

ATP Controls Neuronal Apoptosis Triggered by Microtubule Breakdown or Potassium Deprivation

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Abstract

Background: Early loss of neurites followed by delayed damage of neuronal somata is a feature of several neurodegenerative diseases. Death by apoptosis would ensure the rapid removal of injured neurons, whereas conditions that prevent apoptosis may facilitate the persistence of damaged cells and favor inflammation and disease progression.

Materials and Methods: Cultures of cerebellar granule cells (CGC) were treated with microtubule disrupting agents. These compounds induced an early degeneration of neurites followed by apoptotic destruction of neuronal somata. The fate of injured neurons was followed after co-exposure to caspase inhibitors or agents that decrease intracellular ATP (deoxyglucose, *S*-nitrosoglutathione, 1-methyl-4-phenylpyridinium). We examined the implications of energy loss for caspase activation, exposure of phagocytosis markers, and long-term persistence of damaged cells.

Results: In CGC exposed to colchicine or nocodazole,

axodendritic degeneration preceded caspase activation and apoptosis. ATP-depleting agents or protein synthesis inhibition prevented caspase activation, translocation of the phagocytosis marker, phosphatidylserine, and apoptotic death. However, they did not affect the primary neurite loss. Repletion of ATP by enhanced glycolysis restored all apoptotic features. Peptide inhibitors of caspases also prevented the apoptotic changes in the cell bodies, although the axodendritic net was lost. Under this condition cell demise still occurred 48 hr later in a caspase-independent manner and involved plasma membrane lysis at the latest stage.

Conclusions: Inhibition of the apoptotic machinery by drugs, energy deprivation, or endogenous mediators may result in the persistence and subsequent lysis of injured neurons. In vivo, this may favor the onset of inflammatory processes and perpetuate neurodegeneration.

Introduction

Apoptosis is a mode of cell death by which superfluous or dysfunctional cells are removed from tissues. However, an increased rate of apoptosis may also be a component of neurodegenerative diseases (1,2). Frequently, the apoptotic execution phase is characterized by activation of caspases that cleave a defined subset of cellular

proteins (3). This leads to characteristic morphological changes such as nuclear condensation (4), and to the display of phagocytosis recognition molecules on the cell surface (5,6). The latter process is responsible for the swift removal of dying cells before the plasma membrane lyses (7). Rapid and efficient phagocytosis may also explain the low frequency of apoptotic cells, which can be observed in neurodegenerative disorders (1,8,9). In fact, in Alzheimer's disease, the persistence of dysfunctional (10) and degenerating neurons is more conspicuous than apoptosis (11). Neurons initially undergo cytoskeletal

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changes and lose dendritic projections, until only the cell bodies seem to be left at late stages. Similar changes are common to several axonopathies (11).

The persistence of neuronal remnants in such conditions may be explained by a block or low efficiency of the mechanisms that lead to the activation of the protease families (primarily caspases) involved in exposure of recognition molecules. This would result in a defective disposal of dying cells (5–7). It has been shown that the activation of caspases by the cytochrome-c and Apaf-1-containing apoptosome requires dATP or ATP (12,13). Thus, severe disorders in energy metabolism may preclude the activation of caspases. In agreement with this is the finding that apoptosis of tumor cells is entirely suppressed when cellular ATP is depleted by 50–60% (14,15). Also, in neurons, there is circumstantial evidence for a similar control of cell death by ATP: in cultures exposed to glutamate (16) or 1-methyl-4-phenylpyridinium (MPP⁺) (17), only the neuronal subpopulation that maintains an adequate cellular energy charge dies with apoptotic features.

Thus, we designed this study to test whether manipulation of neuronal ATP levels by drugs or mediators, which affect energy metabolism, would compromise the execution and occurrence of cell death following an initial axonal degeneration. Cell death was triggered in cerebellar granule neurons (CGC) by the cytoskeletal disrupting agents colchicine or nocodazole, which are known to trigger caspase activation (18) and delayed neuronal apoptosis both *in vivo* and *in vitro* (19,20). Finally, we examined the long-term fate of neurons that were treated with microtubule poisons but were incapable of executing apoptosis because of caspase inhibition.

Materials and Methods

Materials

SYTOX and H-33342 were obtained from Molecular Probes (Eugene, OR). The caspase substrate, *N*-acetyl-Asp-Glu-Val-aspartyl-aminotrifluoromethylcoumarine (DEVD-afc), 1-methyl-4-phenyl-pyridinium (MPP⁺), cyclosporine A (CsA), and *S*-nitrosoglutathione (GSNO) were obtained from Biomol (Hamburg, Germany), and (+)-5-methyl-10,11-dihydro-5H-dibenzo[a,d]cyclohepten-5,10-imine (MK801) came from RBI (Biotrend Chemikalien GmbH, Köln, Germany). Caspase inhibitors *N*-acetyl-Asp-Glu-Val-Asp-

aldehyde (DEVD-CHO), *N*-benzyloxycarbonyl-Val-Ala-aspartyl-fluoromethylketone (zVAD-fmk), *N*-acetyl-Tyr-Val-Ala-aspartyl-chloromethylketone (YVAD-cmk), and *N*-benzyloxycarbonyl-aspartyl-2,6-dichlorobenzoyloxymethylketone (ZD-cbk) were obtained from Bachem Biochemica GmbH (Heidelberg, Germany). DEVD-fmk was from Enzyme Systems (Dublin, CA). Fluorescein-labeled annexin V (annexin V) was from Boehringer-Mannheim (Mannheim, Germany). FK506 came from Alexis (Grünberg, Germany), and L-[4,5-³H]-leucine was from Amersham-Buchler (Braunschweig, Germany). 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

Eight-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 (NIH guidelines and European Community Council Directive 86/609/EEC).

Cell Culture

Murine CGC were chosen for these studies and isolated as described previously (21). These cells form a highly complex axodendritic network consisting of over 95% of one single neuronal type. Dissociated neurons were plated on 100 µg/ml (250 µg/ml for glass surfaces) poly-L-lysine (MW >300 kDa) coated dishes at a density of about 0.25×10^6 cells/cm² (800,000 cells/ml; 500 µl/well, 24-well plate) and cultured in Eagle's basal medium (BME, Gibco) supplemented with 10% heat inactivated fetal calf serum (FCS), 20 mM KCl, 2 mM L-glutamine and penicillin-streptomycin, and cytosine arabinoside (added 48 hr after plating). Neurons were used without further medium changes after 8 days *in vitro* (DIV). At that time, the residual glucose concentration in the medium was about 1 mM. The cultures were exposed to colchicine or nocodazole in their original medium for most experiments in the presence of 2 µM MK801 and 2 mM Mg²⁺ to prevent NMDA receptor activation and excitotoxicity (22). All inhibitors were added 30 min before the microtubule disrupters, unless stated otherwise.

ATP Depletion Experiments and Repletion with Glucose

In initial experiments, the ATP depletors [deoxyglucose (DG), MPP⁺, or GSNO] were added 30 min before colchicine exposure and remained present throughout the experiment. To restore ATP, the culture medium was supplemented with glucose (0 or 10 mM) 3 hr before exposure to colchicine. In this setup, the ATP depletors were added 8 hr after colchicine addition. Measurements of ATP and of protein synthesis (leucine-incorporation) were performed after 9.5 hr. Apoptosis and caspase activity were quantitated after 18 hr. In some delayed ATP repletion experiments, colchicine-treated CGC were kept ATP-deprived by deoxyglucose for 20 hr. After that time glucose (10 mM) was added and nuclear condensation was followed over the following 4 hr.

ATP Repletion with Alternative Energy Substrates

To test alternative energy substrates that would not increase the cellular capacity to generate NADPH via the pentose-phosphate shunt, cells were incubated with deoxyglucose plus succinate, glutamine, glycerol, or several other substrates. Colchicine was added 60 min after the energy substrates. ATP was determined after 10 hr and apoptosis after 16 hr.

Viability Assays

Apoptosis and secondary lysis were routinely quantified by double staining of neuronal cultures with 1 μ g/ml H-33342 (membrane permeant, blue-fluorescent chromatin stain) and 0.5 μ M SYTOX (non-membrane permeant, green-fluorescent chromatin stain) as described previously (23). Apoptotic cells were characterized by scoring typically condensed nuclei. About 600–1000 cells were counted in nine different fields in 2–3 different culture wells, and experiments were repeated in at least three different preparations. In addition, the percentage of viable cells was quantified by their capacity to reduce 3-(4,5-dimethylthiazole-2-yl)-2,5-diphenyltetrasodium bromide (MTT) after incubation with 0.5 mg/ml MTT for 60 min. As an additional parameter to assess plasma membrane integrity, CGC were loaded with 0.5 μ M calcein-AM for 5 min (23). Oligonucleosomal DNA fragmentation was quantified as described before by an ELISA method (24).

Immunocytochemistry

CGC were grown on glass-bottomed culture dishes, fixed after the experiment with 4% paraformaldehyde, and permeabilized with 0.1% Triton X-100 in phosphate-buffered saline (PBS). To monitor cytoskeletal alterations, we stained neurons with an anti- β III tubulin monoclonal antibody (1:300, #clone5G8, Promega, Mannheim, Germany), which recognizes only neuronal tubulin, as described previously (20). As secondary antibody, we used an Alexa[™] (λ_{ex} = 488 nm, λ_{em} = 512 nm) coupled anti-mouse IgG antibody (1:300, Molecular Probes). CGC were embedded in PBS containing 50% glycerol and 0.5 μ g/ml H-33342 and imaged by confocal microscopy (TCS-4D UV/VIS confocal scanning system; Leica AG, Benzheim and Leica Lasertechnik, Heidelberg, Germany).

Protein Synthesis

Protein synthesis was analyzed as described previously (25). Briefly, ³H-leucine (2 μ Ci) was added to CGC in a volume of 10 μ l, in 24-well plates (150 MBq/ml final). After 90 min incubation, the medium was removed. Cells were washed three times with ice-cold trichloroacetic acid (TCA, 10%) and dried with methanol (-20° C). The cell layers were lysed (0.3 M NaOH, 1% SDS) for 12 hr at 37 $^{\circ}$ C, and incorporated radioactivity was detected by β -scintillation counting.

Visualization of Phosphatidylserine Translocation

Surface phosphatidylserine expression of cultures grown on glass-bottomed culture dishes was analysed by annexin V staining and confocal microscopy (TCS-4D UV/VIS confocal scanning system; Leica AG and Leica Lasertechnik) as described previously (18).

Enzymatic Assays

Caspase-3-like activity (measured by DEVD-afc cleavage) was assayed essentially as described before (26,27). CGC were lysed 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-block, pH 7.5. The fluorimetric assay was carried out in microtiter plates with a substrate concentration of 40 μ M and a total protein amount of 5 μ g. Glucose concentrations in the medium were determined by the hexokinase/glucose dehydrogenase method using a commer-

cial kit (Sigma). ATP was measured luminometrically after lysis of cells in ATP-releasing agent (Sigma) as described earlier (14,22). Total cellular glutathione was determined according to the enzymatic cycling method of Tietze (28). Briefly, cells grown in 12-well plates were lysed at different times with cold 0.1 mM HCl/10 mM EDTA and immediately frozen. After thawing, glutathione was determined from the supernatant by an automated enzymatic analyzer (ACP5040, Eppendorff, Hamburg, Germany). Control cells had a glutathione content of 420 ± 20 pmol/ 10^6 neurons.

Western Blot Analysis

CGC were lysed in RIPA buffer (150 mM NaCl, 50 mM Tris, 1% NP-40, 0.25% sodium deoxycholate, 1 mM EGTA) supplemented with protease inhibitors [1 mM phenylmethylsulfonyl fluoride (PMSF), 1 mM benzamide, 1 mM iodoacetate, 1 mM iodoacetamide, 40 μ M leupeptin, 10 μ g/ml antipain, 5 μ g/ml pepstatin]. Protein was determined using the bicinchoninic acid method (Bio-Rad, München, Germany).

Proteins were separated by SDS-PAGE with 5 μ g protein/lane on 8% polyacrylamide gels for fodrin, and 80 μ g protein/lane on 4–20% polyacrylamide gradient gels for caspase-3/-7, and then blotted onto nitrocellulose membranes (Amersham-Buchler) in a Bio-Rad semidry blotter at 2.5 mA/cm² for 60 min. Blots were blocked in TNT (50 mM Tris pH 8.0, 150 mM NaCl, 0.05% Tween 20) containing 5% dried milk/1% bovine serum albumin (BSA) and then incubated with either anti-fodrin monoclonal antibody (1:500, clone #1622, Chemicon, Temecula, CA), or anti-caspase-3 p17 polyclonal antibody (1:1000, R#MF393, generously provided by Dr. D. W. Nicholson, Merck-Frost, Quebec, Canada) or anti-caspase-7, p17 polyclonal antibody (1:1000, #R151, generously provided by Dr. G. Cohen, MRC Toxicology Unit, Leicester, UK). Specifically stained bands were detected by chemiluminescence (ECL, Amersham) using a polyclonal IgG horseradish peroxidase-coupled secondary antibody (1:1000, Pharmingen). As a positive control for caspase-3 detection, we used recombinant caspase-3 (generously provided by Dr. F. Fackelmayer, Department of Biology, University of Konstanz, Germany), which appeared as a 29 kDa band after isolation from bacterial lysates and which was processed autocatalytically by about 90% [17 kDa band; specific activity 70×10^6 pmol/(min \times mg protein)].

Statistics

Experiments to determine cell viability, ATP, protein synthesis, or caspase activity were run as triplicates and repeated in three- to eight-cell preparations. Statistical significance was calculated on the original data sets by means of the Student's *t*-test. When variances within the compared groups were not homogeneous the Welch test was applied. All Western blots were performed with samples from at least three independent cell preparations.

Results

Caspase Activation following Microtubule Disruption in Neurons

To depolymerize neuronal microtubules, we used the two chemically unrelated drugs—colchicine (1 μ M) and nocodazol (1 μ M) (EC_{50} of both drugs: 50–100 nM). The first alterations of CGC microtubule network became apparent 6 hr after exposure to colchicine or nocodazol in the axodendritic net. After 12 hr, while nuclear morphology was still normal (Fig. 1A), the microtubule organization was severely compromised, and the activity of caspase-3-related caspases (DEVD-afc cleavage activity) started to increase (Fig. 1B). At the same time, the processing of procaspase-3 (Fig. 1C) and procaspase-7 (not shown) was evident on Western blots. After 15 hr, the microtubule network was completely lost, leaving only a small ring enveloping the nucleus (Fig. 2). Meanwhile, caspase activity had greatly increased and a large percentage of nuclei displayed apoptotic chromatin condensation and fragmentation. Between 18 and 24 hr after the exposure to either of the two microtubule disrupters, the increase in caspase activity reached a maximum (40- to 60-fold compared to control neurons), oligonucleosomal DNA fragmentation was increased 25- to 30-fold, and the percentage of apoptotic neurons reached a maximum of about 80%.

To examine whether caspase activation by colchicine and nocodazol was entirely due to microtubule depolymerization, we used taxol, a drug that stabilizes microtubules and antagonizes the effects of colchicine (20) and nocodazole. Treatment of CGC with 1 μ M taxol up to 3 hr after colchicine addition prevented both microtubule breakdown and the increase in caspase activity (Fig. 2). In contrast, pharmacological inhibitors of the *N*-methyl-D-aspartate (NMDA) receptor (2 μ M MK801), of voltage-dependent

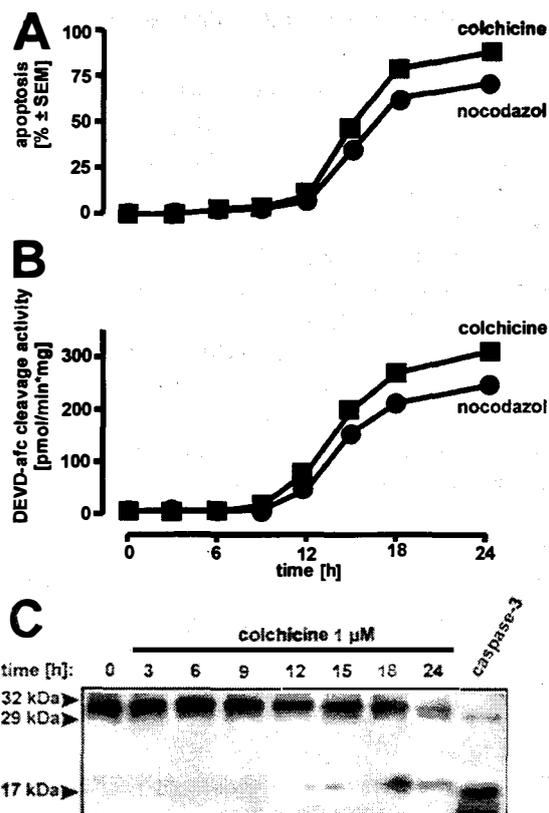


Fig. 1. Time course of caspase activation by microtubule disrupting agents. CGC were incubated with nocodazol (1 μ M) or colchicine (1 μ M). At the time points indicated, the percentage of apoptotic nuclei was counted in cultures stained with H-33342 (A). Enzymatic caspase-activity (DEVD-afc cleavage) (B) and processing of procaspase-3 (C) were determined in parallel cultures. Recombinant caspase-3 (20 ng/lane) was used as positive control for Western blotting. Data are means \pm SEM from seven determinations. Error bars are smaller than the data symbols.

Ca²⁺ channels (10 μ M verapamil) or of calcineurin- and cyclophilin-dependent proteins (2 μ M FK506; 1 μ M cyclosporine A) had no significant effect on colchicine/nocodazol-triggered caspase activation or on apoptosis. Thus, microtubule breakdown seemed to be the predominant upstream event leading to caspase activation in this model.

Prevention of Apoptosis by Caspase Inhibition

We investigated the causal role of caspases in apoptosis induced by microtubule disruption by using different irreversible halomethyl ketone caspase inhibitors and the reversible tetrapeptide aldehyde inhibitor DEVD-CHO. All inhibitors known to

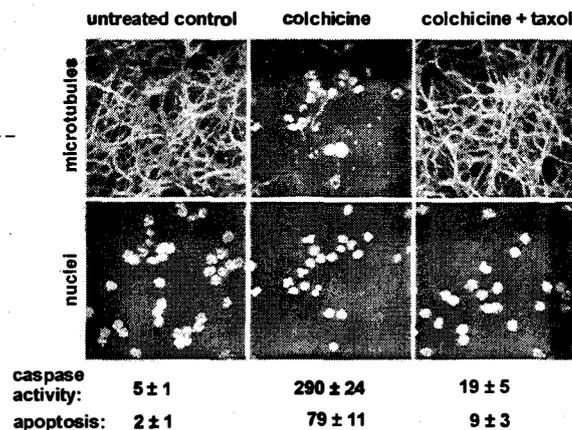


Fig. 2. Prevention of colchicine-induced caspase activation by taxol. CGC were treated with colchicine (1 μ M) in the presence or absence of taxol (1 μ M) for 18 hr. Parallel cultures were used for quantitating the percentage of apoptotic neurons (condensed chromatin) for measuring caspase activity [DEVD-afc cleavage in pmol/(min \times mg)] and for immunocytochemistry (staining of β III-tubulin). Nuclei were stained with H-33342 and correspond to the same field used for imaging the tubulin structure. The images were obtained by confocal microscopy ($\times 63$, NA 1.4 lens), and the width of every panel corresponds to 100 μ m.

block caspase-3-related caspases protected from colchicine-triggered apoptosis in a concentration-dependent manner (zD-cbk, zVAD-fmk: IC₅₀ = 2 μ M; DEVD-fmk: IC₅₀ = 15 μ M; DEVD-CHO: IC₅₀ = 250 μ M; Fig. 3A). Complete protection from apoptosis by zVAD-fmk was still observed when the inhibitor was added up to 10 hr after colchicine. In contrast, YVAD-cmk, which is more specific for caspases related to caspase-1, did not prevent apoptosis at concentrations up to 100 μ M (Fig. 3A).

Caspase activation was also tested by analyzing fodrin proteolysis. Fodrin was cleaved to 150 and 120 kDa fragments after exposure of CGC to colchicine (or nocodazol, not shown), and the proteolysis to the 120 kDa fragment [which is a specific marker for caspase-3 activity (29)] was inhibited by the caspase inhibitors that protected from apoptosis (Fig. 3B). The extent of inhibition of fodrin cleavage with different inhibitors correlated with the protective effect exerted by these agents on apoptosis. This finding further confirms that caspase-3-like caspases are the principal executors of neuronal apoptosis induced by microtubule breakdown.

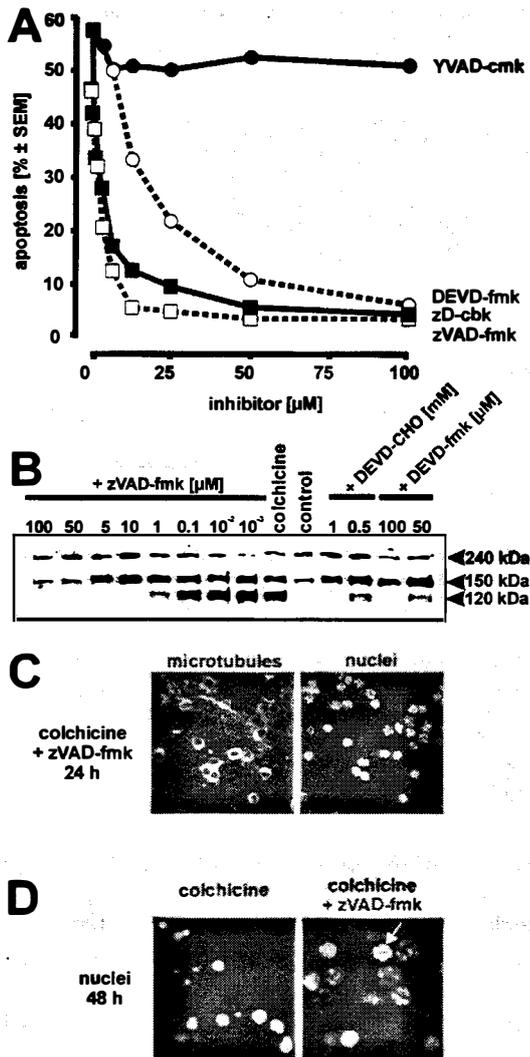


Fig. 3. Inhibition of colchicine-triggered apoptosis by caspase inhibitors. (A) CGC were treated with colchicine (1 μ M) in the presence or absence of different caspase inhibitors (YVAD-cmk, filled circles; DEVD-fmk, open circles; zD-cbk, filled squares; zVAD-fmk, open squares), and the percentage of apoptotic neurons was counted after 16 hr. Data are means \pm SEM from triplicate determinations. Essentially similar data were obtained with nocodazole as stimulus. (B) CGC were treated as above in the presence of different caspase inhibitors. At the end of the incubation cell lysates were prepared and analyzed by Western blot for cleavage of fodrin. (C) CGC were incubated for 24 hr with colchicine (1 μ M) in the presence of zVAD-fmk (100 μ M). Tubulin structure and nuclear morphology were imaged by confocal microscopy. The width of one image corresponds to 100 μ m. (D) CGC were incubated for 48 hr with colchicine (1 μ M) in the presence or absence of zVAD-fmk (100 μ M). Nuclei were stained with H-33342 and imaged by confocal microscopy. Cells that were treated with zVAD-fmk displayed marginated, but not condensed, chromatin (indicated by the arrow; for comparison apoptotic nuclei of cells that did not receive the caspase inhibitor are also shown). Note the scale difference (the image width corresponds to 25 μ m).

Delayed, Caspase-independent Cell Death in the Presence of zVAD-fmk

Inhibition of caspases by zVAD-fmk prevented chromatin condensation and death (as determined by calcein retention; MTT reduction), but did not protect neurons from the loss of the axodendritic net caused by colchicine or nocodazole (Fig. 3C). Between 48 and 72 hr after addition of colchicine, neuronal somata of CGC treated with zVAD-fmk were completely deprived of any neurites and died. This demise was characterized by nuclear alterations different from those observed in the absence of the inhibitor. Chromatin appeared to form lumps beneath the nuclear envelope (Fig. 3D), and these nuclear alterations were followed by a loss of plasma membrane integrity as indicated by the late penetration of non-vital dyes (not shown). Repetitive treatment with 100 μ M zVAD-fmk (cumulative concentration: 300 μ M; measured half-life in neuronal cultures >48 hr) did not prevent this form of cell death.

Prevention of Neuronal Apoptosis by ATP Depletors

We then investigated the effect of ATP depletion on caspase activity and CGC apoptosis. To avoid secondary excitotoxic effect (18,22), these experiments were all performed in the presence of the NMDA-receptor blocker MK801. When CGC were exposed to the glucose analogue 2-deoxyglucose (DG, 1–3 mM), the intracellular ATP concentration dropped below 50% within minutes and reached a level of 20% compared to untreated controls after 1 hr; then it further decreased during the following 20 hr (Fig. 4A). This treatment had no effect per se on neuronal viability (calcein uptake, SYTOX exclusion), neuronal morphology (phase contrast), chromatin structure (H-33342 staining), or mitochondrial membrane potential (tetramethylrhodamine methylester uptake) for at least 24 hr [Fig. 4B (22)]. Essentially similar findings were obtained when the two inhibitors of oxidative phosphorylation GSNO or MPP⁺ were used as ATP depletors [Fig. 4B (22,30)].

Pretreatment of CGC with DG abolished neuronal apoptosis triggered by microtubule disruption (Fig. 4B). Even delayed addition of DG up to 10 hr after colchicine exposure still prevented apoptosis. Thus, ATP-dependent steps were crucial after this time. GSNO and MPP⁺ also prevented colchicine-induced apoptosis to a similar extent (Fig. 4B).

We tested whether ATP depletion would prevent apoptosis in another paradigm of neuro-

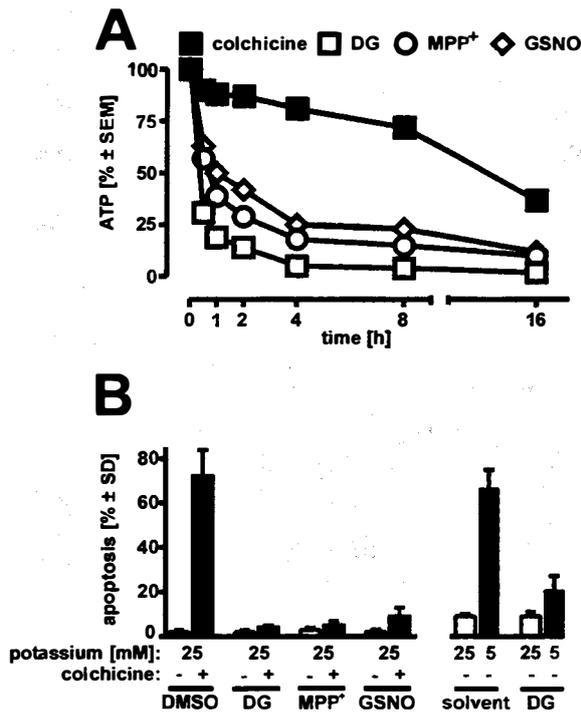


Fig. 4. Prevention of colchicine-triggered apoptosis by ATP depletion. (A) CGC were incubated with deoxyglucose (DG, 2 mM), MPP⁺ (50 μ M), GSNO (100 μ M), or colchicine (1 μ M), and cellular ATP was measured after the times indicated. ATP content in control neurons was 2.8 nmol/mg protein, and data are given as percent of untreated control cells. Data are means \pm SEM from quadruplicate measurements. (B) Left: CGC were incubated with colchicine (0 or 1 μ M) for 24 hr in the continuous presence of DG (2 mM), GSNO (100 μ M), MPP⁺ (50 μ M), or solvent control DMSO (dimethylsulphoxide) in normal culture medium containing 25 mM potassium. Right: medium was exchanged for a controlled salt solution (27,31) containing 0.5 mM glucose and either 5 or 25 mM potassium. In addition, cells were incubated with 0 or 2 mM DG. The percentage of apoptotic neurons was determined after staining with H-33342. Data are means \pm SD from three different experiments.

nal apoptosis. One of the best-characterized models in CGC is withdrawal of potassium (31). In this setup, apoptosis was significantly reduced by deoxyglucose when the salt solution used for the withdrawal situation contained low glucose concentrations (0.5 mM), but not after supplementation with the normally used high glucose concentrations (10 mM) (Fig. 4B).

Restoration of Caspase Activity and Cell Death by Repletion of Intracellular ATP

The ATP depletion elicited by DG, GSNO, and MPP⁺ was then antagonized by increasing the

glucose concentration in the culture medium. Supplementation of the medium with 10 mM glucose bypassed the effects of DG and restored cellular ATP levels as well as colchicine-triggered caspase activity and apoptotic death (Fig. 5A). As a corollary, the prevention of procaspase-3 processing by DG was also reversed by glucose (Fig. 5B).

Glucose may be metabolized glycolytically to generate ATP. Alternatively, it may be channeled into the pentose-phosphate shunt that provides cells with NADPH and therefore increases their capacity to cope with oxidative stress. To get clarity on the metabolic role of glucose, we also used alternative energy substrates that are not metabolized in the pentose-phosphate shunt. All substances that repleted ATP [10 mM glycerol (65%), 20 mM succinate (65%), 5 mM glutamine (60%)] restored the ability of CGC to undergo apoptosis (55%, 65%, 50%, respectively, compared to restoration with glucose). Nonrepleting substances (creatine, fructose, 1,6-fructose-bisphosphate) had no effect on apoptosis.

Glutathione measurements showed that the cellular content of this redox buffer is not significantly altered by incubation with DG or DG plus colchicine for up to 14 hr. This further suggests that DG acts by ATP depletion and not by increasing oxidative stress in neurons. We also found that the capacity of cells to initiate apoptosis is retained in CGC for at least 20 hr after incubation with DG and colchicine: when glucose was added to cultures after that time, apoptosis started immediately, reaching 27% after 60 min and 72% after 4 hr. Thus, it seems unlikely that caspases had been inactivated, e.g., by oxidative or nitritive stress.

Since CGC have a very high glycolytic capacity (32), increased glucose concentrations also compensated the ATP depletion affected by the mitochondrial inhibitors MPP⁺ and GSNO. As in the case of DG, caspase activity, caspase-3 processing, and the rate of apoptosis were modulated in parallel (Fig. 6).

When we examined the morphological features of degenerating neurons after exposure to colchicine, we noticed that ATP depletion (Fig. 5C) prevented all the features of neuronal demise, which were also precluded by the caspase inhibitors. The most relevant finding was that treatment with DG prevented colchicine-triggered exposure of phosphatidylserine (PS) on the cell surface, a key factor for recognition of apoptotic cells by phagocytes (5–7). Glucose

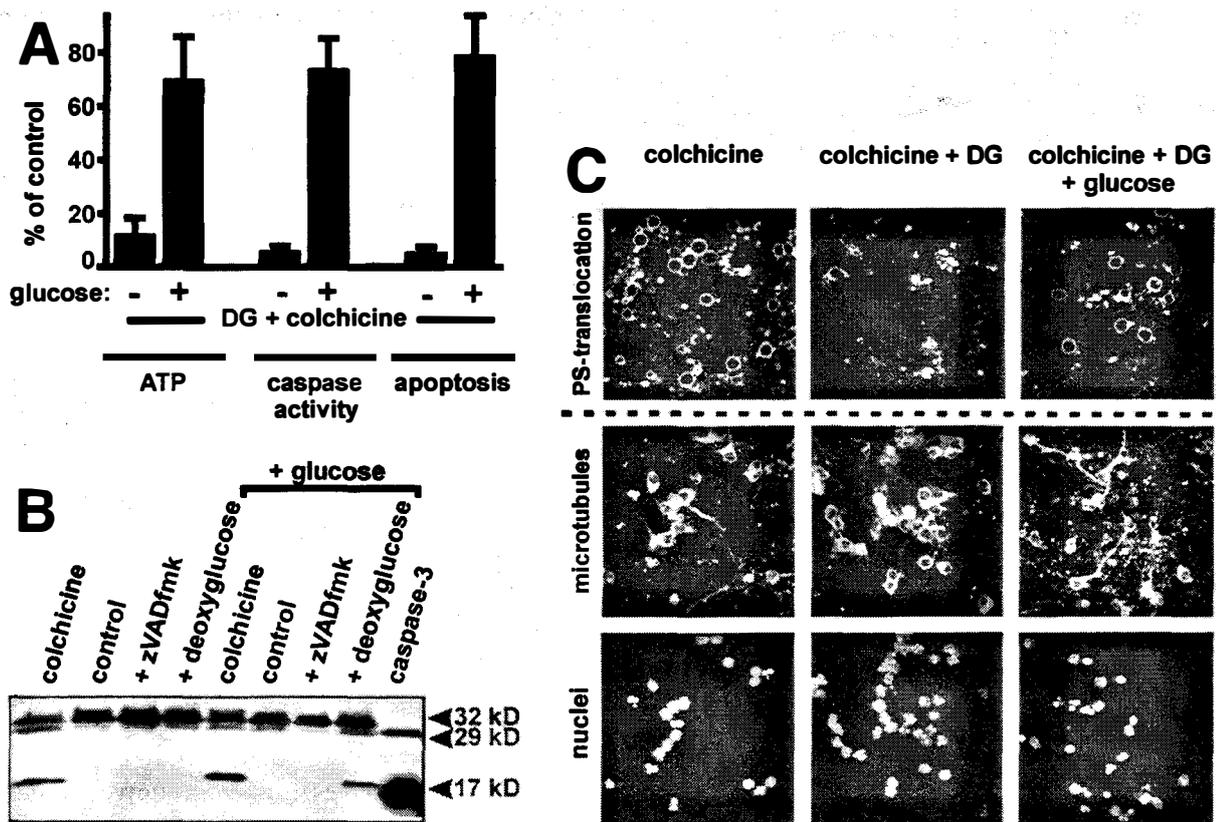


Fig. 5. ATP repletion, colchicine-triggered apoptosis, and caspase activity in DG-treated CGC. (A) Medium of CGC was supplemented with glucose (10 mM) or not. Neurons were then treated with colchicine (1 μ M). Deoxyglucose (DG, 2 mM) was added to all cultures 8 hr after colchicine treatment. ATP was measured 9.5 hr after addition of colchicine (1.5 hr after DG); caspase activity and apoptosis were quantitated after 18 hr exposure to colchicine (8.5 hr after DG). As a control, neurons were incubated with colchicine alone. Data from this incubation (see Figs. 1, 4) were used as 100% reference values. All data shown are expressed as a percentage of colchicine-alone values and are means from three different experiments. (B) CGC were incubated in

the absence or presence of glucose and of different inhibitors as described above. After 18 hr, cell lysates were prepared and analyzed by Western blot for caspase-3 processing. Recombinant caspase-3 was used as standard. (C) CGC were exposed to colchicine for 18 hr either directly in the presence of DG, or in the presence of DG plus glucose. Phosphatidylserine (PS) translocation was stained in live neurons with annexin V and imaged by confocal microscopy. Plasma membranes of >90% of all neurons were impermeable to ethidiumhomodimer at that time point. Microtubules and chromatin structure were examined in parallel cultures after fixation and staining with anti- β III-tubulin and H-33342.

clearly reversed the inhibition of apoptotic-related morphological features and the inhibition of PS exposure (Fig. 5C).

Control of Apoptosis by ATP due to Translation-dependent and Translation-independent Mechanisms

Protein synthesis depends on high ATP concentrations, and colchicine-induced apoptosis may require ongoing translation. We found that both cycloheximide (CHX) and actinomycin D (ActD) prevented caspase activation and apoptotic death. The two inhibitors of protein synthesis arrested the neuronal degeneration process at a

morphological stage similar to that observed with DG or zVAD-fmk (Fig. 7A). Pronounced protection from apoptosis by delayed addition of CHX was observed up to about 10 hr after colchicine addition. As shown in Figure 7B, the time point beyond which neurons could not be rescued from apoptosis by CHX treatment was similar to that observed for caspase inhibition and for ATP-dependent steps.

In view of these findings, we considered that ATP depletion may prevent apoptosis, at least in part, by a block of translation. To test this possibility, we measured the inhibitory effect on translation of different concentrations of DG or

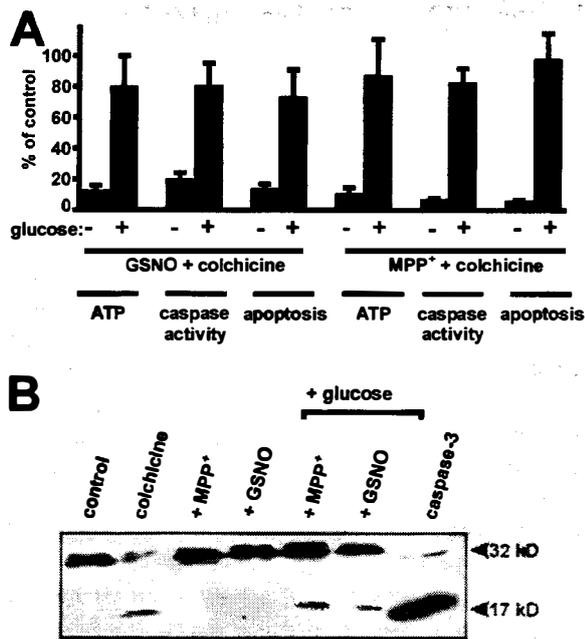


Fig. 6. ATP depletion, colchicine-triggered apoptosis, and caspase activity in CGC treated with GSNO or MPP⁺. (A) Medium of CGC was supplemented with glucose (10 mM) or not. Neurons were then treated with colchicine (1 μ M). GSNO (100 μ M) or MPP⁺ (50 μ M) was added to the cultures 8 hr after colchicine. ATP was measured 9.5 hr after addition of colchicine (1.5 hr after MPP⁺/GSNO); caspase activity and apoptosis were quantitated after 18 hr exposure to colchicine (8.5 hr after MPP⁺/GSNO). As a control, neurons were incubated with colchicine alone. Data from this incubation (see Fig. 4) were used as 100% reference values. All data shown are expressed as percentage of colchicine-alone values and are means from three different experiments. (B) CGC were incubated in the absence or presence of glucose and of different inhibitors as described above. After 18 hr, cell lysates were prepared and analyzed by Western blot for caspase-3 processing. Recombinant caspase-3 was used as standard.

CHX. The resulting data were compared with the corresponding concentration of either inhibitor required to prevent colchicine-triggered apoptosis (Fig. 7C). High DG concentrations (≥ 5 mM) were able to completely shut down protein synthesis, and may thus have inhibited apoptosis exclusively by this mechanism. On the other hand, low DG concentrations (≤ 1 mM) inhibited apoptosis, although inhibition of translation was minimal ($< 25\%$). Such a small extent of translational inhibition did not prevent apoptosis when CHX was used instead. Thus, ATP depletion by DG blocked apoptosis and caspase activation by an additional mechanism unrelated to the inhibition of translation.

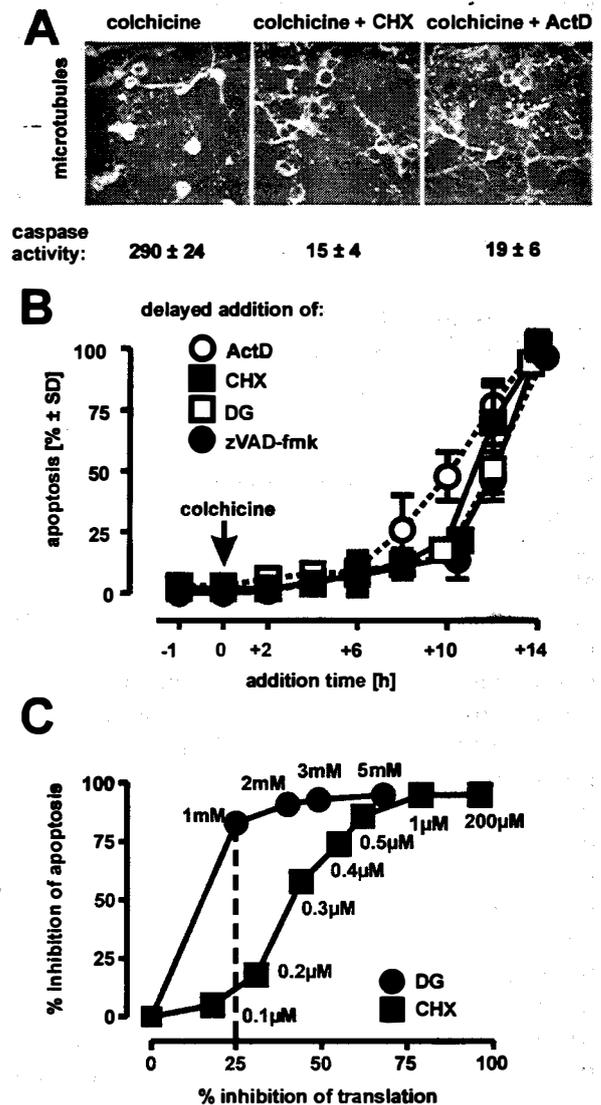


Fig. 7. Apoptosis prevention and translational block: a comparison of CHX and DG effects.

(A) CGC were exposed to colchicine for 18 hr in the presence or absence of CHX (1 μ M) or ActD (150 nM) until caspase activity (pmol/(min \times mg)) was determined enzymatically and tubulin structure was examined by immunostaining. (B) Different inhibitors (2 mM DG, 1 μ M CHX, 100 μ M zVAD-fmk, 150 nM ActD) were added to CGC cultures either 1 hr before colchicine (-1) or at different times after colchicine as indicated on the ordinate. All experiments were stopped 18 hr after colchicine addition. Data are presented as percent of the apoptosis, observed with colchicine alone. Experiments were performed 3-4 times and averages of all experimental data are shown. (C) CGC were incubated with different concentrations of CHX or DG at 8 hr after exposure to colchicine (1 μ M). Inhibition of translation by DG or CHX was measured in the period of 90-180 min after their addition; inhibition of apoptosis was measured 9 hr later. The different inhibitor concentrations are indicated in the figure. Data are averages of three experiments.

Discussion

Defects in mitochondrial energy generation have been frequently associated with neurodegenerative diseases (33–36). Such defects may have either a genetic basis (37,38) or be due to metabolic inhibition of oxidative phosphorylation, for instance, by nitric oxide (30,39–42). It is well established that ATP depletion facilitates damage by many stimuli and sensitizes neurons, particularly to excitotoxicity (34,43). In extreme cases, ATP depletion itself may be the main trigger for subsequent neuronal damage (22). Under such conditions, cell death generally involves sets of proteases different from caspase-3-related enzymes, regardless of whether it is apoptotic or necrotic (1,2,22). Also, such cell damage usually occurs very rapidly and affects neuronal somata at early stages.

Here we suggest that ATP deficiency occurring during neurodegeneration may have a second implication: to prevent the execution of apoptosis and delay cell death, at least under conditions when excitotoxicity is not involved. A preventive effect of lowered energy production on nuclear damage in the irradiated thymus was observed many years ago (44), and ATP depletion in lymphoid cells or in the liver is known to block the execution of apoptosis triggered by cytokines or other agents (14,15,45). The implications of lowering ATP levels during the execution of apoptosis have been less clear in neurons. After exposure of neurons to substances that trigger both ATP depletion and cell death (rotenone, glutamate, MPP⁺), apoptosis is prevalent under conditions of relatively minor ATP loss, whereas necrosis occurs when the ATP loss is more severe (1,17). In the present study, apoptosis and the associated caspase activation were triggered by agents that themselves have only minor effects on energy generation or consumption. Thus, ATP levels could be modulated independently from the pro-apoptotic stimuli. We used three different ATP depletors, which did not compromise neuronal viability. The first was deoxyglucose, which competes with glucose for both cellular uptake and glycolytic metabolism. This can be considered a model for reduced glycolytic flux in neurons and it offers the additional advantage of easy reversibility of the effect of deoxyglucose by increasing glucose supply to the neurons. The second ATP depletor, MPP⁺, is a model toxin for the induction of Parkinson's disease (46) and it was used here as a selective inhibitor of mitochondrial complex I (47). MPP⁺

depletes ATP in CGC kept under low-glucose conditions within minutes (22), but is otherwise nontoxic over periods of ≤ 24 hr, providing that secondary excitotoxic phenomena are prevented (22). Also, MPP⁺ does not affect glycolysis, so ATP could be replenished by providing high glucose concentrations to the medium. Under such conditions, CGC maintain high ATP levels in the complete absence of oxidative phosphorylation (32). Finally, neurons were exposed to the NO donor and endogenous NO carrier GSNO. This substance depletes ATP by inhibiting oxidative phosphorylation (30,39). As in the case of MPP⁺, the specificity of the effect can be probed by replenishing cellular ATP by enhanced glycolysis (30). It has been argued that in some cell types, NO donors may directly affect caspases, and therefore prevent apoptosis (48,49). However, this is not the case in CGC exposed to GSNO (18). As shown in the present study, the prevention of caspase activation and apoptosis by NO donors can be explained by their ability to deplete ATP, since the effects of GSNO on apoptosis were easily reversed by glucose.

The execution of several, currently known apoptotic degradation processes can proceed in the absence of ATP. For instance, nuclear condensation, PS exposure, protease activities, and DNase activities themselves do not require energy. However, one central step upstream of these activities is the activation of execution caspases within the so-called apoptosome complex. *In vitro*, reconstitution of this complex is only possible in the presence of ATP or dATP (12,13), and depletion of ATP by 50–70% in living cells prevents caspase activation (14,15,30). Conversely, glucose supplementation restores caspase activation and the apoptotic phenotype. Thus, inhibition of caspase activation at the level of the apoptosome is a plausible mechanism, which can explain the effects of ATP depletion observed in our system. Accordingly, we found that procaspases-3 and 7 were not processed in ATP-depleted neurons.

In addition to the apoptosome complex, a second target that is influenced by ATP depletion involves translation-dependent processes. In several nonexcitotoxic models of programmed neuronal death [e.g., growth factor withdrawal, potassium deprivation (50,51)], caspase activation and apoptosis are prevented by inhibitors of protein synthesis. Here we showed that colchicine- or nocodazole-induced apoptosis is also sensitive to agents or conditions that inhibit protein synthesis. High concentrations of DG or MPP⁺ de-

pleted ATP by >90%, and shut down protein synthesis by $\geq 95\%$. However, lower concentrations of deoxyglucose, which depleted ATP to a lesser extent and hardly caused any inhibition of protein synthesis (<25%), still prevented caspase activation. Thus, ATP depletion may affect neuronal apoptosis indirectly due to inhibition of protein synthesis and directly due to its effect at the level of caspase processing by the apoptosome.

Intraventricular colchicine injection in the brain is a well-characterized inducer of neuronal stress (52) and stress responses (53). In addition to microtubule depolymerization and cell cycle arrest, colchicine triggers apoptosis in many tumor cells (54) as well as in postmitotic CGC in vitro (18,20) and in vivo (19). Exposure of CGC to colchicine has several secondary effects, such as induction of NO synthase, *fos*-expression, and disturbance of Ca^{2+} homeostasis, all of which precede typical apoptotic processes, such as DNA fragmentation, activation of caspases, and cleavage of fodrin (20,18,22). Our present findings show that caspase activation and the subsequent apoptotic events are strictly dependent on microtubule depolymerization itself. This is suggested by the comparable effects of colchicine and nocodazol, which are chemically unrelated but cause microtubule disruption. In addition, block of microtubule depolymerization by taxol, which functionally antagonizes colchicine, prevented caspase activation and apoptosis. This confirms earlier suggestions that microtubule integrity may indeed be one of the general sensors important for the control of the cell death machinery (20,55). Recent data showing an effect of microtubule disruption on mitochondrial permeability transition (56) and on the phosphorylation state of Bcl-2 (55,57) suggest a link between microtubule integrity and suppression of caspase-activating mechanisms.

Although caspase inhibitors did not prevent microtubule breakdown and the loss of neuronal projections, the somata remained viable (as analyzed by MTT reduction, leucine incorporation, and calcein retention) for ≥ 36 hr. Thereafter, neurons started to die slowly with features that differed morphologically from those of classical apoptosis. These cells, while undergoing some kind of nuclear condensation and chromatin margination (as observed in pre-apoptotic stages), did not expose phosphatidylserine on the outer surface (data not shown) and underwent late cell lysis. This is consistent with previous observations that caspase inhibition can block apoptosis, but not entirely pre-

vent delayed cell death (14,30,58,59). Thus, our findings suggest that inhibition of caspases can explain the uncoupling of degenerative processes in neuronal projections from the eventual death of the cell. In particular, the observation that caspase inhibition hinders PS exposure but does not prevent late cell lysis raises the question as to whether pharmacological inhibition of caspases is always desirable in in vivo neurodegenerative conditions. Inhibition of caspases by synthetic peptides, as well as prevention of caspase activation because of a concomitant energy deprivation, would allow the persistence and late dissolution of severely damaged cells (57). Such conditions may foster the onset of inflammatory responses and a progression of disease.

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