Alternating metabolic pathways in NGF-deprived sympathetic neurons affect caspase-independent death

Louis K. Chang  
*Washington University School of Medicine in St. Louis*

Robert E. Schmidt  
*Washington University School of Medicine in St. Louis*

Eugene M. Johnson Jr.  
*Washington University School of Medicine in St. Louis*

Follow this and additional works at: https://digitalcommons.wustl.edu/open_access_pubs

Part of the *Medicine and Health Sciences Commons*

Recommended Citation  
https://digitalcommons.wustl.edu/open_access_pubs/627

This Open Access Publication is brought to you for free and open access by Digital Commons@Becker. It has been accepted for inclusion in Open Access Publications by an authorized administrator of Digital Commons@Becker. For more information, please contact engeszer@wustl.edu.
Alternating metabolic pathways in NGF-deprived sympathetic neurons affect caspase-independent death

Washington University School of Medicine, Saint Louis, MO 63110

Mitochondrial release of cytochrome c in apoptotic cells activates caspases, which execute apoptotic cell death. However, the events themselves that culminate in caspase activation can have deleterious effects because caspase inhibitor–saved cells ultimately die in a caspase-independent manner. To determine what events may underlie this form of cell death, we examined bioenergetic changes in sympathetic neurons deprived of NGF in the presence of a broad-spectrum caspase inhibitor, boc-aspartyl-(OMe)-fluoromethylketone. Here, we report that NGF-deprived, boc-aspartyl-(OMe)-fluoromethylketone–saved neurons rely heavily on glycolysis for ATP generation and for survival. Second, the activity of F$_0$F$_1$ contributes to caspase-independent death, but has only a minor role in the maintenance of mitochondrial membrane potential, which is maintained primarily by electron transport. Third, permeability transition pore inhibition by cyclosporin A attenuates NGF deprivation–induced loss of mitochondrial proteins, suggesting that permeability transition pore opening may have a function in regulating the degradation of mitochondria after cytochrome c release. Identification of changes in caspase inhibitor–saved cells may provide the basis for rational strategies to augment the effectiveness of the therapeutic use of postmitochondrial interventions.

Introduction

Apoptosis is a form of cell death that occurs during normal development and in pathological situations. Cells undergoing apoptosis exhibit certain characteristics, including cytoplasmic shrinkage, nuclear blebbing, and chromatin condensation (Kerr et al., 1972). The caspase family of cysteine proteases has a critical role in executing apoptosis (Cryns and Yuan, 1998). In the intrinsic, or mitochondria-dependent, pathway of apoptosis, caspase activation is regulated mainly by the release of cytochrome c from the intermembrane space of mitochondria (Li et al., 1997). The Bcl-2 family of proteins contributes to regulating cytochrome c release by integrating and conveying pro- and antiapoptotic signals to the mitochondria (Adams and Cory, 1998). Once in the cytosol, cytochrome c initiates a cascade of caspase activation by promoting the oligomerization of APAF-1 and the activation of procaspase-9 (Li et al., 1997). Other mitochondrial proteins, such as Smac/DIABLO (Du et al., 2000; Verhagen et al., 2000) and HtrA2 (Suzuki et al., 2001), are also released into the cytosol during apoptosis and may contribute to the regulation of caspase activity.

The critical function of caspases in apoptosis is underscored by observations that caspase inhibition prevents the appearance of many markers of apoptosis in neurons, including certain biochemical (Miller et al., 1997; Stefanis et al., 1999) and ultrastructural changes (Oppenheim et al., 2001). However, in many, if not all, systems, caspase inhibition does not prevent the ultimate death of these cells. This caspase-independent cell death often occurs with a delayed time course (Miller et al., 1997; Stefanis et al., 1999). The initiation of the apoptotic cell death pathway leads to a number of caspase-independent changes, including release of death-promoting factors, such as apoptosis-inducing factor (Susin et al., 1999), Endo G (Li et al., 2001), and HtrA2 (Suzuki et al., 2001), and changes in mitochondrial structure (Mootha et al., 2001). In fact, microinjection of neutralizing antibodies to AIF protects cortical neurons from some forms of caspase-independent death (Cregan et al., 2002). However, which of these changes critically regulate caspase-independent death, and whether these mechanisms vary in different models, is not known.

Here, we examined the bioenergetic status of NGF-deprived sympathetic neurons that were prevented from completing apoptosis by a caspase inhibitor. Removal of NGF from these neurons in vitro triggers a classic apoptotic death that recapitulates naturally occurring cell death that ensues within the superior cervical ganglion in vivo during the first week of life. This apoptotic death requires macromolecular...
Abbreviations used in this paper: ANT, adenine nucleotide translocase; BAF, boc-aspartyl-(OMe)-fluoromethylketone; CsA, cyclosporin A; COX IV, cytochrome oxidase subunit IV; CsA, cyclosporin A; JC-1,5,5'-tetrachloro-1',1',3',3'-tetracyethylrhodamine; VDAC, voltage-dependent anion channel.

Results
NGF-deprived, BAF-saved neurons maintain slightly lower levels of ATP
NGF-deprived sympathetic neurons that are prevented from completing apoptosis by caspase inhibition exist in a depressed metabolic state, as determined by a number of morphological and biochemical parameters (Deshmukh et al., 1996; Chang and Johnson, 2002). To determine the effect of caspase inhibition on the energetic status of neurons, we measured total cellular ATP by the luciferase/luciferin method in sympathetic neuronal cultures that were either maintained in NGF or deprived of NGF in the presence of BAF. As seen in Fig. 1 A, the total amount of ATP increased in NGF-maintained cultures over 9 d, whereas it decreased in NGF-deprived, BAF-saved cells. Most of the decrease in ATP in NGF-deprived, BAF-saved neurons, electron transport remains the primary mechanism by which this electrochemical gradient is formed in these cells. Finally, CsA does not affect these changes, but does attenuate loss of mitochondria that occurs in NGF-deprived, BAF-saved neurons. These findings may have important implications for strategies that target caspsases for therapeutic intervention.
lular ATP suggests ongoing metabolism in NGF-deprived, BAF-saved cells consistent with their ability to survive long periods of NGF deprivation (Chang and Johnson, 2002).

**NGF-deprived, BAF-saved neurons require glycolysis, but not oxidative phosphorylation, to generate ATP**

To identify which bioenergetic pathways are used to generate ATP within these cells, we determined whether acutely inhibiting either glycolysis or oxidative phosphorylation altered cellular ATP levels in NGF-maintained and NGF-deprived, BAF-saved neurons. We used total ATP, which is only a gross measure of cellular energy charge, to identify large perturbations in energy balance that were likely to affect cell survival. As seen in Fig. 2, inhibition of glycolysis, achieved by incubating cells in glucose-free medium supplemented with 5 mM 2-deoxyglucose and 1 mM pyruvate, for 2 h in NGF-maintained cells decreased total ATP to 66% of control. Treatment of cells with the F$_0$F$_1$ ATPase inhibitor, oligomycin, in standard, glucose-containing medium was used to inhibit ATP production by oxidative phosphorylation. In NGF-maintained cells, 5 µg/ml oligomycin caused a decrease in ATP levels to 75% of the control. These data indicate that both glycolysis and oxidative phosphorylation contribute to ATP production in NGF-maintained neurons. To determine the contributions of these pathways to ATP production in NGF-deprived, BAF-saved cells, cultures were synchronized by depriving them of NGF in the presence of BAF for 3 d, such that essentially all cells have released cytochrome c but few cells have become committed to die (Chang and Johnson, 2002). In NGF-deprived, BAF-saved cells, inhibiting glycolysis for 2 h caused a decrease in ATP levels to 32% of control, which is much greater than in NGF-maintained neurons (P < 10$^{-7}$). In contrast, inhibiting oxidative phosphorylation in NGF-deprived, BAF-saved cells had no effect on cellular ATP (Fig. 2). Thus, unlike NGF-maintained neurons, NGF-deprived, BAF-saved neurons do not require oxidative phosphorylation to maintain maximal ATP levels, but instead rely more heavily, if not solely, on glycolysis for ATP production.

**Glucose deprivation kills NGF-deprived, BAF-saved, but not NGF-maintained, neurons**

One prediction from the finding that NGF-deprived, BAF-saved cells rely more heavily on glycolysis for ATP generation than NGF-maintained neurons (Fig. 2) is that NGF-deprived, BAF-saved cells should be more dependent on glycolysis for survival and more sensitive to glucose deprivation. To test this hypothesis, the effects of altering the concentration of glucose in the medium on NGF-maintained and NGF-deprived, BAF-saved cells on survival were determined. As schematized in Fig. 3 A, the latter was performed by first depriving cultures of NGF in the presence of BAF for 2 d, to generate a synchronized population of neurons that had released cytochrome c but had not committed to die. At this time, cells were switched from the standard medium, which contains 5 mM glucose, to a medium containing 0, 5, or 25 mM glucose for 8 d, after which the proportion that had committed-to-die was determined by readdition of a medium containing NGF and 5 mM glucose for 7 d. As seen
in Fig. 3 B, survival in the presence of NGF was identical in a medium containing 0, 5, or 25 mM glucose medium. Because NGF-deprived, BAF-saved cells become committed-to-die during the course of the experiment, only roughly 40% of NGF-deprived, BAF-saved neurons incubated in 5 mM glucose after 10 d can be rescued by NGF (Chang and Johnson, 2002). In contrast, only 11% of NGF-deprived, BAF-saved cells maintained in a medium lacking glucose for the final 8 d of the 10-d period could be rescued. Thus, glucose deprivation decreases by roughly 75% the number of BAF-saved cells that are rescued by NGF. Increasing the glucose concentration to 25 mM did not increase the proportion of cells that could be rescued, suggesting that insufficient glucose in the standard culture medium does not underlie caspase-independent death of NGF-deprived sympathetic neurons. These data demonstrate that NGF-deprived, BAF-saved neurons were more sensitive to glucose deprivation than were NGF-maintained neurons, which were remarkably resistant to this insult.

**Oligomycin kills NGF-maintained neurons, but protects NGF-deprived, BAF-saved neurons**

Because NGF-maintained neurons, but not NGF-deprived, BAF-saved cells, require oxidative phosphorylation to maintain maximal ATP levels, a second prediction from the data in Fig. 2 is that inhibition of oxidative phosphorylation should be more detrimental to NGF-maintained neurons than to NGF-deprived, BAF-saved neurons. To examine the effect of inhibiting oxidative phosphorylation on Commitment 2, cells were deprived of NGF in the presence of BAF for 2 d and maintained for an additional 8 d in the presence of 5 μg/ml oligomycin to block oxidative phosphorylation, and subsequently rescued with NGF (Fig. 4 A). Consistent with our previous report (Chang and Johnson, 2002) and data in Fig. 3, the number of NGF-deprived, BAF-saved cells that could be rescued after 2 and 10 d of NGF deprivation decreased from roughly 85% of control to 35%, indicating that caspase-independent death occurred during this period. After 8 d of treatment with oligomycin, only 10% of NGF-maintained neurons survived (Fig. 4 B). Surprisingly, treatment with oligomycin increased the number of BAF-
saved neurons that could be rescued by NGF, from 35 to roughly 55% of control. Because only 85% of NGF-deprived, BAF-saved cells could be rescued at the time of oligomycin treatment, almost half of the cells that could be saved were protected by oligomycin. To illustrate the magnitude of this effect, CsA, a robust inhibitor of Commitment 2 (Chang and Johnson, 2002), increased the proportion of cells that were rescued to a similar degree when added after 2 d of NGF deprivation (Fig. 4 B). Thus, although oligomycin is toxic to NGF-maintained sympathetic neurons, it protects NGF-deprived, BAF-saved cells. These findings strongly suggest that activity of the F$_0$F$_1$ ATPase contributes to caspase-independent death of sympathetic neurons.

**Electron transport maintains ΔΨ$_m$ after cytochrome c release**

Inhibition of F$_0$F$_1$ does not decrease ATP levels in cells that have released cytochrome c (Fig. 2), arguing that it is not generating ATP at the expense of the mitochondrial proton gradient. However, the F$_0$F$_1$ ATPase inhibitor, oligomycin, prevents caspase-independent death, suggesting that the activity of this enzyme complex has a role in the death of these cells. As its name suggests, the F$_0$F$_1$ ATPase can hydrolyze ATP to transport protons actively against the electrochemical gradient. In this mode of operation, F$_0$F$_1$ activity would contribute to ΔΨ$_m$, which is maintained, albeit to a lesser degree, in NGF-deprived, BAF-saved neurons after cytochrome c has been released (Chang and Johnson, 2002). Because cytochrome c mediates electron transport from complexes III to IV, the loss of mitochondrial cytochrome c would be expected to impair electron transport chain–mediated maintenance of ΔΨ$_m$. To determine the mechanism by which caspase inhibitor–saved cells maintain ΔΨ$_m$, we examined the effect of inhibiting either electron transport or F$_0$F$_1$ on ΔΨ$_m$ in NGF-deprived, BAF-saved neurons by using 5,5’6,6’-tetraethylbenzimidazolylcarbonyl cyanide iodide (JC-1), a cell-permeable, ΔΨ$_m$-sensitive dye (Smiley et al., 1991). JC-1 exists as a fluorescent monomer, but reversibly forms aggregates, which have different spectral properties than the monomeric form, in the matrix of polarized mitochondria (Nicholls and Ward, 2000). Sympathetic neurons that were either maintained in NGF or deprived of NGF in the presence of BAF for 3 d were loaded with JC-1 and treated with the 2 μM rotenone and 2 μM antimycin A, to block electron transport at sites I and II, respectively, and/or 5 μg/ml oligomycin, to block both forward and reverse operation of F$_0$F$_1$. The effects of these treatments on ΔΨ$_m$ were determined by comparing the JC-1 ratios obtained immediately before and 15 min after drug or vehicle addition. Exposure to the protonophore, carbonyl cyanide m-chlorophenylhydrazone (CCCP), decreased the JC-1 ratio in NGF-maintained cells, reflecting complete mitochondrial depolarization (Fig. 5). Oligomycin had very little effect on ΔΨ$_m$ in NGF-maintained neurons after oligomycin treatment. Treatment of NGF-maintained neurons with rotenone and antimycin A caused complete mitochondrial depolarization, demonstrating that electron transport is responsible for maintenance of ΔΨ$_m$ in NGF-maintained neurons. Inhibition of electron transport largely, but not completely, depolarized mitochondria in NGF-deprived, BAF-saved neurons because the degree of mitochondrial depolarization achieved by treatment with antimycin and rotenone was less than that with CCCP. This electron transport inhibitor–insensitive contribution to ΔΨ$_m$ persisted for at least 45 min after drug addition, arguing against the possibility that inhibition of electron transport dissipates ΔΨ$_m$ more slowly in NGF-deprived, BAF-saved cells (unpublished data). A combination of rotenone, antimycin A, and oligomycin completely depolarized mitochondria in NGF-deprived, BAF-saved cells, suggesting that the residual ΔΨ$_m$ after inhibition of electron transport was maintained by reversal of F$_0$F$_1$ and hydrolysis of ATP (Fig. 5). Thus, although reversal of F$_0$F$_1$ has a minor function in the maintenance of ΔΨ$_m$ in NGF-deprived, BAF-saved, but not NGF-maintained, neurons, electron transport is the primary mechanism by which ΔΨ$_m$ is maintained before and after cytochrome c release in sympathetic neurons.

**CsA attenuates loss of mitochondrial proteins**

As seen in Fig. 2, CsA did not preserve the ability of mitochondria in NGF-deprived, BAF-saved neurons to generate ATP. Therefore, to investigate alternate mechanisms by which PTP inhibition by CsA might block Commitment 2, we asked whether CsA attenuated the loss of mitochondria observed in caspase inhibitor–saved neurons that may correlate with commitment-to-die (Xue et al., 2001). Western blots (Fig. 6 A) demonstrate that the levels of two mitochondrial proteins, voltage-dependent anion channel (VDAC) and cytochrome oxidase subunit IV (COX IV) decreased over time in NGF-deprived, BAF-saved cells, but less dra-
between BAF and BAF of three independent experiments. Asterisks denote that differences to the amount of tubulin in each sample. Data shown are mean ± SD of three independent experiments. Western blot. Blots were simultaneously probed with antibodies directed against tubulin, VDAC, and cytoCOX IV. (B) Levels of mitochondrial proteins. Mitochondrial proteins in NGF deprivation in the presence of BAF or BAF + 10 μM CsA were examined by Western blot. Blots were simultaneously probed with antibodies directed against tubulin, VDAC, and cytoCOX IV. (B) Levels of VDAC and COX IV after 9 d of NGF deprivation were normalized to the amount of tubulin in each sample. Data shown are mean ± SD of three independent experiments. Asterisks denote that differences between BAF and BAF + CsA are statistically significant (P < 0.05).

Control neurons maintained in NGF (Fig. 7 A) had large, round nuclei (N) with prominent nucleoli. The cytosol contained many mitochondria (Fig. 7 B, m) and occasional electron-dense, membrane-limited late autophagic vesicles. Neurons deprived of NGF in the presence of BAF for 10 d displayed extensive atrophy (Fig. 7 C), as expected from their light microscopic appearance. The most prominent change was the appearance of numerous electron-dense bodies in the cytosol (Fig. 7 D, black arrows), which may represent lipid droplets (Martin et al., 1988) or autolysosomes that have engulfed a large amount of lipid membranes or other electron-dense material (Xue et al., 1999). Similar structures were also occasionally seen in NGF-maintained neurons, although these were often smaller and had limiting membranes (Fig. 7 B, white arrows). These structures were present at a much greater frequency in sections of NGF-deprived, BAF-saved neurons (0.51 ± 0.08 per μm² of cytosol, ± SEM, in 11 sections from different neurons), than in NGF-maintained neurons (0.07 ± 0.04 per μm², n = 6). Cells deprived of NGF in the presence of BAF and CsA also displayed cytoplasmic atrophy and convolution of the nuclear membrane (Fig. 7 E). However, two key differences were observed between the appearance of these cells and those treated with BAF alone. First, these electron-dense bodies were much less abundant in cells treated with BAF and CsA (0.15 ± 0.04 per μm², n = 11) than in neurons saved with BAF only (Fig. 7 E). Second, abundant multilamellar vesicles were present throughout the cytosol of these cells (Fig. 7, E and F, black arrowheads). These multilamellar vesicles were abundant throughout multiple sections of NGF-deprived neurons treated with BAF and CsA (0.59 ± 0.11 per μm², n = 11), but rarely present in NGF-deprived, BAF-saved cells (0.09 ± 0.03 per μm², n = 11) and completely absent in NGF-maintained neurons (n = 6). Although the precise nature of these structures is uncertain, they resemble autolysosomes seen in NGF-deprived, caspase inhibitor–saved neurons (Xue et al., 1999). When normalized to surface area, NGF-deprived, BAF-saved cells with (0.22 ± 0.08, ± SEM) or without CsA treatment (0.17 ± 0.08) had slightly fewer mitochondria than NGF-maintained neurons (0.33 ± 0.09), but this difference was not statistically significant. In NGF-deprived, BAF-saved cells, there was a trend toward more mitochondria in CsA-treated cells, but this difference was not statistically significant.

Discussion

Here, we examined changes in metabolism in sympathetic neurons prevented from completing apoptosis by inhibition of caspases and how these changes influenced caspase-independent death. We present three major findings. First, NGF-deprived, BAF-saved neurons rely on glycolysis for ATP generation and for survival. Second, activity of F₀F₁ contributes to caspase-independent death, but has only a minor role in the maintenance of ΔΨm, which is maintained primarily by electron transport. Finally, CsA preserves mitochondrial proteins, suggesting that PTP opening has a function in regulating the degradation of mitochondria after cytochrome c release.
NGF-deprived, BAF-saved cells rely largely on glycolysis to generate ATP

NGF-maintained neurons are remarkably resistant to glucose deprivation (Fig. 3 B), despite the fact that acute inhibition of glycolysis decreased ATP levels within these cells (Fig. 2). The medium used in this experiment lacks pyruvate, but contains nine amino acids that can serve as carbon sources for intermediates in the citric acid cycle. This argues that in sympathetic neurons, the citric acid cycle and oxidative phosphorylation are able to generate sufficient ATP for survival. However, NGF-deprived, BAF-saved sympathetic neurons rely on glycolysis to generate ATP (Fig. 2) and have a marked sensitivity to glucose deprivation. In these cells, blocking glycolysis for 2 h decreased ATP levels by over 70%. This increased reliance on glycolysis renders NGF-deprived, BAF-saved neurons vulnerable to glucose deprivation, an insult to which NGF-maintained neurons are completely insensitive. Thus, unlike in NGF-maintained cells, mitochondria within NGF-deprived, BAF-saved cells cannot generate ATP to compensate for the loss of glycolysis, arguing that the mitochondrial hit has compromised certain mitochondrial functions.

Given the striking effect of glucose deprivation on ATP levels, that even a small proportion of NGF-deprived, BAF-saved cells survive prolonged glucose deprivation is surprising. These surviving cells may be a subpopulation of cells that are able to maintain oxidative phosphorylation. That no change in total ATP was detected after treatment of NGF-deprived, BAF-saved cells with oligomycin is likely a reflection of the small size of this subpopulation.

This increased sensitivity to glucose deprivation suggests that insufficient ability to generate ATP underlies caspase-independent death of NGF-deprived sympathetic neurons. In fact, a decrease in glucose transport occurs soon after removal of NGF from sympathetic neurons (Deckwerth and Johnson, 1993). However, increasing the concentration of glucose to 25 mM had no effect on Commitment 2, suggesting that availability of glucose in the culture medium was not the factor that limited survival.

Interestingly, oxidative phosphorylation continues to generate ATP in UV-irradiated HeLa cells that have released cytochrome c from their mitochondria (Waterhouse et al., 2001). In this model system, although mitochondrial cytochrome c is lost, ATP levels are initially maintained but decrease ~10 h after cytochrome c release (Waterhouse et al., 2001). Because caspase-independent death in our model occurred with a much slower time course, the time points used in our experiments might not have detected a similar short-term phenomenon. Alternatively, these different findings could reflect the cell type– and stimulus-specific nature of
caspase-independent events, such as degradation of cytosolic cytochrome c or its equilibration between the cytosol and mitochondrial intermembrane space.

**F₀F₁ ATPase does not contribute to ATP generation after cytochrome c release**

Oligomycin protected NGF-deprived, BAF-saved cells (Fig. 4 B), suggesting that activity of F₀F₁ contributes to caspase-independent death. Although we cannot rule out the possibility that oligomycin may have additional activities, it is striking that the toxicity was selective for NGF-maintained cells. Inhibition of oxidative phosphorylation did not decrease the amount of ATP within NGF-deprived, BAF-saved sympathetic neurons (Fig. 2), arguing that F₀F₁ activity does not generate ATP in these cells. Therefore, reverse operation of F₀F₁ to hydrolyze ATP may underlie its role in caspase-independent death. Consistent with this mode of operation, oligomycin dissipated the antimycin, rotenone-insensitive portion of ∆Ψᵐ in NGF-deprived, BAF-saved cells (Fig. 5).

Inhibition of complexes I and II was sufficient to disrupt the majority of ∆Ψᵐ in NGF-deprived, BAF-saved sympathetic neurons (Fig. 5), suggesting that electron transport is the primary mechanism by which ∆Ψᵐ is maintained in cells that have released cytochrome c. How does electron transport continue after the loss of mitochondrial cytochrome c, which is a required component of the electron transport chain? At least two conceivable mechanisms exist. First, enough residual cytochrome c may exist in the intermembrane space after it is “released” from the mitochondrion to continue electron transport. In apoptotic HeLa cells saved with a caspase inhibitor, cytochrome c equilibrates throughout the cytosol and intermembrane space to a concentration sufficient to mediate electron transport and maintenance of ∆Ψᵐ (Waterhouse et al., 2001). However, in sympathetic neurons, cytochrome c appears to be rapidly degraded in the cytosol after its release from the mitochondria, as determined by immunocytochemistry and Western blot of subcellular fractions of cell lysates (Putcha et al., 2000). A second possibility is that an alternate electron carrier substitutes for cytochrome c. This electron carrier could act as an intermediate electron carrier to recapitulate the electron transport chain, or as a terminal electron recipient to form a truncated electron transport chain. Although the nature of this hypothetical electron carrier is not known, precursors to reactive oxygen species (ROS) can act as terminal electron acceptors (Cai and Jones, 1998), allowing continued electron transport and generation of a proton gradient by complexes I and III. ROS, generated by abnormal electron transport, could be directly damaging to caspase inhibitor–saved neurons. The involvement of oxidative damage in caspase-independent death is a topic of current investigation.

Oligomycin collapsed the residual ∆Ψᵐ remaining after inhibition of electron transport (Fig. 5). However, oligomycin alone did not increase the JC-1 ratio in NGF-maintained neurons (Fig. 5). Under conditions of oxidative phosphorylation, oligomycin would normally be expected to cause a slight mitochondrial hyperpolarization because it prevents dissipation of the proton gradient by inhibiting F₀F₁ (Scott and Nicholls, 1980). We did not observe this in

![Diagram of mitochondrial events](image-url)

**Figure 8. Proposed events that account for the protective effect of oligomycin and CsA in neurons subsequent to cytochrome c release by the "mitochondrial hit."** In NGF-maintained sympathetic neurons (A), electron transport generates ∆Ψᵐ, which is used by the ∆Ψᵐ ATPase to generate ATP. In NGF-deprived, BAF-saved cells (B), mitochondrial cytochrome c (Cc) is lost by permeabilization of the outer mitochondrial membrane, possibly by a channel that includes the pro apoptotic Bcl-2 family member BAX. Despite this, electron transport continues, at least through complexes I and III, contributing to ∆Ψᵐ. Oxidation of electron transport intermediates could be mediated by residual mitochondrial cytochrome c, or by the generation of reactive oxygen species (ROS), which could themselves be detrimental. Reverse operation of F₀F₁ also contributes to ∆Ψᵐ by hydrolyzing ATP. The importance of the permeability transition pore (PTP), which is composed of the voltage-dependent anion channel (VDAC), adenine nucleotide translocase (ANT), and cyclophilin D (CyD), is evidenced by the ability of CsA to inhibit caspase-independent death (Chang and Johnson, 2002; Fig. 4). Although precisely how PTP opening contributes to caspase-independent cell death is not known, it is possible, but purely speculative, that the opening of the PTP could allow the F₀F₁ to hydrolyze cytosolic ATP generated by glycolysis, on which the cell depends for survival. Although this model can account for inhibition of caspase-independent death by both oligomycin and CsA, it suggests that mechanisms to equilibrate adenine nucleotides are compromised in NGF-deprived, BAF-saved cells.
our experiments on NGF-maintained cells, even though we were able to detect mitochondrial hyperpolarization caused by exposure to nigericin, which permeabilizes the plasma membrane to protons and increases the proton gradient across the mitochondrial inner membrane, in both NGF-maintained and NGF-deprived, BAF-saved neurons (unpublished data). The failure to observe the predicted effect of oligomycin alone may be because of insensitivity in the upper ranges of this assay, because one cannot rigorously demonstrate the linearity of this assay in this setting. However, clear qualitative differences occur between the nature of ΔΨm in NGF-maintained and NGF-deprived, BAF-saved neurons.

If reverse operation of F0F1 contributes to caspase-independent death, how does it do so? Circumstantial evidence suggests that cytochrome c release, and the events leading up to it, have effects on mitochondria that are detrimental to cell survival, even in models of cell death in which cytochrome c release occurs by selective permeabilization of the outer mitochondrial membrane (Von Ahsen et al., 2000). The F0F1 ATPase generates ATP under normal conditions, but hydrolyzes ATP in an attempt to maintain mitochondrial membrane potential (ΔΨm) after cytochrome c release in apoptotic GT1–7 cells (Rego et al., 2001). The continued ATP hydrolysis by F0F1 is likely to be detrimental to a cell because it would deplete cellular ATP.

The polarity of the F0F1 ATPase in the inner mitochondrial membrane dictates that it can only hydrolyze ATP within the mitochondrial matrix. Under normal conditions, ATP and ADP are freely exchanged between the mitochondrial matrix and the cytosol via the adenine nucleotide translocase (ANT). However, this may not hold true during cell death, as ANT function is compromised in lymphocytes undergoing growth factor deprivation–induced apoptosis (Vander Heiden et al., 1999). If adenine nucleotide equilibration is compromised in NGF-deprived, BAF-saved sympathetic neurons, it is possible that opening of the PTP renews the pool of ATP within the mitochondrial matrix by providing equilibration of ATP levels between the cytosol and the matrix, schematized in Fig. 8. In such a scenario, PTP opening allows the F0F1 ATPase access to ATP that has been generated in the cytosol by glycolysis, which is required by these cells for survival (Fig. 3 B). Thus, inhibiting PTP opening with CsA or directly inhibiting ATP hydrolysis by reverse operation of F0F1 with oligomycin limits the deleterious effects of the mitochondrial hit by preserving glycolytic ATP. Consistent with this hypothesis, CsA and oligomycin inhibited Commitment 2 to virtually the same degree (Fig. 4 B). Although purely speculative at this point, because it rests on the supposition that adenine nucleotide transport is altered in NGF-deprived, BAF-saved cells, we favor this model because it accounts for the similarity in protection against caspase-independent death of both oligomycin and CsA.

If reversal of F0F1 activity is an important event in caspase-independent death, why is the effect of oligomycin on ΔΨm so small (Fig. 6)? At least two possible explanations exist. First, reverse operation of F0F1 may occur at a slow, but sustained, rate in NGF-deprived sympathetic neurons. In this scenario, oligomycin would have only a minor effect on ΔΨm over the course of minutes, but a great effect on survival when applied for over a week. Consistent with this, the rate of reverse activity of F0F1 in anoxic skeletal muscle is lower than predicted, suggesting the existence of mechanisms to limit ATP hydrolysis by F0F1 (St. Pierre et al., 2000). Alternatively, F0F1 may reverse in only a small proportion of cells at any given time. F0F1 reversal may begin asynchronously throughout the population and followed by loss of ΔΨm, because reverse activity of F0F1 contributes to caspase-independent death. Thus, the small effect of oligomycin on ΔΨm, as measured on a population basis by this JC-1 assay, may be entirely accounted for by this small, transient subpopulation of cells. The techniques used in these studies cannot differentiate between these two possibilities.

**CsA attenuates NGF deprivation–induced loss of mitochondrial proteins**

Although CsA does not alter protein synthesis or prevent somal atrophy of NGF-deprived, BAF-saved cells (Chang and Johnson, 2002), CsA increases the amount of mitochondrial protein within these cells (Fig. 6 B), arguing that CsA directly or indirectly prevents the degradation of mitochondria. In other model systems, including a different paradigm of NGF deprivation–induced death of sympathetic neurons, which differs in some important respects with our system (Fletcher et al., 2000), selective mitochondrial elimination is mediated by autophagy (Tolkovsky et al., 2002). The striking difference in the appearance of cytoplasmic vesicles in NGF-deprived, BAF-saved cells treated with CsA further supports the conclusions that mitochondria within apoptotic cells saved with a caspase inhibitor are eliminated by autophagy. That CsA interferes with this process is not unprecedented because inhibition of PTP opening by CsA prevents autophagy of mitochondria in serum-deprived, glucagon-treated hepatocytes (Elmore et al., 2001). Thus, the PTP opening may be the trigger for the removal of mitochondria in NGF-deprived, BAF-saved neurons, marking them for degradation. Mitochondria are likely to be critical for the ability of a cell to survive after trophic factor readdition because oxidative phosphorylation is required for survival in the presence of NGF (Fig. 4 B). However, the loss of mitochondria in caspase inhibitor–saved cells may not cause caspase-independent death because mitochondria removed from these cells may be dysfunctional or damaged. Mitochondrial damage or dysfunction may have a role in regulating selective mitochondrial elimination (James et al., 1996; Elmore et al., 2001). Thus, the selective elimination of mitochondria may be a result, but not a cause, of impaired mitochondrial function. In support of this, CsA protects NGF-deprived, BAF-saved neurons from caspase-independent death (Chang and Johnson, 2002) and attenuated the loss of mitochondrial proteins (Fig. 7), but did not preserve oxidative phosphorylation within these neurons (Fig. 2).

Because degradation of intracellular organelles liberates free amino acids, mitochondrial elimination may represent an attempt to improve the energetic status of caspase inhibitor–saved cells. Although not producing ATP, these “damaged” mitochondria may be maintaining other critical func-
tions in the short term within these cells. The long-term effects of this effort, however, are deleterious because the potential short-term benefits jeopardize the ability of the cell to survive the ongoing, death-inducing stimulus and to recover after the stimulus is terminated.

Implications for postmitochondrial regulation of apoptosis

Much attention has been given to cytochrome c release as a critical point of regulation of apoptosis. However, clearly, mechanisms exist to regulate apoptotic machinery at points distal to cytochrome c release, such as inhibitors of apoptosis (Deveraux and Reed, 1999) and heat shock proteins (Beere et al., 2000; Bruey et al., 2000; Saleh et al., 2000). Regardless of the method by which caspase activation is prevented, caspase-independent sequelae of activating the cell death pathway, such as those examined in this work, will cause the eventual demise of these cells. Like caspase inhibitors, therapeutic use of inhibitors of apoptosis and heat shock proteins will be limited by the ability of the cell to survive these caspase-independent events. Thus, identifying the specific events that contribute to caspase-independent cell death may enhance our ability to exploit postmitochondrial strategies to prevent pathological cell death.

Materials and methods

Unless otherwise noted, all reagents were obtained from Sigma-Aldrich. Timed-pregnant Sprague-Dawley rats were obtained from Harlan Sprague-Dawley.

Cell culture

Sympathetic neurons from postnatal day 0–1 rats were maintained in a medium containing 50 ng/ml NGF (AM50) as described previously (Deshmukh et al., 1996). Cells were deprived of NGF by washing cells in a medium lacking NGF (AM0), followed by culture in AM0-containing neutralizing antibody to NGF. Commitment-to-die was measured by determining the proportion of NGF-deprived neurons that were rescued by NGF readdition. At the time of the rescue, cultures were washed extensively to remove residual anti-NGF and maintained in AM50. After 7 d, cells were washed and fixed in 4% PFA for at least 12 h at 4°C. Cultures were stained with 0.05% Toluidine Blue in TBS (10 mM Tris and 0.9% NaCl, pH 7.6) and counted by using an inverted microscope (Eclipse TE300; Nikon) without knowledge of treatment group. All values are represented as a percentage of the mean number of cells in NGF-maintained sister cultures.

To determine the effect of altering glucose concentration on survival, cultures on AM50 and AM0 were made with glucose-free MEM (W洗ingtton University Tissue Support Center) and dialyzed FBS. In some cases, glucose was added to bring the final concentration either to 5 mM, the concentration of glucose in standard MEM, or 25 mM. These alterations in formulation of the medium did not affect cell viability (Fig. 3 B, respectively to remove residual anti-NGF and maintained in AM50. After 7 d, cells were washed and fixed in 4% PFA for at least 12 h at 4°C. Cultures were stained with 0.05% Toluidine Blue in TBS (10 mM Tris and 0.9% NaCl, pH 7.6) and counted by using an inverted microscope (Eclipse TE300; Nikon) without knowledge of treatment group. All values are represented as a percentage of the mean number of cells in NGF-maintained sister cultures.

Determination of ATP

ATP was measured by using a luciferase-based assay (Bioluminescent Somatic cell assay kit; Sigma-Aldrich) according to the manufacturer’s instructions. All manipulations were performed on ice with ice-cold solutions. Sister cultures of 10,000 neurons per well in 24-well plates were washed once with PBS, lysed in a 1:1 dilution of the supplied releasing agent in water, and immediately frozen at –70°C after 0, 3, 6, or 9 d of treatment. After all samples from an individual experiment were collected, luciferase reagent was added to an aliquot of each lysate, and the amount of ATP was determined with a microplate luminometer (model TR717; Applied Biosystems). At each time point, the amount of ATP in sister cultures plated in anti-NGF was subtracted from all values to determine the amount of neuronal ATP. All values are expressed as a percentage of the average amount of ATP in cultures at time 0 for each experiment. By using known amounts of ATP, a standard curve was generated for each experiment to ensure that all values were in the linear range of the assay. In some experiments, an aliquot of each lysate was used for determination of total protein with the BCA method (Pierce Chemical Co.). The amount of neuronal protein in each sample was determined by subtracting the protein in sister cultures plated in anti-NGF.

To determine the source of intracellular ATP, cells were maintained in NGF or deprived of NGF in the presence of BAF for 3 d and treated with a standard medium, glucose-free medium with 5 mM 2-deoxyglucose and 1 mM pyruvate, or a standard medium with 5 μg/ml oligomycin for 2 h.

Determination of ∆Ψ

Sympathetic neuronal cultures were grown in 96-well, opaque-walled, clear-bottom plates (Corning Costar). NGF-maintained cultures and those that had been deprived of NGF in the presence of BAF for 3 d were washed once and loaded with 3.3 μM JC-1 (Molecular Probes) in PBS with 1 g/liter glucose for 30 min at 37°C after which the cells were washed twice. JC-1 fluorescence was measured with a fluorescent plate reader (Fluoroskan II; Titertek) by using excitation/emission pairings of 485/538 nm, corresponding to the monomeric form of JC-1, and 544/590 nm, corresponding to the aggregated form of JC-1. The relative ∆Ψ was determined by calculating the ratio of the aggregated form of JC-1 to its monomeric form. Baseline readings were taken before vehicle or drug addition. The relative change in ∆Ψ in response to treatment was determined by normalizing the JC-1 ratio of each well 15 min after vehicle or drug addition to the baseline ratio for each well. Because JC-1 is a relatively slow equilibrating dye, changes in the JC-1 ratio were observed over the first 10–12 min after drug addition, but then remained stable over the next 30 min. Vehicle treatment in either NGF-maintained or -deprived, BAF-saved neurons slightly increased the JC-1 ratio over this period likely caused by diffusion of the monomeric form out of the cells. At the end of each experiment, CCCP was added to each well and a reading was taken after 15 min. In all cases, the JC-1 ratio decreased to a value similar to that of cells initially treated with CCCP, suggesting that this reflects maximal mitochondrial depolarization. In some experiments, sister cultures that were deprived of NGF in the absence of BAF were rescued with NGF to monitor cytochrome c release (Patcha et al., 1999). In all cases, <5% of cells were rescued, demonstrating that at least 95% of the cells had released cytochrome c.

Western blotting

At the appropriate times, cultures were lysed in lysis buffer containing 100 mM Tris, pH 6.8, 4% SDS, 20% glycerol, and 5% β-mercaptoethanol. Proteins were resolved on Novex Tris-glycine gels (Invitrogen) and transferred to PVDF membranes (Millipore). Blots were blocked with 5% dry milk in TBS-T (10 mM Tris, pH 7.5, 100 mM NaCl, and 0.1% Tween 20) before incubating with a mixture of mouse antiblutinin, mouse anti-VDAC, and mouse anti–COX IV (Molecular Probes) in TBS-T with 5% milk overnight at 4°C. After washing, blots were incubated with HRP-conjugated anti–mouse secondary and visualized with SuperSignal Pico (Pierce Chemical Co.). The intensity of bands was determined with UnScan-It (Silk Scientific).

Electron microscopy

 Cultures were grown on collagen-coated Permanox LabTek chamber slides (Nalge Nunc). After treatments, cells were fixed for 4 h in 3% glutaraldehyde in 100 mM phosphate buffer, pH 7.3, containing 0.45 mM Ca2+. Cultures were fixed after in buffered OsO4, dehydrated in graded alcohols, and embedded in Epon. Ultrathin sections were cut and examined with an electron microscope (model 1200; JEOL). For the purposes of quantification, photomicrographs taken at 10,000× were examined without knowledge of the treatment group. Cytoplasmic structures with or without limiting membranes that were at least half filled with electron-dense material were considered to be “electron-dense bodies.” Membrane-limited, multilamellar structures in the cytosol that were less than half filled with electron-dense material were counted as “multilamellar whorls.”

We thank Mohanish Deshmukh and Patricia Osborne for helpful discussion and critical review of this paper. This work was supported by the National Institutes of Health grants AG 12957 and NS 38651 (to E.M. Johnson).

Submitted: 19 February 2003
Revised: 5 June 2003
Accepted: 5 June 2003
Oligomycin inhibits Commitment 2 in rat sympathetic neurons | Chang et al. 255

References


