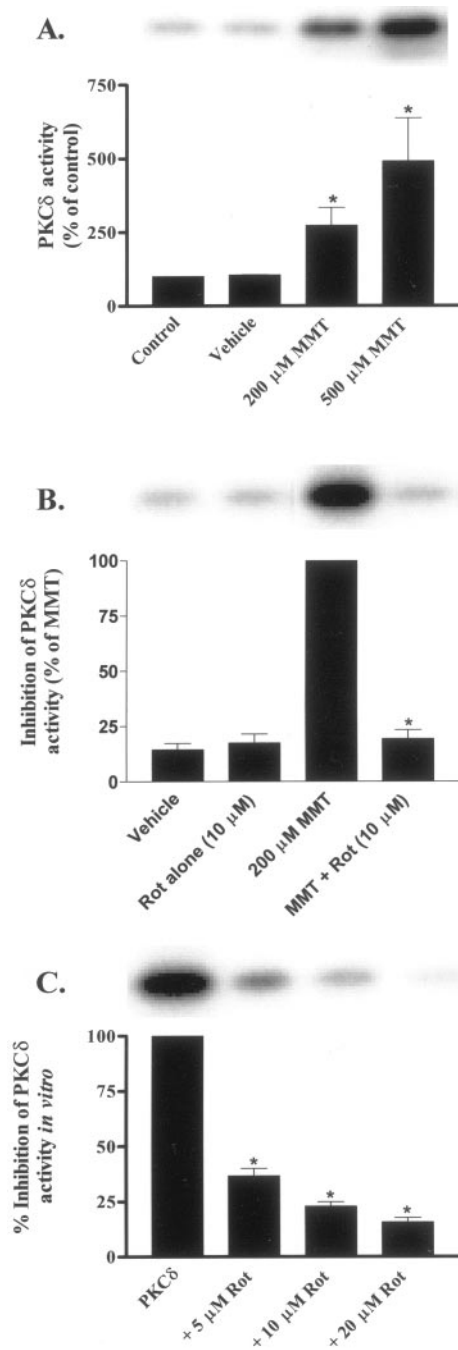
**Figure 8.** Rottlerin pretreatment blocks proteolytic cleavage of PKC $\delta$  in MMT-treated PC12 cells. Subconfluent undifferentiated PC12 cells were treated with 200  $\mu$ M MMT with or without the inclusion of rottlerin (5–20  $\mu$ M). Rottlerin was added 90 min before the addition of MMT. Cytosolic fractions were obtained as described in Materials and Methods and were separated by 10% SDS-PAGE, transferred to nitrocellulose membrane, and PKC $\delta$  was detected using an antibody directed against its catalytic subunit. The immunoblots were visualized using ECL detection agents from Amersham. To confirm equal protein loading in each lane, the membranes were re probed with  $\beta$ -actin antibody. C, PKC $\delta$  catalytic subunit; R, PKC $\delta$  regulatory subunit.

This inhibitor has been used to implicate PKC $\delta$  in a variety of cellular events, including apoptosis (Chen et al., 1999; Reyland et al., 1999; Dempsey et al., 2000; Way et al., 2000; Basu et al., 2001; Vancurova et al., 2001). To further confirm the inhibitory potency of rottlerin on PKC $\delta$  activity, we tested various concentrations of rottlerin on PKC $\delta$  enzyme activity using an *in vitro* kinase assay. PKC $\delta$  was immunoprecipitated from MMT-treated cytosolic fractions using PKC $\delta$  specific polyclonal antibody and incubated with rottlerin *in vitro* for 15 min before the addition of histone H1 and [ $^{32}$ P]ATP. For the *in vitro* reaction, we used same rottlerin concentrations that blocked MMT-stimulated PKC $\delta$  kinase activity in intact PC12 cells. Rottlerin at 5, 10, and 20  $\mu$ M inhibited PKC $\delta$  activity *in vitro* by 63, 77, and 84%, respectively (Fig. 9*C*), and is consistent with rottlerin inhibition of MMT-induced PKC $\delta$  activity in intact PC12 cells (Fig. 9*B*). Our data are also in agreement with previously published values for direct inhibition of PKC $\delta$  by rottlerin *in vitro* kinase assays (Gschwendt et al., 1994; Way et al., 2000; Vancurova et al., 2001).

#### Activation of caspase-3 after intracellular delivery of PKC $\delta$ catalytic fragment

To further confirm the existence of a positive feedback loop between caspase-3 and proteolytic cleavage PKC $\delta$ , we investigated the effect of intracellular delivery of PKC $\delta$  catalytic fragment on caspase-3 activity in PC12 cells. We used a recently developed lipid-mediated delivery system to introduce the catalytically active PKC $\delta$  fragment into the cytoplasm of PC12 cells (Zelphati et al., 2001). The cells were treated with the delivery reagent with or without the PKC $\delta$  catalytic fragment for 4 hr at 37°C. As shown in Table 1, PC12 cells delivered with PKC $\delta$  catalytic fragment showed increases in caspase-3 activity to 341% of reagent control. Neither the intracellular delivery of heat-inactivated PKC $\delta$  catalytic fragment nor the delivery reagent alone produced any increases in caspase-3 enzymatic activity. These results strongly suggest that catalytic fragment of PKC $\delta$  is capable of mediating caspase-3 activation, further supporting our hypothesis that proteolytic cleavage of PKC $\delta$  can augment caspase-3 activity by a positive feedback loop during MMT treatment.





**Figure 9.** Rottlerin inhibits PKC $\delta$  kinase activity in intact cells and *in vitro*. *A*, Dose-dependent increase in PKC $\delta$  activity. *B*, Rottlerin suppresses MMT-induced increase in PKC $\delta$  kinase activity in intact cells. *C*, Rottlerin inhibits PKC $\delta$  kinase activity *in vitro*. Subconfluent undifferentiated PC12 cells were treated with 200  $\mu$ M MMT for 1 hr at 37°C with or without the inclusion of rottlerin (Rot; 5–20  $\mu$ M). Rottlerin was added 30 min before the addition of MMT. For *in vitro* inhibition of PKC $\delta$  activity, rottlerin (5–20  $\mu$ M) was added to the immunoprecipitated samples from MMT-treated cells and incubated for 30 min before the addition of substrate (histone H1) and [ $\gamma$ - $^{32}$ P]ATP. The immunoprecipitation kinase assay was performed as described in Materials and Methods. The bands were quantified by a PhosphoImager after scanning the dried gel and expressed as a percentage of control (untreated cells) (*A*), percentage of MMT treatment (*B*), or percentage of PKC $\delta$  kinase activity (*C*). The data represent an average of three individual measurements from two separate experiments  $\pm$  SEM. Asterisks (\* $p$  < 0.05) indicate significant differences compared with control, MMT-treated cells, or PKC $\delta$  kinase activity.

**Table 1. Activation of caspase-3 after intracellular delivery of PKC $\delta$  catalytic fragment in PC12 cells**

Treatment	Caspase-3 activity (fluorescence units/mg protein/hr)	% Reagent control
Reagent control	25,293 $\pm$ 5726	100
Catalytic active PKC $\delta$ fragment	86,296 $\pm$ 34,116	341 $\pm$ 135*
Heat inactivated PKC $\delta$ catalytic fragment	29,814 $\pm$ 6851	118 $\pm$ 27

\*Asterisk indicates significant difference ( $p$  < 0.05) compared with reagent control. The data are given as the mean  $\pm$  SEM from two separate experiments performed in triplicate.

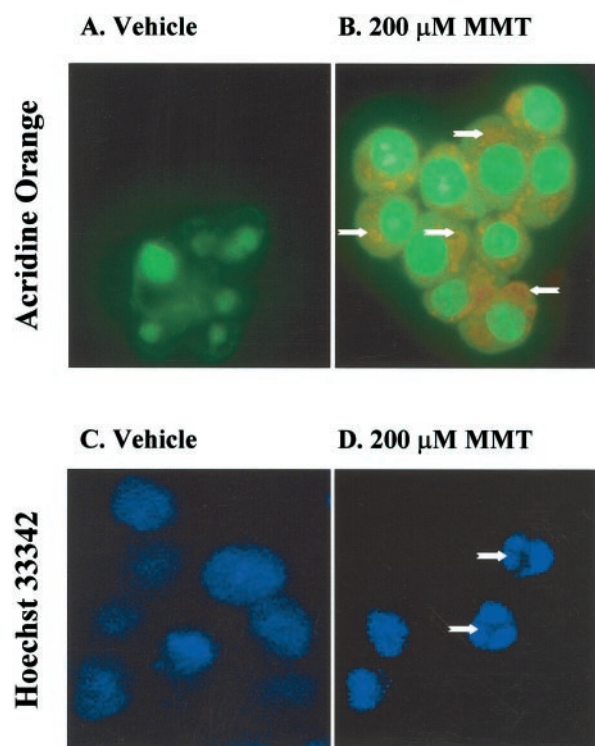
### *In situ* fluorometric detection of apoptosis

To understand the functional consequence of the activation of many apoptotic factors, we tested whether MMT induces DNA fragmentation. Chromosomal breakdown of DNA into 200 bp nucleosomal fragments and DNA condensation are hallmarks of cells undergoing apoptosis. We used *in situ* fluorometric analysis to identify apoptotic cells using acridine orange and Hoechst 33342 to detect nuclear condensation and DNA damage after MMT treatment. In these experiments, we found the majority of PC12 cells exposed to 200  $\mu$ M MMT for 1 hr showed enhanced red fluorescence and reduced green fluorescence, suggesting that acridine orange dye is bound to single-stranded or highly condensed DNA (Fig. 10*B*), whereas little or no enhanced red fluorescence was seen in vehicle-treated cells (Fig. 10*A*). Similarly, the nucleus of PC12 cells exposed to 200  $\mu$ M MMT for 1 hr and subsequently stained with Hoechst 33342 dye showed nuclear condensation as the dye bound to the highly condensed DNA (Fig. 10*D*). The Hoechst 33342 staining of vehicle-treated cells showed a weak and diffused staining, indicating that there is no nuclear condensation in these cells (Fig. 10*C*).

### Oxidative stress, caspase-3, and PKC $\delta$ mediate MMT-induced DNA fragmentation

To further confirm the results obtained by *in situ* fluorometric detection of live apoptosis and to assess the involvement of caspases and PKC $\delta$  in mediating apoptosis, a quantitative DNA fragmentation assay was performed. PC12 cells treated with 200  $\mu$ M MMT showed DNA fragmentation within 1 hr of exposure (Fig. 11). MMT treatment resulted in more than a twofold increase over the levels of basal (vehicle-treated) DNA fragmentation. Pretreatment with an ROS inhibitor, SOD (100 U/ml), almost completely blocked MMT-induced DNA fragmentation (Fig. 11*A*), indicating that SOD is capable of blocking MMT-induced apoptosis. To further confirm the anti-apoptotic effect of SOD in MMT treatment, a cell-permeable SOD mimetic, MnTBAP, was used. Pretreatment with 10  $\mu$ M MnTBAP also almost completely attenuated MMT-induced apoptosis (Fig. 11*A*). SOD, but not MnTBAP, when treated alone also significantly attenuated the basal apoptosis in vehicle-treated PC12 cells.

Pretreatment for 30 min with PKC $\delta$  inhibitor rottlerin (10  $\mu$ M) completely prevented 200  $\mu$ M MMT-induced DNA fragmentation (Fig. 11*B*). Similarly, pretreatment with caspase inhibitors Z-DEVD-FMK (50  $\mu$ M) or Z-VAD-FMK (100  $\mu$ M) almost completely blocked MMT-induced DNA fragmentation (Fig. 11*B*). Rottlerin, Z-DEVD-FMK, and Z-VAD-FMK when treated alone did not significantly attenuate the basal apoptosis in

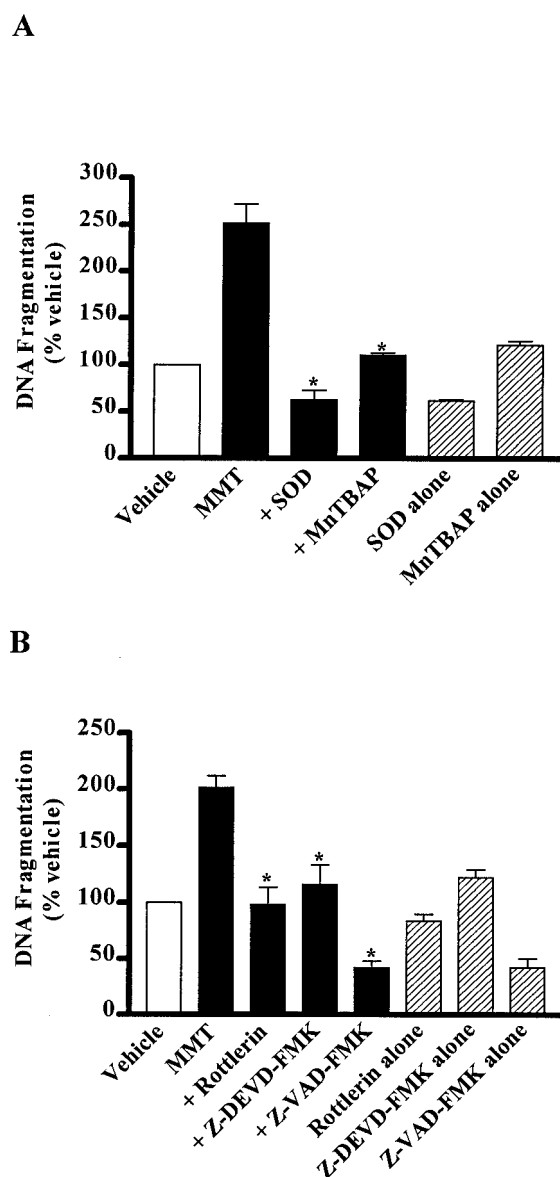


**Figure 10.** MMT treatment increases apoptosis *in situ*. *A, C*, Vehicle-treated cells; *B, D*, 200  $\mu$ M MMT-treated cells. PC12 cells were grown on laminin-coated slides for 2–3 d and then exposed to 200  $\mu$ M MMT for 1 hr. *A, B*, For acridine orange staining, cells were treated with acridine orange (10  $\mu$ M) for 15 min in the dark at RT after exposure to MMT. Arrows indicate enhanced red fluorescence and reduced green fluorescence in MMT-treated cells, which are undergoing apoptosis, whereas little or no enhanced red fluorescence was seen in vehicle-treated cells. *C, D*, For Hoechst 33342 staining, cells were stained with Hoechst 33342 (10  $\mu$ g/ml) for 3 min in dark after exposure to MMT. Arrows indicate apoptotic cells containing condensed chromatin.

vehicle-treated PC12 cells, suggesting that caspase-3 and PKC $\delta$  are participants specifically in MMT-stimulated, and not basal programmed cell death.

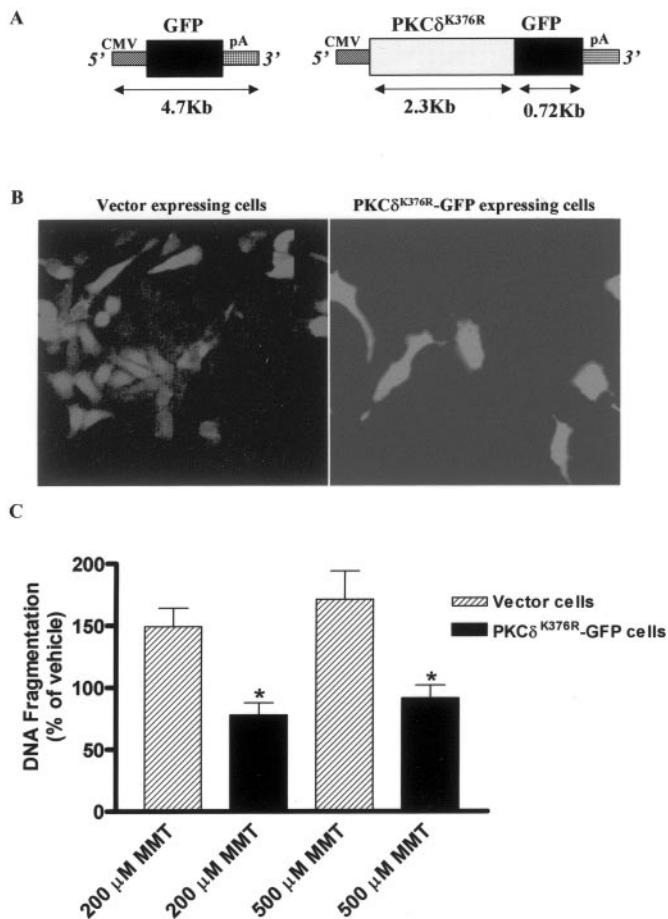
#### MMT treatment does not induce apoptosis in mesencephalic cells overexpressing mutant PKC $\delta^{K376R}$ protein

Pretreatment with a PKC $\delta$ -specific inhibitor rottlerin significantly reduced MMT-induced DNA fragmentation, supporting the idea that the catalytic activity of PKC $\delta$  enzyme is vital for induction of apoptosis. If the kinase activity of PKC $\delta$  is essential for apoptosis, then overexpression of a kinase inactive PKC $\delta$  mutant protein should suppress MMT-induced DNA fragmentation, which occurs downstream of caspase-3 dependent PKC $\delta$  activation. Alternatively, overexpression of a kinase inactive PKC $\delta$  mutant protein may not interfere with ROS production, an event that occurs before caspase-3 dependent PKC $\delta$  activation. To explore these possibilities, we engineered a rat-immortalized mesencephalic (1RB<sub>3</sub>AN<sub>27</sub>) cell line to express a dominant-negative PKC $\delta$  mutant by stably transfecting with plasmids pPKC $\delta^{K376R}$ -GFP (in which a lysine at 376 position is mutated to arginine) and pEGFP-N1 (Fig. 12*A*). The plasmid pPKC $\delta^{K376R}$ -GFP codes for a catalytically inactive PKC $\delta$  mutant fused to GFP and pEGFP-N1 plasmid encodes the green fluorescent protein alone, which was used as a vector control. Figure 12*B* shows stable GFP expression in cell lines transfected with kinase inactive mutant



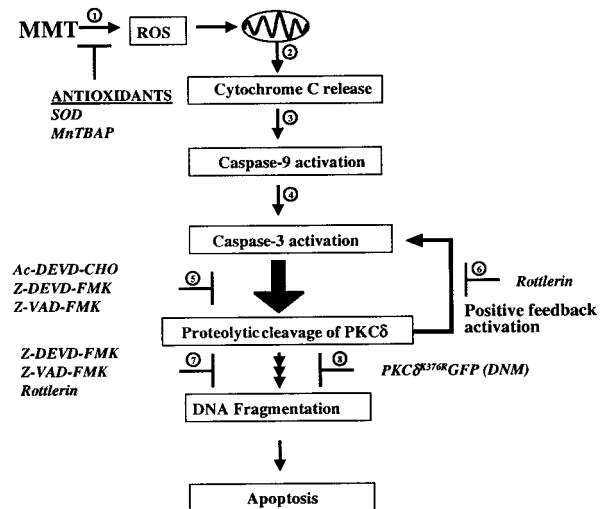
**Figure 11.** Suppression of MMT-induced apoptosis in PC12 cells. *A*, ROS inhibitors, SOD and MnTBAP. *B*, Caspase-3 inhibitors, Z-VAD-FMK and Z-DEVD-FMK and PKC $\delta$  inhibitor, rottlerin. Subconfluent cultures of undifferentiated PC12 cells were treated with MMT (200  $\mu$ M) with or without the inclusion of the following inhibitors: ROS inhibitors SOD (100 U/ml) or MnTBAP (10  $\mu$ M); caspase inhibitors Z-VAD-FMK (100  $\mu$ M) or Z-DEVD-FMK (50  $\mu$ M); and PKC $\delta$  inhibitor rottlerin (10  $\mu$ M). Inhibitors were added 30 min before addition of MMT. Cells were harvested 1 hr after MMT treatment. Apoptosis was assayed using ELISA assay as described in Materials and Methods. The data are expressed as percentage of apoptosis observed in vehicle-treated cells. The data represent the mean  $\pm$  SEM of six individual measurements from three separate experiments. Asterisks (\* $p$  < 0.01) indicate significant differences when compared with cells exposed to 200  $\mu$ M MMT.

PKC $\delta^{K376R}$ -GFP and GFP alone. Antibody directed against GFP detected  $\sim$ 100 and 27 kDa bands in cell lines expressing kinase inactive mutant PKC $\delta^{K376R}$ -GFP and GFP alone, respectively. Similarly, antibody directed against PKC $\delta$  detected  $\sim$ 100 and 72 kDa bands in cell line expressing PKC $\delta^{K376R}$ -GFP fusion, whereas only a 72 kDa band was detected in cells expressing GFP alone. The 100, 72, and 27 kDa bands obtained in Western blots correspond to the expression of intact mutant PKC $\delta^{K376R}$ -GFP



**Figure 12.** Overexpression of catalytically inactive PKC $\delta$  protein blocks MMT-induced apoptosis in immortalized dopaminergic neuronal cell line (1RB<sub>3</sub>AN<sub>27</sub>). *A*, Plasmid description, pEGFP-N1 construct codes for the green fluorescent protein (GFP) mRNA transcribed under the 5' human cytomegalovirus (CMV) immediate early promoter, and the mRNA is stabilized with the 3' SV40 mRNA polyadenylation signal (pA) and was used as vector control. PKC $\delta^{K376R}$ -GFP construct codes for the kinase inactive PKC $\delta$ -GFP fusion transcript. *B*, Stable expression of GFP and PKC $\delta^{K376R}$ -GFP fusion protein in 1RB<sub>3</sub>AN<sub>27</sub> cells. The cells were viewed under a fluorescence microscope, and images were obtained with a SPOT digital camera. *C*, Subconfluent cultures of undifferentiated 1RB<sub>3</sub>AN<sub>27</sub> cells stably expressing vector or PKC $\delta^{K376R}$ -GFP fusion protein were treated with MMT (200 and 500  $\mu$ M) for 3 hr. Apoptosis was assayed using ELISA assay as described in Materials and Methods. The data are expressed as percentage of apoptosis observed in vehicle-treated cells. The data represent a mean  $\pm$  SEM of four to six individual measurements from two separate experiments. Asterisks (\* $p$  < 0.01) indicate significant differences when compared with MMT-treated cells.

fusion protein, native PKC $\delta$  and GFP protein, respectively. In apoptotic measurements, MMT-induced DNA fragmentation was completely abolished in cells stably expressing kinase inactive PKC $\delta$  protein but not in GFP-alone (vector) transfected 1RB<sub>3</sub>AN<sub>27</sub> cells (Fig. 12C). However, MMT-induced ROS production was not significantly different between the kinase inactive PKC $\delta$ -GFP and GFP-alone expressing cell lines. A 15 min exposure of 1RB<sub>3</sub>AN<sub>27</sub> cells stably expressing kinase inactive PKC $\delta$ -GFP and GFP alone to 200  $\mu$ M MMT resulted in a  $157 \pm 20\%$  and  $148 \pm 11\%$  increase in ROS production, respectively. These results suggest that the kinase activity of PKC $\delta$  is essential for MMT-induced DNA fragmentation. These data also indicate that the suppression of apoptosis in PKC $\delta$  dominant-negative cells



**Figure 13.** A model describing the sequence of cell death signaling events in MMT-induced apoptosis. 1, Increased ROS production can be blocked by pretreatment with antioxidants, superoxide dismutase and MnTBAP; 2, cytochrome C is released into the cytosol from the mitochondria; 3, cytosolic cytochrome C activates caspase-9; 4, caspase-9 activates caspase-3; 5, caspase-3 mediates proteolytic cleavage of PKC $\delta$ , which can be blocked by pretreatment with the caspase inhibitors Ac-DEVD-CHO, Z-DEVD-FMK, and Z-VAD-FMK; 6, pretreatment with rottlerin, a PKC $\delta$  inhibitor, reduces caspase-3 activity indicating a possible feedback activation; 7, both caspase-3 and PKC $\delta$  inhibitors block MMT-induced DNA fragmentation; and 8, dopaminergic cells stably overexpressing catalytically inactive PKC $\delta$  [dominant-negative mutant (DNM) PKC $\delta^{K376R}$ GFP] completely blocked MMT-induced DNA fragmentation. In conclusion, our data suggest that caspase-3-dependent proteolytic activation of PKC $\delta$  plays a key role in MMT-induced dopaminergic cell death.

was not caused by a change in the amount of ROS generated in the PKC $\delta$ -GFP overexpressing cells versus GFP-vector cells.

## DISCUSSION

We recently reported that exposure of PC12 cells to MMT induces dopamine depletion and cytotoxic cell death in a dose- and time-dependent manner (Wagner et al., 2000). The present study extends these observations by demonstrating that MMT induces apoptosis in dopamine-producing cells through ROS production and activation of a series of specific cell death signaling events, including release of cytochrome C into the cytosol, activation of caspase-9 and caspase-3, proteolytic cleavage of PKC $\delta$ , and nuclear DNA breakdown. To our knowledge, this is the first report demonstrating that caspase-3-dependent proteolytic cleavage of PKC $\delta$  mediates oxidative stress-induced apoptotic cell death in dopaminergic cells after exposure to an environmental neurotoxicant.

In this study, MMT treatment elevated intracellular ROS levels over 45 min in a time- and dose-dependent manner. ROS generation was observed as early as 5 min after MMT exposure, indicating that ROS generation precedes the cytotoxic response. ROS has been shown to induce cytochrome C release from mitochondria in both neuronal and non-neuronal systems by activation of mitochondrial transition pore opening, which results in swelling and rupturing of mitochondrial membrane (Liu et al., 1996; Petit et al., 1996; Blackstone and Green, 1999; Hollensworth et al., 2000; Lee and Wei, 2000). We observed an accumulation of cytosolic cytochrome C in PC12 cells within 15 min after MMT treatment, suggesting that ROS may be an initial signal for

the release of cytochrome C. Our data are also consistent with the actions of other dopaminergic toxins, 1-methyl-4-phenylpyridinium (MPP<sup>+</sup>) (Leist et al., 1998; Cassarino et al., 1999) and 6-hydroxydopamine (6-OHDA) (Dodel et al., 1999) in their ability to induce ROS-mediated cytochrome C release. Cytochrome C, once released into the cytoplasm, forms a complex with apoptotic protease activating factor, and together they activate a series of caspases.

Activation of caspases by cytosolic cytochrome C is an early and essential step in the apoptotic-signaling pathway (Earnshaw et al., 1999; Jellinger, 2000). Several lines of evidence indicate that caspase-3 plays a major role in the regulation and execution phase of both *in vitro* and *in vivo* models of apoptosis (Cohen, 1997; Schultz and Andreasen, 1999). In this study, we demonstrate that MMT exposure to PC12 cells results in a dramatic activation of caspase-3, indicating that caspase-3 may play a key role in MMT-induced dopaminergic degeneration. Our data are further supported by a recent study in which caspase-3 activation was observed in neuronal cultures after MPP<sup>+</sup> and 6-OHDA treatment (Dodel et al., 1999). The importance of caspase-3 activation as an indicator of apoptosis is further underscored by a recent study from Hartmann et al. (2000), who demonstrated caspase-3 to be a vulnerability factor and a critical effector of apoptotic death in dopaminergic neurons in both MPTP mouse model and in human patients with Parkinson's disease.

Biochemical consequences of caspase-3 activation are proteolytic cleavage of cellular targets associated with apoptosis. Poly (ADP-ribose) polymerase, a DNA cleaving enzyme, has been established as one of the important apoptotic substrates of caspase-3 (Earnshaw et al., 1999; Schultz and Andreasen, 1999). In the present study, we demonstrated that PKC $\delta$  is an emerging putative endogenous substrate for caspase-3 and show that MMT exposure induces PKC $\delta$  cleavage and increases PKC $\delta$  activity in a dose- and time-dependent manner in dopaminergic cells. Additionally, MMT does not induce cleavage of PKC $\alpha$ , suggesting that the cleavage is isoform specific. Proteolytic cleavage of PKC $\delta$  in MMT-treated PC12 cells is blocked by specific caspase inhibitors, indicating that the cleavage is mediated by caspase-3. Proteolytic cleavage of PKC $\delta$  by caspase-3 results in persistent activation of PKC $\delta$  in cytosol, which might initiate a myriad of vital signaling cascades. In a previous study, proteolytic cleavage of PKC $\delta$  was observed in KCl-deprived cerebellar granule cell apoptosis, however, this study did not characterize the caspase-3 dependency of PKC $\delta$  cleavage (Villalba, 1998). Recent studies have additionally implicated the persistently active catalytic fragment of PKC $\delta$  in apoptotic cell death in non-neuronal systems (Earnshaw et al., 1999; Schultz and Andreasen, 1999).

We further determined a possible interaction between PKC $\delta$  and caspase-3 activation in MMT-treated PC12 cells using the PKC $\delta$ -specific inhibitor rottlerin. Pre- and post-rottlerin treatment effectively blocked MMT-induced caspase-3 activation in PC12 cells in a dose-dependent manner, suggesting a positive feedback modulatory role of PKC $\delta$  on caspase-3 activity. Although rottlerin suppressed MMT-induced caspase-3 activity in both pre- and post-treatments (Fig. 7), the inhibition was more pronounced in post-treatment at higher concentrations. The reason for the pronounced inhibition is not completely clear at the present time, and we attribute that this might be caused by action of rottlerin on other cellular targets including other kinases (Davies et al., 2000; Way et al., 2000). However, there was no significant difference in caspase activity between pre- and post-treatments of rottlerin at lower dose 5  $\mu$ M, the concentration at

which a pronounced inhibition of PKC $\delta$  activity *in vitro* (Fig. 9C) was observed. Furthermore, delivery of the catalytically active PKC $\delta$  fragment alone into PC12 cells increased the caspase-3 activity, confirming the presence of such a positive feedback mechanism. It appears that maximal caspase-3 activity requires the kinase activity of cleaved PKC $\delta$  fragment and is made possible by the existence of a positive feedback activation loop. A positive feedback loop between PKC $\delta$  and caspase-3 activation has recently been shown to exist in an etoposide-induced salivary cell apoptosis model (Reyland et al., 1999). Thus, the existence of such a positive feedback loop discovered independently by two research groups in two different apoptotic models, MMT-induced PC12 apoptosis (this study) and in etoposide-induced salivary cell apoptosis, suggests that this may be an important regulatory mechanism, allowing for the amplification of apoptotic signaling processes. Further studies are needed to understand the cellular mechanisms of caspase-3 regulation by PKC $\delta$  and their role in neuronal apoptosis.

DNA fragmentation and condensation resulting from intranucleosomal cleavage have long been considered biochemical hallmarks of apoptosis and are terminal events in the apoptotic process (Cohen, 1997). *In situ* fluorometric experiments using two different fluorescent dyes revealed that MMT exposure of PC12 cells induces chromatin condensation in the nucleus. We also took advantage of a recently developed ELISA method that provides a better quantitative measurement of DNA fragmentation. MMT exposure to PC12 cells induced DNA fragmentation, which could be suppressed under conditions where caspase or PKC $\delta$  activities were inhibited. Suppression of MMT-induced DNA fragmentation by either caspase inhibitors or rottlerin in the present study indicates that both caspase-3 and PKC $\delta$  activities are essential for MMT-induced DNA fragmentation. Furthermore, caspase-3-mediated promotion of DNA fragmentation may be amplified via feedback activation of caspase-3 by the catalytically active PKC $\delta$  fragment. To additionally confirm the role of PKC $\delta$  in MMT-induced apoptosis, we conducted dominant-negative experiments by stably expressing catalytically inactive PKC $\delta$  protein (PKC $\delta$ <sup>K376R</sup>) in immortalized rat mesencephalic neurons. MMT treatment produced a significant increase in DNA fragmentation in vector control cells, whereas MMT failed to induce DNA fragmentation in catalytically inactive PKC $\delta$  over-expressing cells, thus confirming the key functional role of PKC $\delta$  in MMT-induced apoptotic cell death.

Although the events downstream of PKC $\delta$  and those that lead to apoptosis remain unclear, recent studies from many research groups have shown that catalytically active PKC $\delta$  fragment can regulate the activity of a host of cell signaling molecules such as scramblase, a membrane phosphatidylserine translocator (Frasch et al., 2000), DNA protein kinase, a DNA repair enzyme (Bharti et al., 1998), heat-shock proteins-25/27 (Maizels et al., 1998), histone H2B (Ajiro, 2000), and lamin kinase (Cross et al., 2000). In addition, PKC $\delta$  has been shown to phosphorylate other signaling molecules such as MAP kinases (Chen et al., 1999), Jak2, a tyrosine kinase (Kovanen et al., 2000), and Stat3, signal transducers and activators of transcription (Jain et al., 1999). Most recently, it has been demonstrated that PKC $\delta$  activates redox-sensitive transcription factor, NF- $\kappa$ B, and thereby promotes apoptosis in neutrophils (Vancurova et al., 2001). Furthermore, PKC $\delta$  has been shown to translocate to cytosol and a variety of cellular organelles to initiate apoptosis (Sawai et al., 1997; Chen et al., 1999; Dal Pra et al., 1999; Li et al., 1999; Dempsey et al., 2000; Majumder et al., 2000). Hence, constitu-

tively active PKC $\delta$  fragment can promote loss of cellular regulatory function in many of its substrates, resulting in rapid apoptosis. Currently, our laboratory is focusing on identifying critical cellular targets of PKC $\delta$  that might contribute to apoptotic cell death in dopaminergic cells.

In conclusion, we demonstrate for the first time that an environmental neurotoxicant, MMT, induces dopaminergic degeneration by a novel oxidative stress-mediated apoptotic mechanism in which caspase-3-dependent proteolytic cleavage of PKC $\delta$  plays a critical role (Fig. 13). Our data also demonstrate a positive feedback amplification loop between PKC $\delta$  and caspase-3, which has a regulatory role in the promotion of apoptosis. Further research into identifying molecules that participate in this loop might provide very exciting information regarding cell signaling and neuronal apoptosis. Finally, this study emphasizes the importance of characterizing oxidative stress-induced cell signaling molecules after neurotoxicant exposure to better understand the role of environmental risk factors in the pathogenesis of Parkinson's disease.

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