

1-Methyl-4-Phenylpyridinium Induces Autocrine Excitotoxicity, Protease Activation, and Neuronal Apoptosis

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ABSTRACT

The pathogenesis of several neurodegenerative diseases may involve indirect excitotoxic mechanisms, where glutamate receptor overstimulation is a secondary consequence of initial functional defects of neurons (e.g., impairment of mitochondrial energy generation). The neurotoxin 1-methyl-4-phenylpyridinium (MPP⁺) and other mitochondrial inhibitors (e.g., rotenone or 3-nitropropionic acid) elicited apoptosis in cerebellar granule cell cultures via stimulation of autocrine excitotoxicity. Cell death, increase in intracellular Ca²⁺ concentration, release of cytochrome c, and all biochemical and morphological signs of apoptosis were prevented by blockade of the *N*-methyl-D-aspartate receptor with noncompetitive, glycine-site or glutamate-site inhibitors. In addition, MPP⁺-induced apoptosis was

reduced by high Mg²⁺ concentrations in the medium or by inhibiting exocytosis with clostridial neurotoxins. Two classes of cysteine proteases were involved in the execution of cell death: caspases and calpains. Inhibitors of either class of proteases prevented cell death, cleavage of intracellular proteins (i.e., fodrin), and the appearance of typical features of apoptosis such as phosphatidylserine translocation or DNA fragmentation. However, protease inhibitors did not interfere with the initial intracellular Ca²⁺ concentration increase. We suggest that MPP⁺ as well as other mitochondrial inhibitors trigger indirect excitotoxic processes, which lead to Ca²⁺ overload, protease activation, and subsequent neuronal apoptosis.

Excitotoxic mechanisms (i.e., excessive stimulation of glutamate receptors with a resultant disturbance of cellular [Ca²⁺]_i homeostasis) may be involved in stroke and possibly in slowly developing neurodegenerative diseases (Choi and Rothman, 1990). However, glutamate receptor stimulation is rarely a primary event in neurotoxicity. Glutamate release and glutamate-triggered excitotoxicity are rather secondary consequences of other defects or metabolic disturbances. A frequent initiating condition is energy depletion, and mitochondrial dysfunction is among the most generalized causes favoring the development of different neurodegenerative diseases (Beal, 1996). For example, Huntington's disease is modeled by exposing specific neuronal subpopulations to mitochondrial toxins (Ferrante *et al.*, 1997), and ischemic damage can be examined *in vitro* after mitochondrial substrate

depletion due to oxygen/glucose deprivation (Choi and Rothman, 1990).

In these models, a close relationship has been established between energy deficiency and excitotoxicity. The ATP loss resulting from decreased mitochondrial energy generation would lead to impaired function of ion pumps and partial hypopolarization of neurons, thereby releasing the voltage-dependent Mg²⁺ block of the NMDA-R-gated Ca²⁺ channel. This would make the receptor/channel hypersensitive to glutamate stimulation. NMDA-R-mediated influx of Na⁺ and Ca²⁺ then would increase energy demand and ATP depletion, enhance depolarization, trigger further [Ca²⁺]_i increase, and eventually result in further glutamate release. This putatively self-propagating process finally leading to a loss of [Ca²⁺]_i homeostasis and excitotoxicity has been termed the "energy-linked excitotoxicity hypothesis" (Henneberry *et al.*, 1989; Zeevalk and Nicklas, 1990). Distal mechanisms (downstream of [Ca²⁺]_i increase) leading to neuronal death under such conditions may differ from those operating

This study was supported by the DFG Grants Ni519/1-1 and Ni519/2-1 and the EEC Grants ENV4-CT96-0169, BMH4CT97-2410, and 12029-97-06 FIED ISP D.

ABBREVIATIONS: [Ca²⁺]_i, intracellular Ca²⁺ concentration; ΔΨ, mitochondrial membrane potential; AP5, 5-aminophosphovalerat; calp II, acetyl-leucyl-leucyl-L-methionyl; calp III, z-Val-L-phenylalaninal; CCCP, carbonylcyanide-chlorophenylhydrazone; CGC, cerebellar granule cells; CSS, controlled salt solution; DCK, 5,7-dichlorokynurenate; DIV, days *in vitro*; EGTA, ethylene glycol bis(β-aminoethyl ether)-*N,N,N',N'*-tetraacetic acid; HEPES, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid; DNQX, dinitroquinoxalinedione; MK801, (+)-5-methyl-10,11-dihydro-5*H*-dibenzo[*a,c*]cyclohepten-5,10-imine; MPP⁺, 1-methyl-4-phenylpyridinium; MTT, 3-(4,5-dimethylthiazole-2-yl)-2,5-diphenyltetrasodium bromide; NMDA, *N*-methyl-D-aspartate; NMDA-R, *N*-methyl-D-aspartate receptor; NO, nitric oxide; 3-NP, 3-nitropropionic acid; PARP, poly-(ADP-ribose)-polymerase (E.C. 2.4.2.30); BoNT/C, botulinum neurotoxin serotype C; PS, phosphatidylserine; TMRE, tetramethylrhodamine ethylester; VDCC, voltage-dependent calcium channel; z-D-cbk, z-aspartyl-2,6-dichlorobenzoyloxymethylketone.

after simple glutamate receptor stimulation and are largely unknown.

Excitotoxicity may lead to either necrosis or to a more ordered sequence of molecular events resulting in apoptosis (Ankarcrona *et al.*, 1995; Leist and Nicotera, 1998). Recent work in our laboratory has shown caspase activation downstream to the initial $[Ca^{2+}]_i$ increase in a model of indirect excitotoxicity, where NMDA-R activation was triggered by NO (Leist *et al.*, 1997a, 1997d). In parallel, studies *in vivo* have suggested that caspases may be involved in neuronal damage after stroke (Loddick *et al.*, 1996). In all these models, the exact primary insult is little characterized and possibly not limited to mitochondrial dysfunction; therefore, it is still unclear how a primary mitochondrial impairment can lead to excitotoxic apoptosis.

The neurotoxin MPTP induces a Parkinson's disease-like syndrome in humans and animals via its active metabolite, MPP⁺ (Tipton and Singer, 1993). Although MPP⁺ affects only certain neurons *in vivo* due to pharmacokinetic reason, the substance inhibits mitochondrial function in any cultured neuronal or non-neuronal cell or isolated mitochondria. A known molecular target of MPP⁺ is the mitochondrial respiratory chain complex I (NADH-ubiquinone-oxidoreductase) (Nicklas *et al.*, 1985; Kilbourn *et al.*, 1997), and no other main target has been characterized to date, despite extensive studies. *In vivo* (i.e., in the presence of glutamatergic cortical inputs), the primary impairment of mitochondrial respiration by MPP⁺ sensitizes neurons to secondary excitotoxic damage. Thus, blockade of glutamate receptors (Turski *et al.*, 1991; Srivastava *et al.*, 1993) or prevention of the synthesis of endogenous NO (Schulz *et al.*, 1995; Hantraye *et al.*, 1996), a mediator closely linked to excitotoxicity, has been found to reduce MPP⁺ toxicity in animals.

In the current study, we exposed neurons to the mitochondrial poison MPP⁺ or other respiratory chain inhibitors. CGC have been shown to be susceptible to MPP⁺ (Marini *et al.*, 1989). Thus, we used here neuronal cultures, which mimic the very high density of excitable glutamatergic synapses present *in vivo*. Densely plated CGC, differentiated for ≥ 8 days in K⁺-supplemented medium, represent an ideal culture system for this purpose. We examined in this culture system (i) whether excitotoxic mechanisms contribute to MPP⁺ toxicity, (ii) whether such excitotoxicity results in apoptosis, (iii) under which conditions the execution of excitotoxic death triggered by MPP⁺ or other mitochondrial toxins involves caspases, and (iv) which apoptotic features depend on the activity of protease families activated by excitotoxic disruption of $[Ca^{2+}]_i$ homeostasis.

Experimental Procedures

Materials. Fluo 3-acetoxymethyl ester, Fura 2-acetoxymethyl ester, calcein acetoxymethyl ester, TMRE, SYTOX, ethidium homodimer-1, and H-33342 were obtained from Molecular Probes (Eugene, OR). The caspase substrate DEVD-afc, MPP⁺, DCK, DNQX, and 6,7-dichloroquinoxaline-2,3-dione were obtained from BIOMOL (Hamburg, Germany). MK801 came from RBI (Biotrend Chemikalien GmbH, Köln, Germany). succ-LLVY-amc; calpain inhibitors I, II, and III (Ac-Leu-Leu-L-norleucinal, Ac-Leu-Leu-L-methional, and calp III, z-Phe-chloromethylketone (cmk); and the caspase-inhibitors DEVD-CHO, z-VAD-fluoromethylketone (fmk), Ac-YVAD-cmk, or Ac-YVAD-2,6-dimethylbenzoyloxymethylketone (ICE II) and z-D-2,6-dichlorobenzoyloxymethylketone (cbk) (ICE III) were obtained

from Bachem Biochemica (Heidelberg, Germany). DEVD-fmk was from Enzyme Systems (Dublin, CA). Fluorescein-labeled annexin V (annexin V) was from Boehringer-Mannheim (Mannheim, Germany). Clostridial toxins were generously supplied by Dr. C. Montecucco (University of Padova, Padova, Italy). Solvents and inorganic salts were from Merck (Darmstadt, Germany) or Riedel-de Haen (Seelze, Germany). All other reagents not further specified were from Sigma (Deisenhofen, Germany).

Animals. PARP^{-/-} mice (C57Bl/6 \times 129/Sv background) or corresponding wild-type animals were generously provided by Dr. Zhao-Qi Wang (IARC, Lyon, France) (Wang *et al.*, 1995). All animals used for cell preparations were typed by Southern blotting (Wang *et al.*, 1995) to verify the genotype. For other experiments, 8-day-old specific pathogen-free BALB/c mice were obtained from the Animal Unit of the University of Konstanz. All experiments were performed in accordance with international guidelines to minimize pain and discomfort (National Institutes of Health guidelines and European Community Council Directive 86/609/EEC).

Cell culture. Murine CGC were prepared as described previously (Schousboe *et al.*, 1989). Neurons were plated onto 100 μ g/ml (250 μ g/ml for glass surfaces) poly-L-lysine (molecular mass, >300 kDa)-coated dishes at a density of $\sim 0.25 \times 10^6$ cells/cm² (800,000 cells/ml; 500 μ l/well; 24-well plate) and cultured in Eagle's basal medium (GIBCO, Paisley, Scotland) supplemented with 10% heat-inactivated fetal calf serum, 20 mM KCl, 2 mM L-glutamine, 100 units/ml penicillin, and 100 μ g/ml streptomycin. Forty-eight hours after plating, cytosine arabinoside (10 μ M) was added to the cultures. Neurons were routinely used at 8–10 DIV unless otherwise indicated. Glial fibrillary acid protein-positive cells were $<5\%$.

Cytotoxicity assays. Cultures were exposed to MPP⁺ or rotenone in their own medium. The culture medium was exchanged for a CSS (120 mM NaCl, 25 mM HEPES, 25 mM KCl, 1.8 mM CaCl₂, 4 mM MgCl₂) plus 15 mM glucose 4 hr after the start of the incubation, and the cells were left in this medium until toxicity parameters were read (usually 18 hr for MTT or nuclear morphology). Inhibitors were added routinely 30 min before exposure to MPP⁺.

To assess plasma membrane integrity and nuclear morphology, CGC were loaded with 0.5 μ M calcein acetoxymethyl ester for 5 min (cells with intact membranes display green fluorescence) in the presence of 1 μ M ethidium homodimer-1 (cells with broken membranes exhibit nuclear red fluorescence) and 500 ng/ml concentration of the bisbenzimidazole dye H-33342 (cell permeant, blue-fluorescent). Alternatively, apoptosis and secondary lysis were quantified by double staining neuronal cultures with 1 μ g/ml H-33342 and 0.5 μ M SYTOX (non-cell-permeant, green-fluorescent chromatin stain). Apoptotic cells were characterized by condensed highly fluorescent nuclei. About 600–1000 cells were counted in nine different fields in two or three different culture wells, and experiments were repeated in at least three different preparations. In addition, the percentage of viable cells was quantified by their MTT-reducing capacity after incubation with 0.5 mg/ml MTT for 60 min. The viability of untreated control cultures was set to 100%, and the viability of treated cultures was expressed as percentage of formazan absorbance compared with that of control cultures.

Ca²⁺ measurements. The $[Ca^{2+}]_i$ was measured by imaging neurons loaded with fluorescent Ca²⁺ indicators. To monitor dynamic changes of Ca²⁺, CGC were loaded in their original medium with 1 μ M fluo-3 acetoxymethyl ester for 10 min at 37°. The medium then was exchanged for CSS, in which CGC were incubated for 5 min to allow complete de-esterification of fluo-3. The medium was exchanged again for the original neuron-conditioned complete Eagle's basal medium supplemented with 20 mM HEPES (MPP⁺ as stimulus) or for CSS (without Mg²⁺) (50 μ M glutamate or 200 μ M NMDA as stimulus) or for CSS plus 2 μ M MK801 (200 μ M kainate as stimulus). CGC were allowed to equilibrate at room temperature for 10 min before the exposure. Images were collected using the 488-nm excitation and 520-nm emission wavelengths and a Leica DM-IRB microscope equipped with a 40 \times NA 1.0 lens. Data from 10–20 neurons

were recorded at 3–60-sec intervals with a Leica TCS 4D confocal system. Relative mean fluorescence levels from defined areas corresponding to the position of neuronal cell bodies were recorded over the time course of the experiment, and the values were arbitrarily set to 1 at the beginning of each experiment. To determine absolute $[Ca^{2+}]_i$, we used the low K_d indicator Fura 2-acetoxymethyl ester (2.5 μ M; Molecular Probes, Eugene, OR), which reports $[Ca^{2+}]_i$ exactly in the low concentration range (≤ 500 nM). This allowed us to examine whether MK801 would maintain $[Ca^{2+}]_i$ to base-line levels in MPP⁺-treated cells. A Leica M-IRB microscope equipped with a computer-controlled filter wheel (Sutter, Novato, CA), quartz optics, and a Dage-72 (Dage-MTI; Michigan City, IN) charge-coupled device (CCD) camera [756 (H) \times 581 (V) pixels] coupled to a videoscope GEN-III image intensifier was used for imaging ($\lambda_{ex-1} = 340$ nm, $\lambda_{ex-2} = 380$ nm, $\lambda_{em} = 505$ nm). Videomicroscopy data were analyzed using software from Imaging Research (St. Catherine's, Ontario, Canada). $[Ca^{2+}]_i$ was determined by *in situ* calibration using the equation $[Ca^{2+}]_i = K_d \times (R - R_{min}) / (R_{max} - R) \times S_{p2} / S_{b2}$, with $K_{d(25^\circ)} = 264$ nM. To determine R_{min} , cells were washed twice with calibration buffer (120 mM NaCl, 25 mM HEPES, 15 mM glucose, 25 mM KCl, 2 mM MgCl₂, 2 mM EGTA) and equilibrated for 20 min in calibration buffer supplemented with 5 μ M ionomycin. Ca²⁺ (5 mM) and ionomycin (10 μ M) were added to saturate Fura-2 with Ca²⁺ and calculate R_{max} . Autofluorescence was measured after the addition of 5 mM MnCl₂.

Electrophoretic assays. Fodrin proteolysis was analyzed by immunoblot. CGC cultures were lysed in RIPA buffer (150 mM NaCl, 50 mM Tris, 1% Nonidet P-40, 0.25% sodium deoxycholate, 1 mM EGTA) supplemented with protease inhibitors (1 mM phenylmethylsulfonyl fluoride, 1 mM benzamide, 1 mM iodoacetate, 1 mM iodoacetamide, 40 μ M leupeptin, 10 μ g/ml antipain, 5 μ g/ml pepstatin). Before lysis, cultures were stained with 0.5 μ M SYTOX to control the percentage of cells with intact membranes, which was >95% for all samples analyzed. Protein was determined using the bicinchoninic acid method (BioRad, München, Germany), and 5 μ g of protein/lane was loaded onto 8% polyacrylamide gels. Proteins were separated under reducing conditions at 60 mA and then blotted onto a nitrocellulose membrane (Amersham-Buchler, Braunschweig, Germany) in a Bio-Rad semidry blotter at 2.6 mA/cm² for 50 min using a Towbin buffer system. Blots were blocked for 1 hr and then incubated with anti-fodrin monoclonal antibody (clone 1622, 1:1000) from Chemicon (Temecula, CA) diluted in TNT (150 mM NaCl, 50 mM Tris, pH 8.0, 0.05% Tween 20) for 1 hr at room temperature. Specifically stained bands were detected by enhanced chemiluminescence (Amersham) using a peroxidase-coupled secondary antibody. For Western blot analysis of procaspase-3 (14% gel, 60 μ g of neuronal protein/lane), we used a primary rabbit anti-human procaspase-3 polyclonal antibody (1:1000, no. 06753; Upstate Biotechnology, Lake Placid, NY) recognizing the 32-kDa murine procaspase but not the cleavage products.

Field inversion gel electrophoresis was performed as described previously (Ankarcrona *et al.*, 1995; Leist *et al.*, 1997d). About 5 \times 10⁶ cells (corresponding to 3 wells of a 12-well plate) were embedded into 40- μ l agarose blocks. λ -DNA concatemers ($n \times 50$ kbp) were used as molecular weight markers.

Mitochondrial function and integrity. ATP was measured luminometrically after lysis of cells in ATP-releasing agent (Sigma) with a commercial kit (Boehringer-Mannheim) as described previously (Leist *et al.*, 1997c). The $\Delta\Psi$ was monitored by loading cells with the fluorescent indicator TMRE (5 nM; λ_{ex} , 568 nm; λ_{em} , ≥ 590 nm). Under these conditions, fluorescence completely disappeared on loss of $\Delta\Psi$ (50 μ M glutamate or 50 μ M CCCP) in neurons that retained plasma membrane integrity. For the experiments, at least three culture dishes were imaged in at least six cell preparations for each data point presented. For semiquantitative analysis of $\Delta\Psi$, we added red fluorescent beads of 2- μ m diameter and calibrated fluorescence intensity (Molecular Probes, Eugene, OR) to the cultures. The standardized fluorescence of the beads was used as an internal standard to normalize the fluorescence intensities of digitized im-

ages of TMRE-loaded neurons. The fluorescence of neurons treated with 20 μ M CCCP was used as reference for depolarized mitochondria.

The release of cytochrome *c* from mitochondria was analyzed by a selective digitonin permeabilization method. This protocol was established to avoid possible artifacts due to potential mechanical breakage of the outer mitochondrial membrane of apoptotic cells by Dounce homogenization. At the indicated time points, the culture medium was exchanged for permeabilization buffer (210 mM D-mannitol, 70 mM sucrose, 10 mM HEPES, 5 mM succinate, 0.2 mM EGTA, 0.15% BSA, 80 μ g/ml digitonin, pH 7.2, 4 $^\circ$). Cell culture plates were shaken gently for 5 min at 4 $^\circ$, and then the permeabilization buffer was removed and centrifuged for 10 min at 13,000 $\times g$. Protein from the supernatants of this centrifugation was precipitated with 5% trichloroacetic acid and separated on a 15% polyacrylamide gel. All samples were obtained from supernatants derived from 10⁶ CGC. Cytochrome *c* was detected by chemiluminescent detection after blotting on nitrocellulose membranes with a monoclonal antibody raised against pigeon cytochrome *c* (clone 7H8.2C12; Pharmingen, San Diego, CA). At the digitonin concentration used in these experiments, neurons released LDH and became permeable to the fluorescent chromatin dye SYTOX or propidium iodide. Control experiments showed that cytochrome *c* was not released into the supernatant of untreated cultures even after 20-min incubation in the permeabilization buffer. Treatment of these cultures with Triton X-100 led to a maximal release of cytochrome *c* into the supernatant. As an additional control, we stained cytochrome *c* *in situ*, using a monoclonal antibody directed against native cytochrome *c* (clone 6H2.B4). In control neurons treated for 10 min with permeabilization buffer and then fixed with paraformaldehyde, punctate cytochrome *c* staining colocalized entirely with a fixable mitochondrial marker (CMX-rosamine, Mito-Tracker Red; Molecular Probes, Eugene, OR), and the stain intensity was identical to that of untreated cells (i.e., without the addition of permeabilization buffer) stained and fixed in the same way.

Enzymatic assays. Caspase-3-like activity (measured by DEVD-afc cleavage) was assayed as described previously (Leist *et al.*, 1997b) with the following modifications: CGC were pelleted in PBS supplemented with 5 mM EDTA, 1 μ g/ml leupeptin, 1 μ g/ml pepstatin, 1 μ g/ml aprotinin, and 1 mM PEFA-block. Lysis was performed in 25 mM HEPES, 5 mM MgCl₂, 1 mM EGTA, 0.5% Triton X-100, 1 μ g/ml leupeptin, 1 μ g/ml pepstatin, 1 μ g/ml aprotinin, and 1 mM PEFA-

TABLE 1
Modulation of mitochondrial function and apoptosis by respiratory chain inhibitors

Treatment ^a	$\Delta\Psi^b$	ATP ^c	Apoptosis ^d
		%	%
MPP ⁺	–	2.0	>80
MPP ⁺ + MK801	++	5.8	<10
3-NP	–	9.1	>80
3-NP + MK801	++	18.5	<10
DG	+	9.6	<10
DG + pyruvate	++	45.0	<10
MPP ⁺ + DG	–	1.3	>80
MPP ⁺ + DG + MK801	–	1.8	<10
3-NP + DG	–	1.4	>80
3-NP + DG + MK801	–	9.9	<10
MPP ⁺ + methylsuccinate	+	3.7	<10
DG (40 mM)	N.D.	3.3	>80
DG (40 mM) + MK801	N.D.	5.6	<10

^a Neurons were exposed to combinations of MPP⁺ (50 μ M), DG (20 mM), 3-NP (400 μ M), methylsuccinate (5 mM), pyruvate (4 mM), or MK801 (2 μ M) for 3 hr in the original culture medium.

^b $\Delta\Psi$ was measured in cells loaded with 5 nM TMRE. Fluorescence intensity corresponding to 70–100% of controls is indicated by ++, 30–70% by +, and <30% by –.

^c ATP is given as percentage of control cells. Data are mean values from three different cell preparations. Standard deviations were always below 3%.

^d Apoptosis was determined by counting the percentage of condensed nuclei after staining with H-33342.

block, pH 7.5. The fluorimetric assay was carried out in microtiter plates with a substrate concentration of 40 μM and a total protein amount of 5 μg . Cleavage of DEVD-afc was followed in reaction buffer (50 mM HEPES, 10 mM dithiothreitol, 1% sucrose, 0.1% 3-[(3-cholamidopropyl)dimethylammonio]propanesulfonate) over a period of 30 min at 37° with $\lambda_{\text{ex}} = 390 \text{ nm}$ and $\lambda_{\text{em}} = 505 \text{ nm}$, and the activity was calibrated with afc-standard solutions. Calpain activity was determined by a kinetic fluorimetric assay as described previously (Leist *et al.*, 1997d). Glucose concentrations in the medium were determined according to the hexokinase/glucose dehydrogenase method using a commercial kit (Sigma).

Visualization of PS translocation. Cells stained with different fluorescent probes were imaged on a Leica DM-IRBE microscope equipped with a computer-controlled z-stage and connected to a TCS-4D UV/VIS confocal scanning system (Leica AG, Benzheim, and Leica Lasertechnik, Heidelberg, Germany). The staining protocol for fluorescein-conjugated annexin V (detection of PS on the outer leaflet of the plasma membrane) was adapted for neuronal cultures as follows: CGC were grown on glass-bottomed culture dishes and incubated with MPP⁺ or rotenone with or without inhibitors. At the end of the incubation periods, 0.5 $\mu\text{g/ml}$ H-33342 was added to the cultures to later visualize chromatin structure. After a 10-min incubation at 37°, CGC were washed for 10 sec with binding buffer (10 mM HEPES, 140 mM NaCl, 2.5 mM CaCl₂, 10 mM MgCl₂) and subsequently incubated for 2 min in the dark with annexin V diluted 1:100 in binding buffer. After a new wash with binding buffer, stained cultures were immersed in binding buffer supplemented with 0.25 μM ethidium homodimer-1 and visualized by three-channel confocal microscopy (blue, chromatin structure, green, annexin binding, red, membrane integrity) using a 63 \times /NA 1.32 UV-corrected lens.

Statistics. Toxicity experiments were run in triplicate and repeated in three to eight cell preparations. Statistical significance was calculated on the original data sets using the Student's *t* test. When variances within the compared groups were not homogeneous, the Welch test was applied. Western blots and measurements of [Ca²⁺]_i were repeated in at least three independent cell preparations.

Results

A role for the NMDA-R in MPP⁺-induced CGC apoptosis. MPP⁺ caused apoptosis of differentiated (8 DIV) CGC in a concentration range of 25–100 μM (Fig. 1). Toxicity of MPP⁺ was evident in cells exposed to MPP⁺ for 2–4 hr, unlike the slow toxicity observed in an earlier report (Marini *et al.*, 1989). Susceptibility of CGC to such rapidly developing

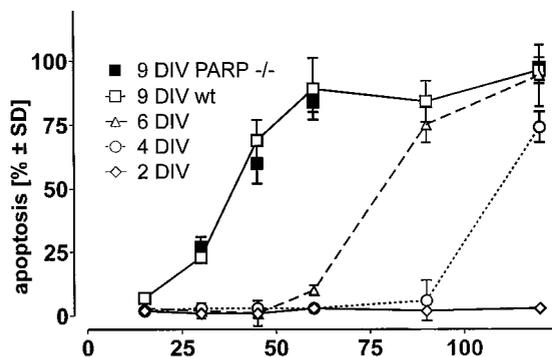


Fig. 1. MPP⁺-induced apoptosis in CGC cultures from wild-type and PARP^{-/-} animals. Cultures from wild-type CGC of different culture age (*wt*) or PARP-deficient mice (PARP^{-/-}) were challenged with MPP⁺ concentrations as indicated. The percentage of neurons with apoptotic nuclei was determined after 7 hr by staining with fluorescent chromatin dyes and counting of the cells. Data are mean \pm standard deviation of triplicate determinations.

MPP⁺-induced apoptosis was strongly dependent on the cell differentiation state. CGC were hardly sensitive toward MPP⁺ during the first 2 DIV, whereas they became progressively more susceptible to induction of apoptosis elicited by MPP⁺ with increasing time in culture, as described elsewhere for other excitotoxic stimuli (i.e., NMDA, NO, or ONOO⁻) (Leist *et al.*, 1997a) (Fig. 1).

The rapid toxicity of MPP⁺ (4-hr exposure) was entirely prevented by the noncompetitive NMDA-R blocker MK801 over the full concentration range used in this study (MPP⁺ \leq 120 μM) (Fig. 2A). In the presence of MK801, MPP⁺ also caused no significant cell death when neurons were continuously exposed for 18 hr. Long preincubation with MK801 was not required because the channel blocker still protected MPP⁺-challenged neurons when it was added up to 15 min after the exposure to MPP⁺. Neurons challenged for 4 hr with 50 μM MPP⁺ in the presence of MK801 and then cultured in conditioned medium without MPP⁺ for an additional 96 hr were virtually unaffected (\geq 95% viable). In contrast, CGC exposed permanently to MPP⁺ at concentrations of \geq 50 μM , in the presence of MK801, showed signs of degeneration after 24 hr and died after 36–72 hr. This suggests that chronic exposure of CGC to MPP⁺ resulted in the activation of additional, nonexcitotoxicity mechanisms.

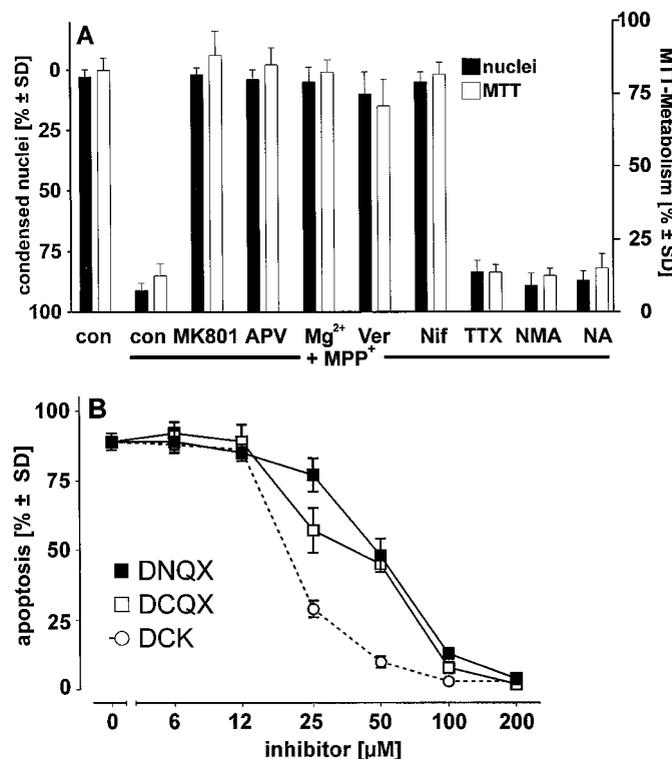


Fig. 2. Prevention of MPP⁺-induced CGC apoptosis by blockade of Ca²⁺ channels. *Top*, CGC were preincubated for 30 min with solvent (0.4% DMSO; *con*) or 2 μM MK801, 500 μM AP5, 5 mM Mg²⁺, 1 μM verapamil (*Ver*), 1 μM nifedipine (*Nif*), 500 nM tetrodotoxin (*TTX*), 400 μM *N*-methyl-L-arginine (*NMA*), or 1 mM nitroarginine (*NA*) before they were challenged with 50 μM MPP⁺. Medium was exchanged for CSS after 4 hr, and cell death parameters were determined after 18 hr. Apoptosis was quantified by scoring condensed neuronal nuclei (note *inverted scale*); mitochondrial function was determined by measuring MTT-reduction capacity. *Bottom*, CGC were incubated as described above with different concentrations of inhibitors of the NMDA-R glycine site (DNQX, 6,7-dichloroquinoxaline-2,3-dione, or DCK) plus 50 μM MPP⁺. Apoptosis was quantified after 18 hr. Data are mean \pm standard deviation of triplicate determinations.

Other NMDA-R antagonists such as AP5, a competitive inhibitor of the glutamate binding site, or high Mg^{2+} concentrations in the medium protected MPP⁺-treated neurons (Fig. 2A). In addition, inhibitors acting on the costimulatory glycine-binding site of the NMDA-R prevented MPP⁺ toxicity (Fig. 2B). The quinoxalinedione compounds used in this study (DNQX and CNQX) inhibit non-NMDA glutamate receptors at low concentrations ($\sim 20 \mu M$) and block NMDA-R action at three or four times higher concentrations. Prevention of MPP⁺ toxicity by these substances at concentrations of $\geq 50 \mu M$ points to the predominant role of the NMDA-activated subtype of glutamate receptors for this effect. DCK, the most selective inhibitor of the NMDA-R glycine site tested here, also was most potent in inhibiting MPP⁺-induced CGC apoptosis. This further suggests a role for the NMDA-R and excitotoxicity in MPP⁺-induced CGC apoptosis.

Activation of VDCC has been shown to contribute significantly to NMDA-triggered neuronal $[Ca^{2+}]_i$ increase (reviewed in Leist and Nicotera, 1997). Therefore, we tested whether pharmacological inhibitors of VDCC had a further modulatory effect on MPP⁺ toxicity. At intermediate MPP⁺ concentrations ($50 \mu M$) and limited exposure times, protection of CGC was observed with either the dihydropyridine-type VDCC blocker nifedipine or with verapamil (Fig. 2A). At MPP⁺ concentrations of $>50 \mu M$, or when exposure was prolonged for >4 hr, block of VDCC was not sufficient to prevent cell death.

It has been described previously that indirect excitotoxic damage in hippocampal cultures triggered by Mg^{2+} withdrawal and NMDA-R hypersensitization was blocked by the Na^+ channel blocker tetrodotoxin (Abele *et al.*, 1990). We therefore examined whether activation of Na^+ channels, which may propagate membrane depolarization, was necessary for MPP⁺ toxicity. In the case of indirect excitotoxicity triggered by MPP⁺ in CGC, tetrodotoxin had no protective effect (Fig. 2A). This is in line with findings that KCl-triggered indirect excitotoxicity in cortical neuronal cultures (Monyer *et al.*, 1992) or glutamate-dependent retinal damage due to inhibitors of energy metabolism (Zeevalk and Nicklas, 1991) is not significantly reduced by tetrodotoxin and thus also does not require the long-distance traveling of action potentials.

Rapid ATP depletion as a possible trigger of MPP⁺ excitotoxicity. Because energy depletion is known to trigger excitotoxic processes in different experimental systems, we examined intracellular ATP levels after treatment of CGC with MPP⁺. ATP declined slowly during the first 30 min of treatment with MPP⁺, and it was dissipated $\geq 90\%$ after 3–4 hr. Treatment with MK801 delayed the loss of ATP by 30–60 min; however, at 3–4 hr after exposure to MPP⁺, ATP also was depleted in these cells (Fig. 3A). These findings suggest that MPP⁺ caused a primary ATP depletion that was largely independent of the secondary excitotoxicity. This also implied that ATP depletion alone was not immediately lethal to neurons but rather sensitized them to glutamate and facilitated glutamate release. Under our culture conditions (residual glucose concentrations of ~ 1 mM in the medium), glycolytic ATP production alone was not sufficient to maintain intracellular ATP concentrations at control levels after exposure to MPP⁺. Accordingly, the addition of 10 mM glucose delayed MPP⁺ toxicity, which suggests that glycolysis re-

mained functional. Also, when we measured the glucose consumption of neurons, the basal rate of 20 nmol/10⁶ cells/hr increased to ~ 400 nmol/10⁶ cells/4 hr in the presence of MPP⁺, regardless of whether MK801 was included in the culture dish.

Prevention of mitochondrial damage by NMDA-R block. In this system, MPP⁺ may damage mitochondria by two ways: (i) by direct radical-induced damage and (ii) by triggering excitotoxicity, which results in Ca^{2+} overload and subsequent mitochondrial failure. We distinguished between these two mechanisms by comparing MPP⁺-triggered mitochondrial effects in the presence and absence of NMDA-R blockers. First, changes in $\Delta\Psi$ were examined by staining CGC with the mitochondrial potential-sensitive dye TMRE (Fig. 3B). CGC pretreated with MK801 (Fig. 3B) or AP5 (not shown) and then exposed to MPP⁺ showed no significant loss of $\Delta\Psi$. In neurons exposed to MPP⁺ alone, ATP depletion was paralleled by a complete loss of $\Delta\Psi$, in analogy with a model of direct excitotoxicity (Ankarcrona *et al.*, 1995). Thus, loss of $\Delta\Psi$ seemed to be due to indirect excitotoxicity and preceded nuclear condensation and loss of membrane integrity. MPP⁺-induced loss of TMRE fluorescence was selective for neurons because astrocytes within the same culture dish lost neither TMRE fluorescence nor viability. To confirm this finding, $\Delta\Psi$ also was monitored with the indicators JC-1 (Ankarcrona *et al.*, 1995) and Mito-Tracker Red (CMX-rosamine). All dyes yielded similar results, but TMRE was most convenient for routine measurements because it was less photolabile than JC-1 and more suitable for monitoring long time periods (24 hr) than CMX-Ros, which reacts with cellular thiols.

To determine mitochondrial outer membrane integrity, we measured release of cytochrome *c* from mitochondria into the cytosol. Treatment of CGC with MPP⁺ caused cytochrome *c* release within 90 min (Fig. 3C). Such mitochondrial damage was downstream to NMDA-R activation because it was entirely prevented by pretreatment of cells with MK801.

NMDA-R block inhibits loss of mitochondrial membrane potential and toxicity triggered by classic mitochondrial inhibitors. In view of these findings, we considered that NMDA-R-dependent toxicity might be triggered by other compounds affecting mitochondria. To evaluate this possibility, we exposed CGC to low concentrations of mitochondrial poisons such as oligomycin, CCCP, or rotenone. The latter blocks the respiratory chain at a site similar to that affected by MPP⁺ (Kilbourn *et al.*, 1997). Toxicity of oligomycin, CCCP, or rotenone was significantly ($\geq 50\%$ in all cases) reduced/delayed by MK801 (not shown). Pretreatment with MK801 completely prevented the toxicity of low concentrations of rotenone (25–50 nM) and maintained $\Delta\Psi$ in short term incubation (Fig. 3D).

Similarly, we found that apoptosis induced by low concentrations of the complex II inhibitor 3-NP or by high concentrations of DG was completely prevented by MK801 (Table 1). 3-NP also stimulated glucose consumption similarly to MPP⁺ in the presence or absence of MK801 (not shown). Moreover, all mitochondrial inhibitors caused a marked ATP depletion, regardless of the presence or absence of MK801. Notably, in all cases, $\Delta\Psi$ was maintained long after the ATP depletion. Residual cytosolic ATP or alternative energy substrates likely contributed to the maintenance of $\Delta\Psi$. This is suggested by the finding that $\Delta\Psi$ was dissipated when mitochondrial inhibitors were added together with deoxyglucose. Nev-

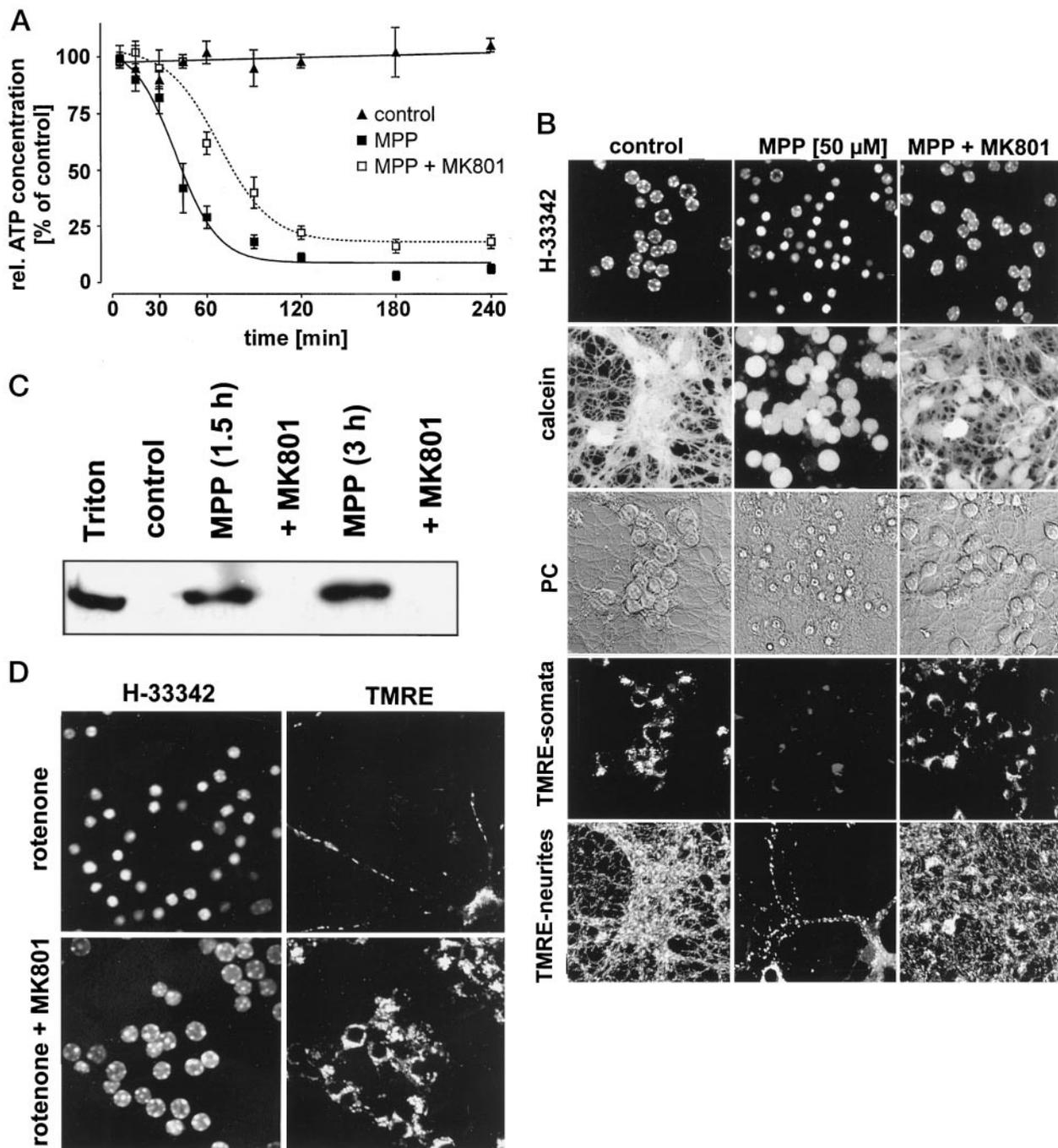


Fig. 3. NMDA-R and prevention of mitochondrial damage caused by MPP⁺. **A**, CGC were preincubated for 30 min with solvent (0.4% dimethylsulfoxide) or 2 μ M MK801 and then challenged with 50 μ M MPP⁺. At the times indicated, the medium was removed, cells were lysed, and ATP was determined luminometrically. Intracellular ATP content is expressed as percentage of that measured in untreated cells (control) at $t = 0$ (2.8 nmol of ATP/mg of protein). Viability and $\Delta\Psi$ were determined in parallel cultures from the same preparation. TMRE fluorescence did not significantly drop in CGC treated with MK801 + MPP⁺, whereas it was fully lost after 60–90 min in cells treated with MPP⁺ only. Viability of MPP⁺ + MK801-treated cells was maintained for ≥ 16 hr, whereas nuclear condensation in MPP⁺-treated neurons started after 45 min. Data represent mean \pm standard deviation of triplicate determinations. **B**, CGC incubated in control medium, with 50 μ M MPP⁺, or with MPP⁺ plus 2 μ M MK801 were stained after 4 hr with H-33342 (chromatin), calcein (cell membrane integrity), and TMRE (mitochondrial $\Delta\Psi$). Samples were imaged by confocal microscopy using fluorescence and polarization contrast (PC). Mitochondria from the axodendritic network (TMRE neurites) and those within the cell body (TMRE somata) were imaged at different focal planes. MPP⁺-treated neurons had condensed nuclei, whereas the plasma membrane (calcein staining) was still intact. TMRE staining in both neurites and somata disappeared. Only astrocytes (fifth row, middle: one astrocyte is shown with the cell body in the lower right corner and processes extending to the left) retained $\Delta\Psi$ (TMRE staining). MK801 prevented both nuclear condensation and loss of mitochondrial function. All experiments were repeated in at least six cell preparations with similar results. *Scale*, width of each micrograph corresponds to 108 μ m in CGC cultures. **C**, CGC were preincubated with or without 2 μ M MK801 and challenged with 50 μ M MPP⁺. Cytochrome *c* release into the cytosol was determined by Western blot. Cells treated with 0.5% Triton X-100 (*Triton*) were analyzed to indicate the maximal cytochrome *c* release. **D**, CGC were incubated with 25 nM rotenone or with rotenone plus 2 μ M MK801. They were stained after 4 hr with H-33342 (chromatin). In rotenone-treated cells, all TMRE staining (mitochondrial $\Delta\Psi$) was lost, and nuclei appeared condensed. Astrocyte nuclei and their mitochondria were unaffected (astrocyte in lower right corner with processes extending up and to the left). MK801 prevented nuclear condensation and loss of TMRE.

ertheless, also when both $\Delta\Psi$ loss and ATP depletion occurred, CGC were protected by MK801. Loss of $\Delta\Psi$ triggered by the complex I inhibitor MPP⁺ could be partially prevented by energizing mitochondria with the complex II substrate methylsuccinate (Table 1).

The role of NO generation and exocytosis in MPP⁺ toxicity. Endogenous NO production contributes to MPTP-induced pathology *in vivo* (Hantraye *et al.*, 1996). Thus, we tested whether endogenously formed NO was involved in our system. Inhibitors of the neuronal NO synthase did not reduce MPP⁺-induced apoptosis, nor did they modify other cytotoxicity parameters (Fig. 2A). One mechanism potentially involved in neuronal excitotoxic death is the activation of the enzyme PARP downstream to NO-mediated DNA damage (Zhang *et al.*, 1994). The mechanism of MPP⁺-induced excitotoxicity in CGC was, however, unrelated to PARP activation because neurons prepared from PARP^{-/-} mice were equally sensitive as those from wild-type mice (Fig. 1). Together, these findings suggested that MPP⁺ has the potential to activate excitotoxic mechanisms independent of NO generation.

A proapoptotic mechanism requiring NMDA-Rs involves the release of NMDA-R agonists from presynaptic terminals. This mechanism has been described in neurons treated with NO donors or with ONOO⁻ (Leist *et al.*, 1997a, 1997d). Prevention of MPP⁺-induced apoptosis by the competitive glutamate antagonist AP5 suggested that the release of NMDA-R agonists is a possible upstream mechanism, which also triggered excitotoxicity in this system. To test this assumption, we used the clostridial toxins BoNT/C and tetanus toxin, which are known to block exocytosis. Pretreatment of CGC with the toxins prevented apoptosis completely when cells were exposed to low MPP⁺ concentrations for a limited time (Fig. 4A). When the intensity of insult was increased (e.g., 100 μM MPP⁺), the clostridial toxins became less effective, although cultures were still completely protected by MK801 or AP5. These data suggest that with high levels of MPP⁺, apoptosis was not mediated by vesicular exocytotic glutamate release. The reversal of the glutamate transporter (Szatkowski and Attwell, 1994) may play a role under these circumstances. A role for increased extracellular glutamate concentrations is further suggested by experiments performed with inhibitors of the glutamate uptake system. α -Amino adipic acid (250 μM) significantly accelerated or enhanced MPP⁺-induced CGC apoptosis (not shown).

Further support for an excitotoxic mechanism at the base of MPP⁺-mediated neuronal apoptosis came from the findings that MPP⁺ triggered a sustained increase of $[\text{Ca}^{2+}]_i$ that was prevented by MK801 (Fig. 4, B and C). Similar data were obtained with rotenone (Fig. 5A). Despite the strong rise in $[\text{Ca}^{2+}]_i$, MPP⁺ challenged neurons retained the Ca^{2+} fluorescent indicators for up to 4 hr (i.e., they retained plasma membrane impermeability to the Ca^{2+} dyes when nuclei had already undergone condensation).

The role of proteases in MPP⁺- or rotenone-induced autocrine excitotoxicity. To further characterize the steps between NMDA-R-mediated $[\text{Ca}^{2+}]_i$ increase and cell death, we examined the role of cellular thiol proteases. Inhibition of either caspases or calpains was sufficient to block apoptosis elicited by MPP⁺ concentrations up to 50 μM (Fig. 6, A and B). The same set of inhibitors also protected CGC from the apoptosis induced by 50 nM rotenone ($\geq 80\%$ cells remained

viable in the presence of each of five different protease inhibitors) (Fig. 6E). Because all available inhibitors lack an absolute specificity for a single type of proteases (Villa *et al.*, 1997), we used three or five structurally different agents for each class of proteases with similar results. Inactive control peptides ($\leq 100 \mu\text{M}$) with similar end groups as the inhibitors were not effective. We also established sensitive *in vitro* assays with purified calpain or recombinant caspase-3 to test cross-reactivity of the inhibitors between the two classes of thiol proteases: for instance, z-D-cbk was found to be a highly potent inhibitor of caspase-3 ($\text{IC}_{50} < 200 \text{ nM}$) without any inhibitory effect on calpains ($\text{IC}_{50} > 200 \mu\text{M}$). Calp II showed exactly opposite characteristics. Thus, both caspases and calpains seemed to be required to mediate apoptosis of CGC challenged with 50 μM MPP⁺. Cells protected by protease inhibitors remained viable for ≥ 24 hr, when they received new medium or CSS. At very high intensities of insult (e.g., 100 μM MPP⁺; > 4 -hr toxin exposure), the protective effect of protease inhibitors was bypassed.

To gather direct evidence for intracellular proteolysis, we measured cleavage of fodrin (nonerythroid spectrin), a cytoskeletal protein, that is cleaved in neurons and various apoptotic cells by different isoenzymes of the caspase family, as well as by calpains (Nath *et al.*, 1996; Leist *et al.*, 1997d). MPP⁺-triggered fodrin proteolysis was prevented/reduced by different caspase and calpain inhibitors in a concentration-dependent fashion. The extent of inhibition of proteolysis at different protease inhibitor concentrations correlated with the extent of protection from apoptosis (Fig. 6C). Fodrin proteolysis also was blocked by MK801, but not by z-F-cmk, an inactive analogue of caspase inhibitors (Fig. 6D).

Protease inhibitors act downstream to the Ca^{2+} influx via the NMDA-R. Two sets of experiments were performed to test whether protease inhibitors indeed acted downstream of NMDA-R-mediated $[\text{Ca}^{2+}]_i$ increase. First, we measured NMDA-R-mediated $[\text{Ca}^{2+}]_i$ increases in CGC in the presence of protease inhibitors. None of the peptide inhibitors had a significant inhibitory effect (Fig. 7). Similar experiments were performed using as agonists glutamate, kainate, or high $[\text{K}^+]$. None of the protease inhibitors blocked $[\text{Ca}^{2+}]_i$ increase due to these stimuli (not shown).

In addition, we examined the protective effect of z-D-cbk and Ac-YVAD-cmk against glutamate and kainate-induced toxicity. Similar to the results published recently by Du *et al.* (1997a), we found small but significant protection (Fig. 6E). Notably, these protective effects were observed only at low concentrations of glutamate receptor agonists (12–50 μM), and in our experiments, only 20% of the neurons were rescued with any concentration of agonist and inhibitor. It seems that the degree of neuronal damage was more easily controlled, and more susceptible to pharmacological intervention, in the indirect excitotoxic model than after direct treatment of neurons with glutamate receptor agonists. This suggests that the role of caspases in excitotoxic neuronal apoptosis is more relevant at very low intensities of challenge or under conditions of indirect excitotoxicity.

The role of proteases and the NMDA-R in MPP⁺-induced chromatin breakdown. Caspase-3 activation has recently been shown to be a direct upstream mechanism leading to oligonucleosomal DNA-fragmentation. In CGC, oligonucleosomal DNA fragmentation was not significantly activated within the first 4 hr after challenge, when most

other processes relevant to cell death had already occurred. In glutamate-challenged CGC oligonucleosomal DNA fragmentation is a late event and may not be evident at all (Ankarcrona *et al.*, 1995). In analogy with the glutamate model, we observed here high molecular weight DNA fragmentation into 600-, 300-, and 50-kbp fragments in CGC treated with MPP^+ . This characteristic feature of apoptotic chromatin degradation was prevented by MK801 or by inhibitors of either calpains or caspases (Fig. 8A). This suggests the involvement of a proteolytic step, downstream to the initial $[Ca^{2+}]_i$ increase, which results in chromatin breakdown. This proteolytic step does not seem to be linked to caspase-3 activation. Consistent with the lack of oligonucleosomal DNA fragmentation and with the proteolysis pattern

of fodrin [caspase-3 forms a 120-kDa fragment, whereas caspase-4, caspase-2, or calpain forms mainly a 150-kDa fragment (Nath *et al.*, 1996)], we did not detect caspase-3-like (DEVD-afc cleavage) activity in CGC challenged with MPP^+ . Notably, DEVD-afc cleavage activity instead was easily detected in the same cell preparation challenged with colchicine or low $[K^+]$ (Leist *et al.*, 1997d). Further evidence for the absence of caspase-3 activation in the MPP^+ model is the lack of processing of procaspase-3 to the active caspase. Procaspase-3 was not cleaved even at a time point where fodrin was already $\geq 90\%$ cleaved and nuclei were condensed (Fig. 8B).

Prevention of MPP^+ -induced PS translocation by NMDA-R blockade and caspase inhibition. Translocat-

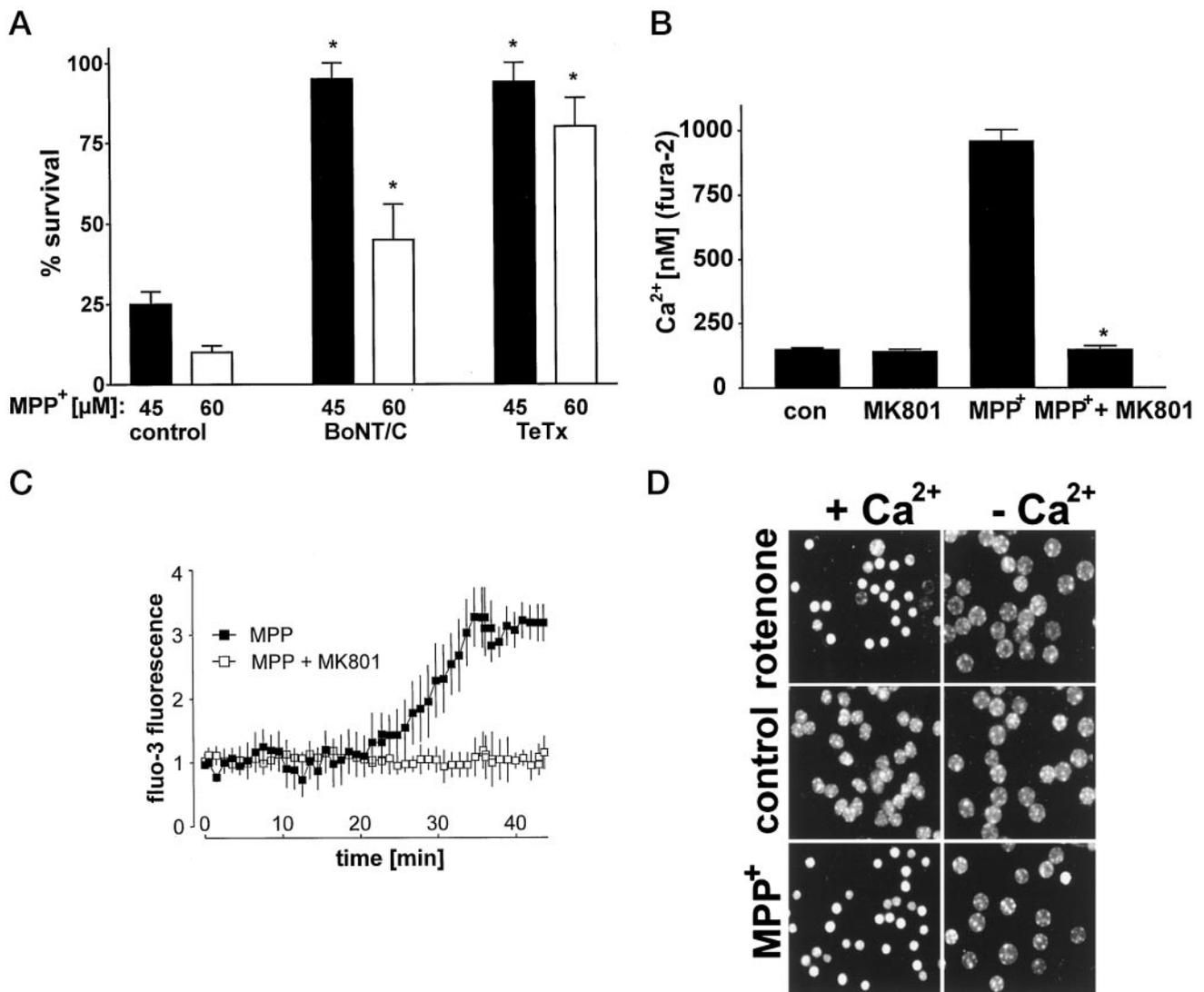


Fig. 4. MPP^+ -triggered exocytosis and autocrine excitotoxicity in CGC. **A**, CGC were preincubated for 10 hr with 20 ng/ml BoNT/C (130 pM) or 500 ng/ml tetanus toxin (3 nM) before they were challenged with MPP^+ . The medium was exchanged 4 hr after challenge for CSS, and survival was determined after 18 hr by MTT assay. Data are mean \pm standard deviation from triplicate determinations. *, $p \leq 0.05$. **B**, CGC were challenged with 50 μM MPP^+ \pm MK801. After 2 hr, CGC were loaded with Fura-2, and $[Ca^{2+}]_i$ was determined by ratiometric video imaging and *in situ* calibration. Data are mean \pm standard deviation from 16–20 neurons. Similar results were obtained in three different cell preparations. The $[Ca^{2+}]_i$ measured in MPP^+ -treated CGC is close to the saturation range of the indicator and therefore may be $\geq 1 \mu M$. *, $p \leq 0.05$. **C**, CGC were pretreated with DMSO alone (solvent) or 2 μM MK801 (MK801) and loaded with fluo-3 acetoxymethyl ester before they were challenged with 120 μM MPP^+ . The increase of $[Ca^{2+}]_i$ in time (relative fluo-3 fluorescence) was measured by confocal microscopy. Data are mean \pm standard error from 10–15 neurons. **D**, CGC were challenged in CSS containing 1.8 mM Ca^{2+} (+ Ca^{2+}) or no nominal Ca^{2+} (- Ca^{2+}) with 100 μM MPP^+ or 100 nM rotenone. After 3 hr, apoptosis was determined by staining of cells with H-33342 and counting of the percentage of condensed nuclei. In three independent experiments, the average apoptosis was $84 \pm 7\%$ (MPP^+ + Ca^{2+}), $88 \pm 8\%$ (rotenone + Ca^{2+}), and $\leq 10\%$ under all other conditions.

tion of PS from the inner to the outer surface of the plasma membrane is an early and specific event in apoptosis (Leist *et al.*, 1997b, 1997d), which seems to be dependent on caspase activation, at least in tumor cells. In CGC, MPP⁺ caused PS translocation, as detected by annexin V staining, within 1–2 hr. MPP⁺-treated neurons stained intensively when the first signs of chromatin condensation were visible and well before the nucleus became fully pyknotic. CGC remained stainable for several hours until membrane permeability was lost and membranes were degraded. Virtually all neurons with condensed chromatin had lost plasma membrane lipid asymmetry ($\geq 90\%$ annexin V-positive neurons at MPP⁺ concentrations of $\geq 50 \mu\text{M}$ in four independent experiments). Often, large spherical membrane blebs were formed from the axons, which also stained positively with annexin V and had an intact membrane. Notably, annexin V binding to neuronal

bodies was completely prevented by MK801 (Fig. 9) and was triggered by direct exposure to glutamate (not shown), which suggests that PS translocation is linked to NMDA-R-mediated Ca²⁺ influx and not to direct MPP⁺ actions. In addition, exposure to rotenone induced PS translocation, which was inhibited by MK801 (Fig. 5B). Annexin V labeling also was inhibited by the caspase inhibitor z-D-cbk, in agreement with the view that caspases also are pivotal for this feature of neuronal apoptosis (Fig. 9).

Discussion

MPP⁺ elicits autocrine excitotoxicity in CGC. Our results show that MPP⁺ induces apoptosis of cultured CGC by eliciting autocrine excitotoxicity. MPP⁺ can induce either apoptosis or necrosis *in vivo* (Tatton and Kish, 1997), and previous studies have shown that either form of cell death can be induced *in vitro*, by mechanisms unrelated to excitotoxicity (Marini *et al.*, 1989; Du *et al.*, 1997b). Low MPP⁺ concentrations predominantly elicit apoptosis, whereas high concentrations trigger necrosis (Hartley *et al.*, 1994; Du *et al.*, 1997b). In animals, removal of glutamatergic inputs (decortication), blockers of glutamate release, or NMDA-R antagonists reduce MPTP- and MPP⁺-induced striatal damage and dopamine depletion (Srivastava *et al.*, 1993) or loss of dopaminergic neurons in the substantia nigra (Turski *et al.*, 1991).

Because binding to complex I (Kilbourn *et al.*, 1997) is the primary known biochemical effect of MPP⁺, putative excitotoxic mechanisms should be secondary to the initial mitochondrial dysfunction. Energy failure sensitizes neurons to excitotoxicity (Henneberry *et al.*, 1989) and fosters the release of excitotoxic mediators (Szatkowski and Attwell, 1994). In addition, mitochondria represent a target of excitotoxic mechanisms (Leist and Nicotera, 1998), and mitochondrial damage can aggravate the initial damage by releasing Ca²⁺, reactive oxygen species, and factors essential for apoptotic protease activation (Leist and Nicotera, 1997). Therefore, the exact contribution of mitochondria to MPP⁺-stimulated excitotoxicity is complicated by the cyclic nature of the events (initial direct inhibition and further mitochondrial damage and ATP depletion caused by excitotoxicity).

Of all the cytotoxic events elicited by MPP⁺ and examined in this study, enhanced glucose consumption and ATP depletion were the only ones that were not ameliorated by NMDA-R blockade and thus seemed to be primary effects of MPP⁺. It has been suggested earlier that mitochondrial inhibition by NO would result in ATP depletion and disturb [Ca²⁺]_i homeostasis even in the presence of glucose (Brorson *et al.*, 1997). This is confirmed by our findings with both MPP⁺ and 3-NP. Therefore, ATP depletion caused by MPP⁺ seems to be the most plausible triggering event for excitotoxicity. Notably, both the collapse of $\Delta\Psi$ and the loss of cytochrome *c* were prevented when CGC were pretreated with the NMDA-R antagonist MK801 (i.e., major components of mitochondrial damage were the result of secondary excitotoxicity and not the direct effects of MPP⁺ on the respiratory chain). In agreement with this view, low concentrations of four other mitochondrial inhibitors did not cause massive mitochondrial failure *per se* (i.e., under conditions when the NMDA-R was blocked) but triggered rapid cell death when the NMDA-R was functional.

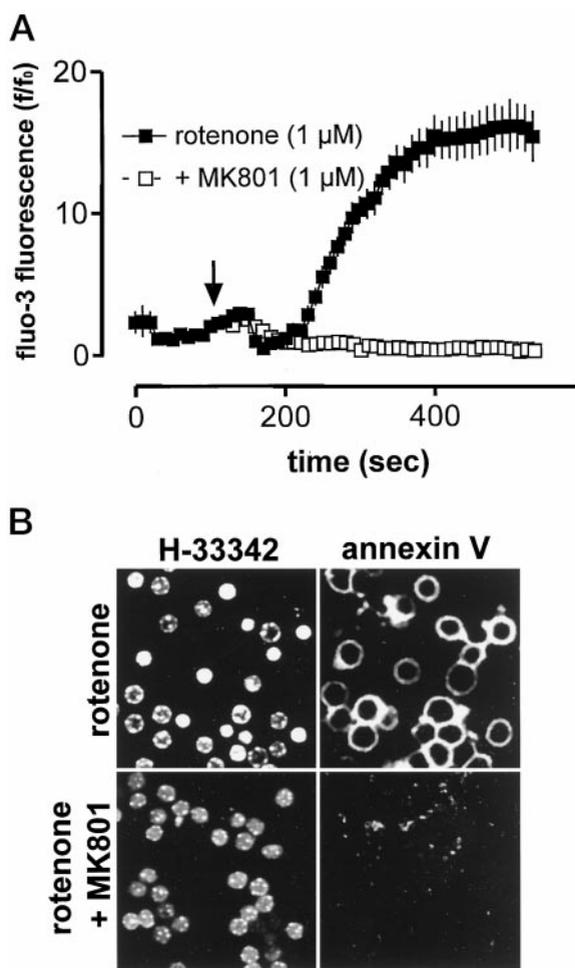


Fig. 5. MK801 prevents rotenone-induced Ca²⁺ increase and PS translocation. A, CGC were incubated with (open symbols) or without (filled symbols) 1 μM MK801. Rotenone (1 μM) was added as indicated (arrow), and the change of [Ca²⁺]_i was followed by confocal imaging of fluo-3 fluorescence. Data represent the mean \pm standard deviation from 15–25 cells in three independent experiments. B, CGC were pretreated with 1 μM MK801 or not treated. At 2 hr after the addition of rotenone (25 nM), cells were stained with fluorescent annexin V (PS translocation) and H-33342 (chromatin structure). Fluorescence images were collected by confocal microscopy. Note that cells with only marginally condensed nuclei are already annexin V positive. Phosphatidylserine translocation (annexin V stainability) was completely prevented by MK801. Data are representative for four experiments in three independent cell preparations.

Release of NMDA-R agonists in CGC treated with MPP⁺ may occur by two different mechanisms (Szatkowski and Attwell, 1994). First, release may occur by exocytosis. This process may be relevant with lower MPP⁺ concentrations and shorter exposure times, when ATP levels are still high enough to promote the fusion of neurotransmitter vesicles with the presynaptic membrane or when the pool of vesicles that has already been primed for exocytosis is released. The reduction in MPP⁺ toxicity by tetanus toxin or botulinum toxin suggests that this mechanism is operative in our system, at least with lower MPP⁺ concentrations. The relevance of this release mechanism for secondary excitotoxicity has also been demonstrated in models of oxygen-glucose depriva-

tion (Monyer *et al.*, 1992) or NO-triggered excitotoxicity (Leist *et al.*, 1997a). Second, glutamate may be released by the reversal of the glutamate transporter. This mechanism is likely to operate in CGC exposed to high MPP⁺ concentrations for longer times. It does not require ATP but rather is triggered by intracellular and extracellular ionic changes, which occur under conditions of ATP loss. Support for this hypothesis comes from the finding that intracellular ATP levels declined slowly immediately after exposure to MPP⁺ but then reached very low levels. At early stages, ATP production may have been maintained by residual mitochondrial production or glycolysis (Budd and Nicholls, 1996). Accordingly, the addition of 10 mM glucose or methylsuccinate to the

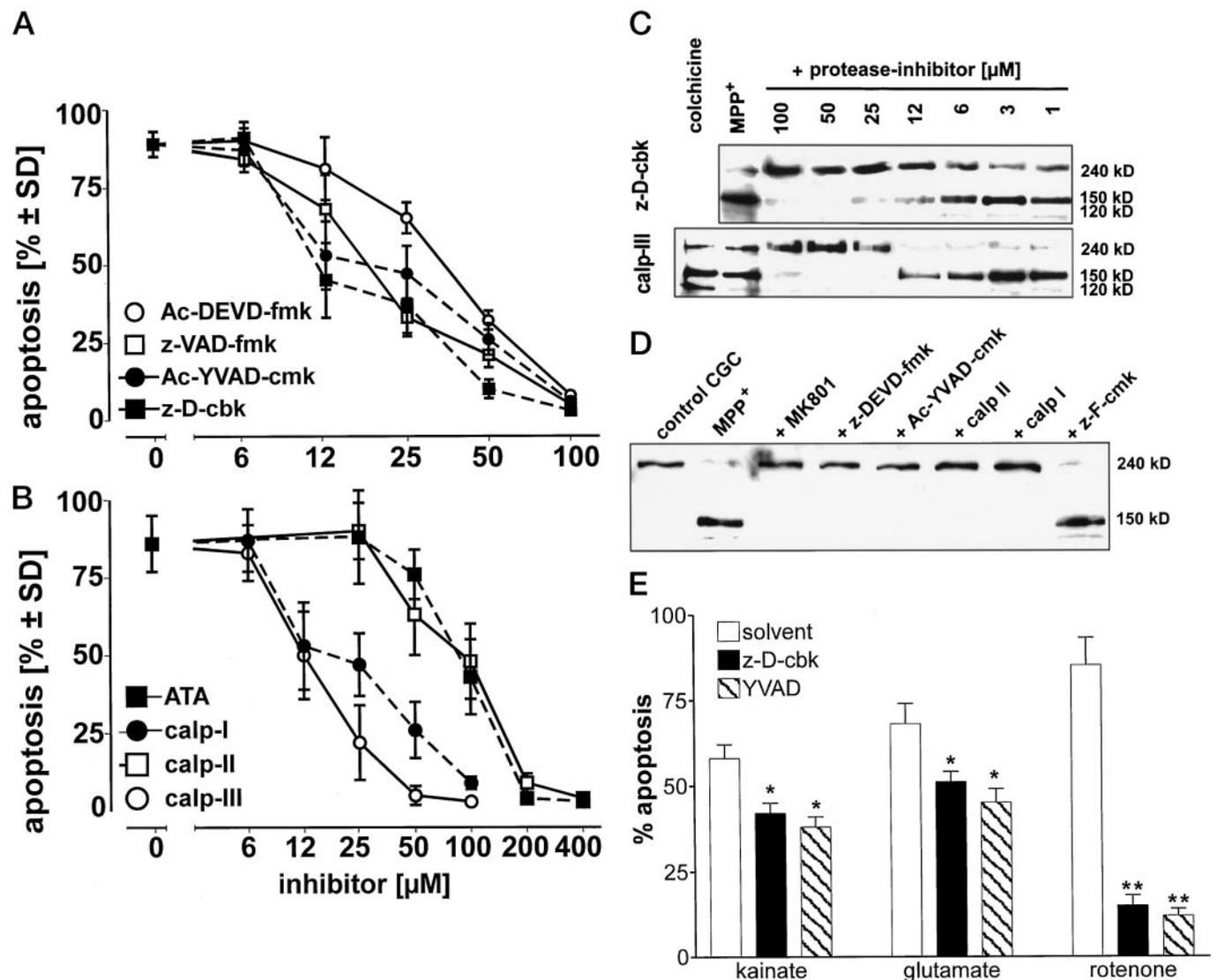


Fig. 6. Caspase or calpain inhibitors prevent MPP⁺ triggered intracellular proteolysis and apoptosis. CGC were incubated with (A) various caspase inhibitors or (B) calpain inhibitors at the concentrations indicated (logarithmic scale). At 30 min later, they were challenged with 50 μM MPP⁺. The percentage of apoptotic neurons was determined by counting the number of cell bodies. Data are mean ± standard deviation of triplicate determinations. C, Inhibition of fodrin proteolysis was studied by Western blotting of extracts from CGC pretreated with seven different concentrations of the caspase inhibitor z-D-cbk (top) or the calpain inhibitor calp III (bottom) and challenged with 50 μM MPP⁺. As positive control for fodrin proteolysis, we used the proapoptotic toxin colchicine (1 μM). Colchicine treatment (12 hr) produced proteolytic fragments of 150 and 120 kDa (bottom, first lane), whereas MPP⁺ (4-hr treatment) mainly produced the 150-kDa fragment. D, CGC were pretreated with various caspase or calpain inhibitors (100 μM) or MK801 (2 μM), or the inactive caspase inhibitor analogue z-F-cmk, before they were challenged with MPP⁺ (50 μM) for 4 hr. Fodrin proteolysis was analyzed by Western blot. E, CGC were pretreated with z-D-cbk (100 μM) or Ac-YVAD-cmk (YVAD; 100 μM) before they were exposed to 50 μM kainate, 25 μM glutamate, or 50 nM rotenone for 90 min. Then, medium was exchanged for CSS plus glucose. Apoptosis was quantified by staining neurons with H-33342 and determining the percentage of condensed nuclei. Data are mean values from triplicate determinations. *, $p < 0.05$; **, $p < 0.01$ solvent versus inhibitor.

culture medium significantly delayed MPP⁺-induced apoptosis and ATP depletion.

Mechanisms of indirect excitotoxicity. Several mechanisms have been implicated in direct excitotoxic neuronal death, including excessive NO production and subsequent activation of PARP (Zhang *et al.*, 1994), mitochondrial alterations (Ankarcrona *et al.*, 1995), and protease activation (Siman and Noszek, 1988; Du *et al.*, 1997a). Although brain NO synthase has an aggravating role in some models of excitotoxicity, NO plays a minor role as a direct mediator of toxicity in CGC (reviewed in Leist *et al.*, 1997a). CGC strongly express nitric oxide synthase and therefore may have developed mechanisms to prevent direct NO toxicity. Consistent with this assumption, MPP⁺ toxicity was not altered by nitric oxide synthase inhibitors. PARP activation is triggered by DNA damage, which may be caused by NO (Zhang *et al.*, 1994). We found no alteration in the cell death rate or mode of cell death in cells prepared from PARP^{-/-} mice, which further supports the lack of involvement of NO in MPP⁺ toxicity. In contrast, in CGC stimulated with MPP⁺, mitochondrial dysfunction and protease activation seem to be key events, which mediate excitotoxic cell death.

Proteases in excitotoxic death. Apoptosis is associated with the activation of a proteolytic cascade, probably involving different sets of proteases, which operate virtually at all stages of the cell death program (i.e., signaling, control, and execution) (Villa *et al.*, 1997). Here, we found that both caspases and calpains were activated downstream to the NMDA-R-mediated [Ca²⁺]_i increase and upstream of PS translocation, nuclear condensation, and DNA fragmentation. Under appropriate conditions, cells pretreated with protease inhibitors survived MPP⁺ challenge for several days. Obviously, different sets of proteases can interact to cause neuronal death; examples include caspases and calpains (Nath *et al.*, 1996; Jordán *et al.*, 1997; current study), different caspases and serine proteases (Stefanis *et al.*, 1997), and caspases plus the proteasome.

Very recent findings describe caspase-3 activation in neurons exposed to MPP⁺ (Du *et al.*, 1997d). In this model, cell

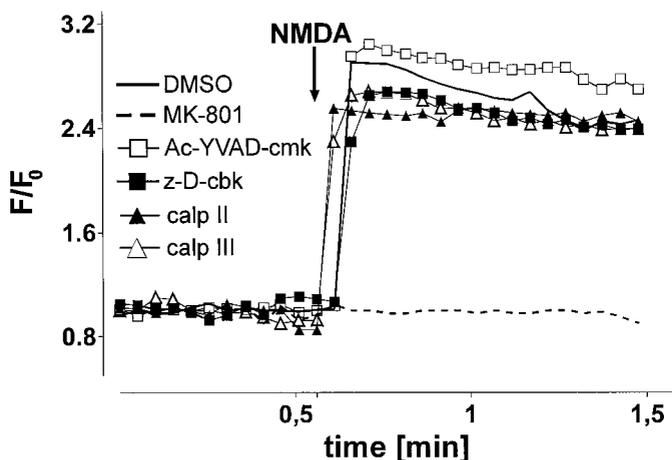


Fig. 7. Protease inhibitors do not inhibit NMDA-R-mediated neuronal Ca²⁺ increase. CGC were pretreated with various protease inhibitors (100 μM) or MK801 and loaded with fluo-3. They were challenged with 200 μM NMDA as indicated, and the fluorescence increase (F/F₀) was recorded by confocal microscopy. Data were obtained from 10–15 neurons, and experiments were repeated in three different CGC preparations.

death was slow (72 hr) and did not involve excitotoxic mechanisms. In contrast, caspase-3 activity did not seem to be involved in our experimental paradigm: (i) procaspase-3 was not proteolytically activated; (ii) MPP⁺ did not elicit DEVD-afc cleavage activity (specific for caspases-3/7), which is instead stimulated by several other proapoptotic agents under similar conditions (Leist *et al.*, 1997d); and (iii) the cleavage pattern of fodrin was not typical for caspase-3.

Excitotoxic apoptosis. Excitotoxic cell death may occur by either apoptosis or necrosis. Seemingly divergent observations are reconciled by the finding that the intensity of insult may determine the mode of cell death (e.g., Ankarcrona *et al.*, 1995; Du *et al.*, 1997a; Leist and Nicotera, 1998) and that excessive Ca²⁺ entry may convert the mode of cell death in some cases from apoptosis to necrosis (reviewed in Leist *et al.*, 1997d). Intracellular ATP levels seem to be critical in determining the shape of cell death (Ankarcrona *et al.*, 1995; Leist *et al.*, 1997c), and ATP may be required at multiple sites. For instance, ATP/dATP is required for the acti-

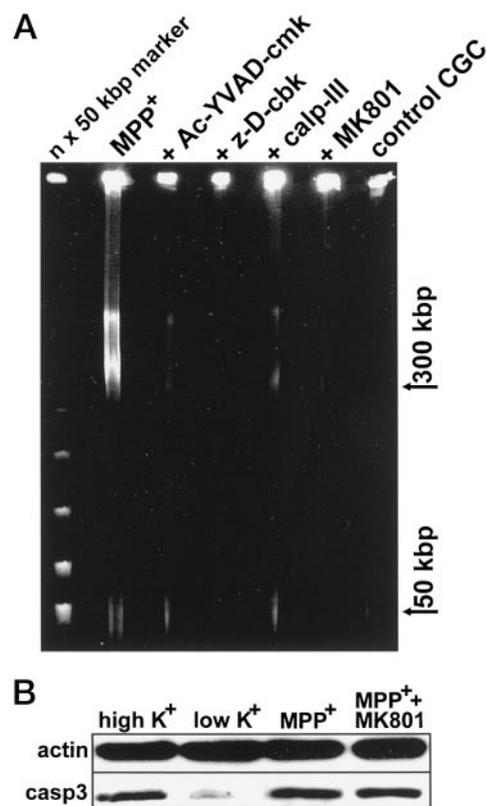


Fig. 8. MPP⁺ triggers NMDA-R- and caspase-dependent apoptotic chromatin degradation but not procaspase-3 proteolysis. A, CGC were preincubated for 30 min with 100 μM Ac-YVAD-cmk, z-D-cbk, calp III, or 2 μM MK801 or solvent control only. Then, they were challenged with 50 μM MPP⁺ or they were left unchallenged (control CGC). After 4 hr, incubations were stopped and DNA fragmentation was analyzed by field-inversion gel electrophoresis. Less than 5% of CGC had lost membrane permeability at that time point. B, CGC were challenged with MPP⁺ in the presence and absence of MK801. Cells were harvested after 3 hr, when fodrin was already maximally cleaved (see Fig. 6) and >80% of apoptotic nuclei were scored. Alternatively, apoptosis was induced by switching cells to low K⁺ medium for 24 hr (low K⁺; ~75% apoptotic neurons). Control cells were incubated for 24 hr in high K⁺ (<10% apoptotic neurons). A decrease of procaspase-3 (*casp3*) due to activation of the protease is seen in the low K⁺ model but not in the MPP⁺ model. Activated murine caspase-3 was not detected by the antibody used. Therefore, the blot membrane was redeveloped after stripping with an anti-actin antibody to confirm equal protein loading.

vation of caspase-3, which is a central protease in many apoptosis models. Here, we show that apoptosis proceeds in the absence of procaspase-3 processing. This also may explain the lack of oligonucleosomal DNA fragmentation, which is a direct consequence of caspase-3 activation. In MPP⁺-treated neurons, other cell death proteases may be activated by mechanisms not directly requiring ATP.

This work and previous evidence suggest that in mild excitotoxicity, caspases are among the direct executors of apoptosis. In addition to their degradative role in cell death, caspases may participate in the multiple positive feedback reactions that characterize secondary excitotoxicity; examples may include cleavage of proteins regulating [Ca²⁺]_i homeostasis or energy generation. Multiple further positive feedbacks link reactive oxygen species, glutamate release, disturbed [Ca²⁺]_i homeostasis, mitochondrial defects, energy depletion, and proteolysis. Establishment of such vicious circles may be blocked at an early stage by inhibiting any of the looping processes. Accordingly, we found that blocking the cycle of events at different sites (clostridial toxins, NMDA-R antagonists, antiproteases, glucose) prevented cell death and the associated biochemical features.

In neuropathological situations, these reciprocal interactions would form the basis of intricate vicious loops that are not likely to be interrupted by a single agent once they are fully established. In line with this view, caspase inhibitors

only partially protect CGC lethally challenged with glutamate receptor agonists. Also, indirect excitotoxicity due to NO or MPP⁺ is prevented by protease inhibitors only when the insult is relatively mild (Leist *et al.*, 1997d; current study). This may become important for the design of therapeutic strategies. Protease inhibitors may become maximally effective in preventing neuronal death only when combined with agents reducing the overall intensity of the insult (e.g., glutamate antagonists). The elucidation of key sequences of these vicious circles and a possible hierarchy of events is the subject of ongoing research.

Acknowledgments

We gratefully acknowledge the excellent technical assistance of H. Naumann and T. Schmitz. We are grateful to Dr. Zhao Qi Wang (IARC, Lyon, France) for the gift of the PARP^{-/-} mice and to Dr. C. Montecucco (University of Padova, Padova, Italy) for the clostridial toxins.

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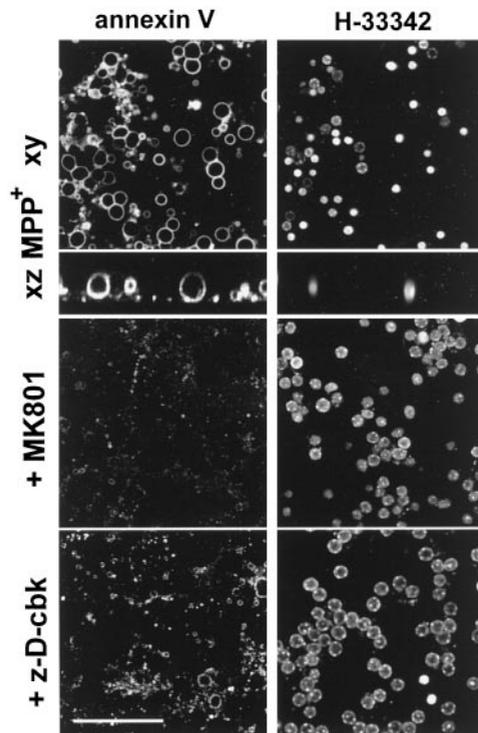


Fig. 9. MPP⁺ triggers NMDA-R- and caspase-dependent apoptotic PS translocation. CGC were incubated with MPP⁺ for 2 hr before they were stained with FITC-annexin V and H-33342 and imaged by confocal microscopy. *Top*, an *xy* section through the cell bodies (4 μ m from coverslip). Apoptotic nuclei seem to be condensed and hyperfluorescent and are surrounded by annexin V-stained membranes. A transversal section (*xz* plane) demonstrates that the entire surface of the apoptotic neurons plus some non-chromatin-containing blebs formed from the axonal projections are surface stained with annexin V. Neurons that had been preincubated with z-D-cbk or MK801 did not condense and did not stain with annexin V.

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