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BAX insertion, oligomerization, and outer membrane permeabilization in brain mitochondria: role of permeability transition and SH-redox regulation

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Abstract

BAX cooperates with truncated BID (tBID) and Ca²⁺ in permeabilizing the outer mitochondrial membrane (OMM) and releasing mitochondrial apoptogenic proteins. The mechanisms of this cooperation are still unclear. Here we show that in isolated brain mitochondria, recombinant BAX readily self-integrates/oligomerizes in the OMM but produces only a minuscule release of cytochrome *c*, indicating that BAX insertion/oligomerization in the OMM does not always lead to massive OMM permeabilization. Ca²⁺ in a mitochondrial permeability transition (mPT)-dependent and recombinant tBID in an mPT-independent manner promoted BAX insertion/oligomerization in the OMM and augmented cytochrome *c* release. Neither tBID nor Ca²⁺ induced BAX oligomerization in the solution without mitochondria, suggesting that BAX oligomerization required interaction with the organelles and followed rather than preceded BAX insertion in the OMM. Recombinant Bcl-xL failed to prevent BAX insertion/oligomerization in the OMM but strongly attenuated cytochrome *c* release. On the other hand, a reducing agent, dithiothreitol (DTT), inhibited BAX insertion/oligomerization augmented by tBID or Ca²⁺ and suppressed the BAX-mediated release of cytochrome *c* and Smac/DIABLO but failed to inhibit Ca²⁺-induced swelling. Altogether, these data suggest that in brain mitochondria, BAX insertion/oligomerization can be dissociated from OMM permeabilization and that tBID and Ca²⁺ stimulate BAX insertion/oligomerization and BAX-mediated OMM permeabilization by different mechanisms involving mPT induction and modulation of the SH-redox state.

Keywords

mitochondria; calcium; BAX; BID; Bcl-xL; permeability transition

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1. Introduction

Apoptosis is an omnipresent form of cell death involved in various neurodevelopmental [1] as well as neuropathological processes, including age-related neurodegenerations [2-6], stroke [7-10], and secondary brain injury following mechanical brain trauma [11,12]. The release of mitochondrial apoptogenic factors, a key step in executing of apoptosis [13], occurs due to a concert action of pro-apoptotic proteins such as BID and BAX [14]. Under normal conditions, monomeric BAX and full-length BID are located in the cytosol [15]. Caspase-8 activated by apoptotic stimuli cleaves BID, producing activated (truncated) BID (tBID) [16]. In turn, tBID activates BAX either “directly” [15,17,18] or “indirectly” [19,20] leading to oligomerization of BAX, its insertion into the OMM, and OMM permeabilization culminating in the release of mitochondrial apoptogenic proteins [18,21].

In addition to tBID, elevated Ca^{2+} enhances the ability of BAX to integrate into the lipid membranes and permeabilize them [22-24]. Ca^{2+} also amplifies BAX ability to permeabilize the OMM [25], though the mechanism of such amplification is unknown. Since elevated Ca^{2+} induces the mitochondrial permeability transition (mPT), a phenomenon accompanied by mitochondrial depolarization and remodeling [26], it is possible that the mPT is involved in augmentation of BAX-mediated OMM permeabilization. In cerebellar granule neurons, trophic factor withdrawal in low- K^+ medium resulted in the mPT that triggered BAX translocation to mitochondria and release of Cyt *c* [27]. In line with this, in the model of ischemia/reperfusion heart injury, inhibition of the mPT either with Ru_{360} , an inhibitor of the mitochondrial Ca^{2+} uniporter [28], or with cyclosporin A (CsA), an inhibitor of the mPT [29], precluded BAX insertion in the OMM and OMM permeabilization [30]. Thus, there is evidence suggesting a synergistic relationship between the Ca^{2+} -induced mPT and BAX in OMM permeabilization.

In the present study, we demonstrated that BAX could readily self-integrate and oligomerize in the OMM, but these events were not accompanied by massive Cyt *c* release. We also found that Ca^{2+} in an mPT-dependent and tBID in an mPT-independent manner augmented BAX insertion and oligomerization in the OMM that correlated with the increased OMM permeabilization. Moreover, we showed that the Ca^{2+} - and tBID-stimulated BAX insertion/oligomerization depended on SH-redox state and could be inhibited by a reducing agent, dithiothreitol (DTT). DTT also attenuated BAX-mediated OMM permeabilization stimulated by Ca^{2+} or tBID, revealing an important role of SH-redox regulation in the release of mitochondrial apoptogenic proteins.

2. Materials and Methods

2.1 Recombinant proteins

Full-length human monomeric BAX with a tag of six histidine residues at the N-terminus was expressed in the pBAD plasmid in *Escherichia coli* [31]. Mouse tBID (BID cut with caspase-8 and separated from N-terminal and the caspase) was obtained from full-length BID as described previously [32]. Recombinant Bcl-xL was produced as described previously [17]. Recombinant BAX, tBID, and Bcl-xL were stored in dialysis buffer containing 25 mM HEPES-NaOH, pH 7.5, 0.2 mM dithiothreitol, 30% glycerol (v/v) at -86°C .

2.2 Isolation and purification of brain mitochondria

Mitochondria from the brains of male Sprague-Dawley rats, 200–250 g (Harlan, Indianapolis, IN, USA) were isolated in mannitol-sucrose medium according to an Institutional Animal Care and Use Committee approved protocol and purified on a discontinuous Percoll gradient as

described previously [33]. Mitochondrial protein was measured by the Bradford method [34], using BSA as a standard.

2.3 Measurements of mitochondrial light scattering

Mitochondrial swelling was evaluated in the standard incubation medium at 37°C by monitoring the scattering of light directed on mitochondrial suspension under 90° to the axis of the photodetector at 525 nm in a 0.4-ml cuvette under continuous stirring using a PerkinElmer LS-55 luminescence spectrometer. The standard incubation medium used in these and other experiments contained 125 mM KCl, 10 mM HEPES, pH 7.4, 0.5 mM MgCl₂, 3 mM KH₂PO₄, 10 μM EGTA, 0.1% bovine serum albumin (free from fatty acids), 3 mM glutamate, and 3 mM succinate.

2.4 Transmission electron microscopy

Electron microscopy of isolated brain mitochondria was performed as described previously [35]. Mitochondria were incubated in the standard 125 mM KCl-based medium at 37°C prior to fixation in 2% paraformaldehyde and 2% glutaraldehyde in 0.05 M phosphate buffer in the same incubation medium at room temperature for 15 minutes. Transmission electron microscopy (TEM) images were taken using a Tecnai G12 BioTwin electron microscope (FEI, Hillsboro, OR) equipped with an AMT 2.6×2.6K digital CCD camera.

2.5 Alkali-resistant BAX insertion

The alkali treatment of mitochondria removes loosely attached proteins but leaves proteins inserted into the OMM [18]. We determined the alkali-resistant fraction of BAX inserted into the OMM using the earlier described method [36]. Briefly, mitochondria treated with BAX (50 or 150 nM) at 37°C for 30 minutes were pelleted at 15,800 g for 5 minutes, and supernatant was used for the Cyt *c* release measurements. Mitochondrial pellets were re-suspended in 0.2 ml of 0.1 M Na₂CO₃, pH 11.5, then incubated for 30 minutes on ice. Samples were centrifuged for 30 minutes at 100,000 g in an Optima L-100K Beckman ultracentrifuge. The pellets were solubilized using 1% 3-[(3-cholamidopropyl) dimethylammonio]-1-propanesulfonate (CHAPS) or 1% [octylphenoxy] polyethoxyethanol (Nonidet P-40, Amresco, Solon, OH) and analyzed by western blotting against BAX and cytochrome oxidase subunit IV (COX IV, loading control).

2.6 Immunoblotting

The release of Cyt *c* and Smac/DIABLO from isolated brain mitochondria was assessed in supernatants obtained through incubation of mitochondria in the standard 125 mM KCl-based incubation medium with or without additions for 30 minutes at 37 °C. For SDS-PAGE, we used 4-12% Bis-Tris gels (Invitrogen, Carlsbad, CA). Western blotting was performed as previously described [37]. In some experiments, alamethicin (30μg/ml) was used to produce the maximal Cyt *c* release. Mitochondrial cytochrome oxidase subunit IV (COX IV) was used as a loading control for the pellet samples. COX IV was detected with mouse monoclonal anti-COX IV antibody, dilution 1:5000 (Invitrogen, Carlsbad, CA). Following SDS-PAGE, proteins were transferred to Hybond™-ECL™ nitrocellulose membrane (Amersham Biosciences, Piscataway, NJ), and blots were incubated with mouse anti-cytochrome *c* antibody (7H8.2C12, PharMingen, San Diego, CA) at 1:3000 dilution or with rabbit anti-Smac/DIABLO antibody (Alexis Biochemicals, San Diego, CA) at 1:1500 dilution for an hour at room temperature in 5% non-fat milk, phosphate-buffered saline, pH 7.2, and 0.15% Triton X-100. Prior to analysis of Smac/DIABLO release, the supernatants were concentrated threefold in the Microcon YM-10 filtering devices (Millipore Corporation, Bedford, MA). In the alkali-resistant BAX insertion experiments, BAX was detected by western blotting with rabbit polyclonal anti-BAX antibody (Upstate, Lake Placid, NY). Recently, it was shown that oxidation of BAX's cysteines

favored formation of disulfide bridges and BAX oligomerization [38,39], so it is possible that formation of disulfide bridges might contribute to BAX oligomerization in our experiments. Correspondingly, to prevent disruption of disulfide bridges and disassembly of BAX oligomers, SDS-PAGE was performed under non-reducing conditions. Anti-BAX antibody was used at 1:2000 dilution for an hour at room temperature in 5% BSA (Jackson ImmunoResearch Laboratories, West Grove, PA), phosphate-buffered saline, pH 7.2, and 0.15% Triton X-100. Blots were developed using goat anti-rabbit or anti-mouse IgG (1:20000) coupled to horseradish peroxidase (Jackson ImmunoResearch Laboratories) and Supersignal West chemiluminescent reagents (Pierce, Rockford, IL). Molecular weight marker SeeBlue® Plus 2 Standards (5 µl), (Invitrogen, Carlsbad, CA) were used to determine the molecular weights of the bands. NIH ImageJ 1.40g software (<http://rsb.info.nih.gov/ij/>) was used to quantify band densities. All immunoblots are representative of at least three independent experiments.

2.7 Analytical gel-filtration

Analytical gel-filtration was carried out on a Superdex 200 HR 10/30 column using FPLC. Prior to injecting into the column, BAX (500 nM) was pre-incubated at 4°C for 24 hours in the solution containing 125 mM KCl, 10 mM HEPES, pH 7.4, and 1% CHAPS. The same solution was used to equilibrate the column. After injecting the column with 150 µl sample, fractions of 0.4 ml were collected and protein was concentrated with trichloroacetic acid/acetone precipitation prior to analysis by western blotting. The column was calibrated using gel-filtration protein standards. Protein standards were Blue Dextran (2000 kDa), ferritin (440 kDa), catalase (232 kDa), albumin (66 kDa), chymotrypsinogen A (25 kDa) (all from Sigma).

2.8 Protein cross-linking

Cross-linkers were dissolved in DMSO right before the experiment. Ethylene glycol bis (succinimidyl succinate) (EGS, 0.2 mM), disuccinimidyl suberate (DSS, 0.2 mM), and bismaleimidohexane (BMH, 0.5 mM) (all from Pierce) were used. The cross-linkers were added to the standard incubation medium supplemented with 50 nM BAX for 15 minutes at 37°C. EGS and DSS were quenched by 20 mM Tris-HCl, pH 7.5, incubating with rocking for 30 minutes at room temperature. BMH was quenched by 50 mM dithiothreitol (DTT) incubating with rocking for 30 minutes at room temperature. Then, non-reducing SDS-PAGE and western blotting were performed.

2.9 Statistics

Statistical analyses of experimental data consisted of a one-way analysis of variance (ANOVA) followed by Bonferroni's *post hoc* test (GraphPad Prism, version 4.0, GraphPad Software, San Diego, CA). The data represent the mean ± SEM of at least three independent experiments.

3. Results

3.1 BAX insertion and oligomerization in the outer mitochondrial membrane

The release of mitochondrial apoptogenic proteins depends on BAX insertion/oligomerization in the OMM [17,18]. How Ca^{2+} and tBID influence BAX insertion and oligomerization in the OMM of brain mitochondria is unknown. In our study, we took advantage of isolated purified brain mitochondria as a well-defined, cell-free model system that allows direct access to the OMM and precise control of the experimental conditions. Importantly, the OMM represents a natural target for pro-apoptotic proteins like BAX and tBID and contains all necessary components involved to the release of mitochondrial apoptogenic proteins. Thus, isolated brain mitochondria represent a powerful experimental model perfectly suited for detailed analysis of BAX insertion and oligomerization in the OMM and OMM permeabilization.

The recombinant BAX used in our study was predominantly monomeric with small amount of dimers (Fig. 1). Neither Ca^{2+} nor tBID triggered BAX oligomerization in the solution prior to adding mitochondria. Thus, BAX oligomerization required interaction of BAX with the OMM and, hence, most likely followed rather than preceded BAX insertion into the OMM. In the untreated mitochondria, the amount of endogenous BAX was below the detection limit of western blotting (Fig. 2a, lane 1). Incubation of mitochondria with BAX alone produced alkali-resistant BAX insertion and oligomerization in the OMM (Fig. 2a, lane 2), indicating that BAX can self-integrate and self-oligomerize in the OMM producing various BAX oligomers. Both Ca^{2+} and tBID significantly increased the amount of inserted/oligomerized BAX (Fig. 2a-c). In these experiments, we used previously established concentration of Ca^{2+} that produced distinct swelling of isolated brain mitochondria but did not cause significant Cyt *c* release in the standard, 125 mM KCl-based incubation medium [33]. In some western blotting experiments, the key samples were run in duplicate to demonstrate reproducibility. Figure 2b shows statistical analysis of BAX insertion based on densitometry data obtained with individual BAX bands (1×BAX-5×BAX) shown in Figure 2a. Thus, BAX could self-integrate/oligomerize in the OMM and both Ca^{2+} and tBID stimulated these processes. Importantly, we did not use cross-linkers in our experiments. In our hands, cross-linkers ethylene glycol bis (succinimidyl succinate) (EGS), disuccinimidyl suberate (DSS), and bismaleimido-hexane (BMH, all from Pierce) triggered BAX oligomerization in the solution without mitochondria (Supplemental Fig. 1) and therefore were unacceptable. In addition, in these experiments we found that BSA-containing blocking solution was preferable for detecting BAX oligomers than non-fat milk.

We used overnight incubation with 1% CHAPS at 4°C to solubilize mitochondrial pellets after alkali treatment (Fig. 2a). For comparison, we also used 1% Nonidet P-40 (NP-40, Amresco, Solon, OH), another non-ionic detergent, and detected the same major bands corresponding to BAX oligomers (Fig. 2c). Importantly, not all exogenous, recombinant BAX was inserted and oligomerized in the OMM. A fraction of exogenous BAX (~35-40%) remained in the incubation medium in the form of monomers and dimers (Supplemental Fig. 2). Figure 2d shows statistical analysis of BAX insertion based on densitometry data obtained with individual BAX bands (1×BAX-5×BAX) shown in Figure 2c. In the experiments with mitochondrial pellets solubilized with NP-40, we tested the hypothesis that the mPT is involved in Ca^{2+} -stimulated BAX insertion/oligomerization in the OMM. A combination of CsA and ADP, inhibitors of the mPT [40-42], added to mitochondria prior to BAX attenuated BAX insertion and oligomerization stimulated by Ca^{2+} (Fig. 2d, compare lanes 2 and 6 and lane 4). On the other hand, CsA and ADP failed to attenuate tBID-stimulated BAX insertion and oligomerization (Fig. 2d, compare lanes 3 and 5), which is consistent with the insensitivity of tBID plus BAX-induced Cyt *c* release to mPT inhibitors [37].

3.2 Effect of different detergents on BAX quaternary structure

In the experiments with NP-40, the amount of large BAX oligomers (>190 kDa) was much smaller than in the experiments with CHAPS (Fig. 2a,c). This suggested that either NP-40 disassembled the large BAX oligomers (>190 kDa), or they were an artifact produced by interaction of BAX with CHAPS. We tested several non-ionic detergents to evaluate their ability to trigger BAX oligomerization. Previously, Antonsson et al. (2000) reported that octyl glucoside (OG) caused BAX oligomerization [43]. However, in our experiments we did not observe BAX oligomerization with OG. The reason for that is unclear but might be related to the difference in experimental conditions. Our experiments revealed that OG, Triton X-100 (Tr X-100), and NP-40 somewhat increased amounts of BAX dimers and produced small amount of BAX trimers but failed to trigger formation of larger BAX oligomers (Fig. 3a-d). CHAPS, on the other hand, readily oligomerized BAX, producing various forms of BAX oligomers (Fig. 3a-d). The reason other investigators did not observe BAX oligomerization in

the presence of CHAPS is not clear but, possibly, this could be due to difference in experimental conditions used for western blotting. For example, in our hands 5% non-fat milk, which is also used by many investigators as blocking solution in western blotting, significantly hindered detection of BAX oligomers produced by CHAPS or by interaction of BAX with mitochondria. Interestingly, in the experiments with the experiments with CHAPS we observed an increase in the total amount of BAX immunoreactive material over time despite equal protein loading in every lane. The reason for this increase is unclear but it is possible that in these experiments monomeric BAX bands were oversaturated and this could obscure redistribution of BAX from monomeric band to the bands corresponding to BAX oligomers.

To confirm that CHAPS induced BAX oligomerization, we performed analytical gel-filtration of BAX in 1% CHAPS solution (Fig. 3*e,f*). In these experiments, we detected BAX in high molecular weight fractions, indicating formation of large BAX oligomers. Notably, UV absorbance measurements in the eluate revealed huge BAX aggregates with molecular weights up to several megaDa (Fig. 3*e,f*). Thus, both SDS-PAGE and analytical gel-filtration confirmed BAX oligomerization in the solution with CHAPS. Overall, these data suggest that in the experiments with alkali-resistant BAX insertion into the OMM, CHAPS might produce an artifact leading to formation of high-molecular weight BAX oligomers (Fig. 2*a*). On the other hand, these results confirmed that NP-40 did not trigger BAX oligomerization and therefore in the following experiments we used NP-40 to solubilize mitochondria.

3.3 tBID and Ca²⁺ increase BAX-mediated Cyt c release: role of the mPT

In the next experiments, we evaluated whether BAX insertion/oligomerization augmented by tBID and Ca²⁺ correlated with increased OMM permeabilization. We examined Cyt *c* release induced by BAX alone or in combination with tBID or Ca²⁺. Notably, in these experiments, isolated brain mitochondria retained OMM integrity and did not release Cyt *c* spontaneously during incubation in the standard 125 mM KCl-based medium for 30 minutes at 37°C (Fig. 4*a*, lane 1). BAX (50 nM) added alone produced a minuscule Cyt *c* release (Fig. 4*a*, lane 3). Larger BAX concentration (150 nM) resulted in a greater Cyt *c* release (Fig. 4*a*, lane 4) comparable with Cyt *c* release observed in our previous study [37]. Importantly, Ca²⁺ added alone to mitochondria failed to produce substantial Cyt *c* release (Fig. 4*a*, lane 2). Similar observations were reported earlier and were linked to insufficient mitochondrial swelling that was not extensive enough to rupture the OMM [33,44,45]. Nevertheless, Ca²⁺ significantly augmented BAX-mediated Cyt *c* release (Fig. 4*a*, lanes 5 and 6). A combination of 50 nM BAX and 20 nM tBID produced a nearly complete Cyt *c* release (Fig. 4*a*, lane 7). Pre-treatment of mitochondria with CsA plus ADP, inhibitors of the mPT [40-42], significantly diminished Cyt *c* release induced by a combination of BAX and Ca²⁺ (Fig. 4*c*, compare lanes 2 and 3). In these experiments, alamethicin (30 µg/ml) was used as a positive control to produce maximal Cyt *c* release (Fig. 4*c*, lane 4). Thus, our data suggested mPT involvement in the Ca²⁺-induced stimulation of BAX-mediated OMM permeabilization. However, it remained unclear whether Ca²⁺ amplified membrane-permeabilizing activity of BAX, or BAX augmented Ca²⁺-induced mitochondrial swelling resulting in OMM damage and Cyt *c* release.

3.4 BAX does not augment Ca²⁺-induced mitochondrial swelling

To address this question, we evaluated mitochondrial volume changes using 90° light scattering assay [46]. The untreated mitochondria did not swell spontaneously during the course of the experiment (Fig. 5*a*). At the end of the experiments, alamethicin (30 µg/ml) was added to produce maximal swelling (Fig. 5*a-d*). BAX (50 nM) alone failed to induce mitochondrial swelling (Fig. 5*b*). On the other hand, Ca²⁺, an inducer of the mPT [47], produced large-amplitude mitochondrial swelling, and CsA plus ADP completely prevented this swelling (Fig. 5*c*). To address the question whether BAX could increase the Ca²⁺-induced swelling, we incubated mitochondria with BAX (50 nM) and then added Ca²⁺ (1.4 mol/mg protein) (Fig.

5d). To quantify our data, we measured the amplitude of mitochondrial swelling induced by Ca^{2+} as a percentage of maximal alamethicin-induced swelling taken as 100% (Fig. 5c). These experiments showed that BAX did not increase the Ca^{2+} -induced mitochondrial swelling (Fig. 5e). Without BAX, Ca^{2+} (1.4 mol/mg protein) produced $61 \pm 5.6\%$ of maximal swelling versus $63.2 \pm 4.9\%$ with 50 nM BAX (mean \pm SEM, N=3).

Transmission electron microscopy (TEM) corroborated the results obtained with light scattering assay. Following Ca^{2+} application, mitochondrial matrices changed from condensed to predominantly swollen (Fig. 5f,g). BAX (50 nM) failed to affect mitochondrial morphology (Fig. 5h) and did not augment mitochondrial swelling induced by Ca^{2+} (Fig. 5i). In these experiments, we used the morphometric analysis described previously [35,48]. Figure 5j shows the results of morphometric analysis of mitochondria incubated with or without Ca^{2+} and BAX. These data indicated that BAX failed to augment the Ca^{2+} -induced swelling. Therefore, the non-specific damage of the OMM appeared unlikely to be the mechanism of the increased Cyt c release following combined application of BAX and Ca^{2+} .

3.5 Alkali treatment and heating are not necessary for BAX oligomerization in the OMM

High pH or heating of BAX samples above 43-47°C could lead to BAX oligomerization [49-51]. Correspondingly, there was a possibility that BAX oligomerization in our experiments resulted from alkali treatment of mitochondria or heating samples before SDS-PAGE (15 minutes, 70°C). To rule out this possibility, we evaluated BAX oligomerization without alkali-treatment of mitochondria and heating of samples for SDS-PAGE. In these experiments, we detected the same pattern of BAX insertion/oligomerization in the OMM as we observed in our standard experiments with alkali treatment of mitochondria and heating of protein samples (Fig. 6). Interestingly, without alkali treatment, we detected a new band with molecular weight ~80 kDa in solubilized untreated mitochondria (Fig. 6, lane 1). This band was completely eliminated by alkali treatment of mitochondria (Fig. 2a, d) and therefore might represent endogenous BAX tetramers loosely attached to the OMM.

3.6 Effect of recombinant Bcl-xL on BAX insertion/oligomerization and Cyt c release

In our experiments, recombinant Bcl-xL significantly inhibited Cyt c release induced by a combination of BAX and Ca^{2+} (Fig. 7a, compare lanes 3 and 4). Figure 7d shows statistical analysis of the Cyt c release. Despite inhibition of Cyt c release, Bcl-xL failed to attenuate BAX insertion and oligomerization in the OMM (Fig. 7b, compare lanes 3 and 4). Figure 7c illustrates statistical analysis of BAX insertion based on densitometry data obtained with individual BAX bands (1×BAX-5×BAX) shown in Figure 7b. Interestingly, using polyclonal anti-BAX antibody, we detected a distinct band with a molecular weight ~30 kDa (Fig. 7b, lane 6), which corresponded to molecular weight of Bcl-xL [52] and was strongly amplified after addition of exogenous Bcl-xL (Fig. 7b, compare lanes 1 and 6). It is possible that this band belonged to exogenous, recombinant Bcl-xL inserted into mitochondrial membranes in alkali-resistant manner.

3.7 Role of SH-redox state in BAX insertion/oligomerization and OMM permeabilization

Oxidation of BAX's cysteines and formation of disulfide bridges between BAX molecules favors BAX oligomerization and OMM permeabilization [38,39]. In our experiments, a reducing agent dithiothreitol (DTT, 20 mM) dismantled BAX dimers in the solution without mitochondria (Supplemental Fig. 3). We hypothesized that tBID- and Ca^{2+} -stimulated BAX insertion/oligomerization in the OMM and Cyt c release might depend on oxidation of SH-groups. Indeed, DTT added into the standard incubation medium significantly diminished BAX insertion/oligomerization stimulated by tBID or Ca^{2+} (Fig. 8b, c, compare lanes 3 and 4 versus 7 and 8). DTT also attenuated insertion/oligomerization of BAX in the absence of tBID or calcium (Supplemental Fig. 4). In addition, DTT inhibited BAX-mediated Cyt c release

stimulated by Ca^{2+} and to a much lesser extent by tBID (Fig. 8a, compare lanes 3 and 7 versus lanes 4 and 8) but failed to inhibit Cyt *c* release induced by tBID alone (Fig. 8a, compare lanes 2 and 6). On the other hand, DTT strongly inhibited the release of Smac/DIABLO, another mitochondrial apoptogenic protein with twice larger molecular weight than Cyt *c* [53,54], induced by tBID alone or by a combination of tBID and BAX (Fig. 8a). Interestingly, a combination of Ca^{2+} and BAX appeared to be ineffective in the release of Smac/DIABLO. Figure 8c shows statistical analysis of BAX insertion shown in Figure 8b. Figure 8d and e show statistical analysis of densitometry data obtained with Cyt *c* and Smac/DIABLO bands respectively. Importantly, DTT (20 mM) failed to inhibit mitochondrial swelling induced by Ca^{2+} (Fig. 8f) indicating that DTT effect could not be attributed to inhibition of the mPT. Thus, these experiments revealed for the first time an important role of the SH-redox state in the regulation of BAX insertion/oligomerization and in BAX-mediated OMM permeabilization in brain mitochondria.

4. Discussion

It has been established in early studies that the extent of Cyt *c* release correlates with the amount of BAX inserted in the OMM [17,18]. In addition, early studies suggested that OMM permeabilization required BAX oligomerization that occurred prior to BAX insertion into the OMM [17,18], whereas monomeric BAX neither integrated into the OMM nor released Cyt *c* [36]. In our study for the first time we clearly demonstrated that recombinant monomeric BAX readily self-integrated into the OMM of brain mitochondria and self-oligomerized. We found no evidence for tBID- or Ca^{2+} -induced oligomerization of BAX in the solution prior to interaction with mitochondria. Accordingly, our results suggest that BAX most likely integrates into the OMM as a monomer and that interaction of BAX with the OMM is necessary for BAX oligomerization. Our findings are consistent with reports showing that BAX insertion into the OMM or liposomal membrane preceded the oligomerization step [55,56]. Importantly, the amount of BAX inserted into the OMM in the absence of tBID or calcium was relatively high (see Fig. 2a, lanes 2 and 4, and Fig. 2b, lane 1). On the other hand, the amount of BAX oligomers (except dimers) in the BAX preparation was below the detection limit of western blotting (Fig. 1). Therefore, the amount of BAX inserted and oligomerized in the OMM did not correspond to the amount of BAX oligomers in the BAX preparation.

In our experiments, BAX self-insertion and oligomerization in the OMM resulted in a minute release of Cyt *c*. Our observation echoes early findings and several recent reports indicating that BAX translocation to mitochondria does not necessarily cause massive OMM permeabilization [57-61]. Additional factors appeared to be required for unleashing the permeabilizing activity of the membrane-inserted and oligomerized BAX. Earlier, Epan et al (2002) reported that the negative curvature in membranes that is essential for OMM permeabilization was promoted by tBID [62]. Correspondingly, in our experiments the lack of massive OMM permeabilization by BAX alone could be explained by the lack of changes in the membrane curvature. In our experiments, tBID and Ca^{2+} augmented BAX insertion/oligomerization in the OMM and strongly amplified membrane-permeabilizing activity of BAX. The Ca^{2+} -dependent amplification of BAX activity is of particular interest. Bearing in mind that BAX can cause Ca^{2+} efflux from the endoplasmic reticulum [63-66] and, hence, increase the likelihood of the Ca^{2+} -induced mPT [67], the Ca^{2+} -induced stimulation of BAX insertion/oligomerization in the OMM leading to enhanced OMM permeabilization might represent a feed-forward amplification loop ensuring successful, irreversible progression of the apoptotic program.

Previously, it was shown that Ca^{2+} stimulated BAX-mediated Cyt *c* release from isolated liver mitochondria [25]. However, the mechanism of this stimulation was not investigated further. In our study with isolated brain mitochondria, we demonstrated that the Ca^{2+} -induced

amplification of the BAX-mediated Cyt *c* release occurred parallel to augmented alkali-resistant BAX insertion/oligomerization in the OMM, and that both BAX insertion/oligomerization in the OMM and BAX-mediated Cyt *c* release were facilitated by mPT induction. Thus, our results suggest augmented BAX insertion/oligomerization a mechanistic link between the Ca²⁺-induced mPT and increased BAX-mediated Cyt *c* release. In contrast to Ca²⁺, tBID-stimulated BAX insertion, oligomerization, and Cyt *c* release appeared to be mPT-independent, but in this case augmented BAX insertion/oligomerization also correlated with the increased Cyt *c* release.

Anti-apoptotic Bcl-2, a close relative of Bcl-xL [68,69], can inhibit pro-apoptotic BAX activity by heterodimerizing with BAX [70,71] or by binding tBID and hence precluding tBID-dependent activation of BAX [56,72]. Whether Bcl-xL/BAX heterodimerization affected BAX insertion/oligomerization in the OMM or inhibited already inserted and oligomerized BAX remained unclear. In our experiments, recombinant anti-apoptotic protein Bcl-xL failed to prevent BAX insertion and oligomerization in the OMM. However, Bcl-xL strongly inhibited Cyt *c* release induced by a combination of BAX and Ca²⁺. Earlier, we showed that recombinant Bcl-xL inhibited Cyt *c* release induced by a combination of tBID and monomeric BAX [73]. Thus, our results support a scenario in which Bcl-xL inhibits inserted/oligomerized BAX and emphasize the fact that BAX insertion/oligomerization in the OMM could be dissociated from OMM permeabilization. How Bcl-xL restrains the inserted/oligomerized BAX from permeabilizing the OMM has yet to be determined. It seems conceivable that Bcl-xL could bind to the inserted/oligomerized BAX and physically block or disrupt the BAX pore, leading to inhibition of the BAX-mediated OMM permeabilization.

It is well established that apoptosis induced by different stimuli is often accompanied by an increase in ROS generation, and that suppression of ROS generation might protect cells against apoptosis [74-77]. Following ROS attack, critical SH-groups of different proteins might be oxidized leading to formation of intra- and inter-molecular disulfide bridges [78]. The exact role of the SH-redox state in the membrane-permeabilizing activity of BAX is not clear yet but it is possible that changes in intracellular SH-redox state could influence BAX conformation and thus stimulate BAX insertion/oligomerization in the OMM. Indeed, D'Alessio et al. (2005) demonstrated that oxidation of cysteine residues of BAX resulted in formation of disulfide bridges, causing conformational changes that favored BAX dimerization and translocation to mitochondria [38]. In our study, a reducing agent DTT inhibited tBID- and Ca²⁺-stimulated BAX insertion/oligomerization in the OMM, but only in the latter case DTT significantly suppressed Cyt *c* release. This suggests that Ca²⁺-stimulated BAX-mediated Cyt *c* release depends on oxidation of SH-groups whereas the tBID-stimulated BAX-mediated Cyt *c* release does not. It is conceivable that disruption of disulfide bridges between BAX molecules with DTT underlies a decrease in BAX insertion/oligomerization in the OMM affecting OMM permeability. Alternatively, DTT could antagonize the Ca²⁺-induced mPT [79] and hence hinder BAX-mediated Cyt *c* release. However, in our experiments DTT failed to inhibit mitochondrial swelling induced by Ca²⁺ ruling out this possibility.

The lack of correlation between diminished BAX insertion/oligomerization and virtually unchanged Cyt *c* release observed with tBID in the presence of DTT suggests that even small amounts of BAX inserted and oligomerized in the OMM might be sufficient for massive Cyt *c* release as proposed recently [80]. However, in our experiments, self-insertion and self-oligomerization of BAX in the OMM failed to induce massive Cyt *c* release, signifying a need for additional factors. It is also conceivable that the size of BAX pores formed with tBID remains large enough to pass Cyt *c* even in the presence of DTT whereas conductance of the Ca²⁺-activated BAX pores declines more significantly with DTT making the pores less passable for Cyt *c*. Our experiments with Smac/DIABLO release support this hypothesis. Smac/DIABLO is approximately twice larger than Cyt *c* [53,54]. While producing sizable Cyt

c release, a combination of BAX and Ca²⁺ failed to induce Smac/DIABLO release suggesting BAX pore size a limiting factor. DTT, which failed to inhibit tBID-stimulated BAX-mediated Cyt *c* release, at the same time strongly decreased the release of Smac/DIABLO. It is possible that reduction of disulfides with DTT affects not only insertion and oligomerization of BAX and, correspondingly, the number of BAX pores in the OMM, but also the size of the BAX pores. Thus, in addition to the amount of BAX inserted/oligomerized in the OMM, modulation of SH-redox state might influence molecular architecture of BAX oligomers that could be critical for effective OMM permeabilization.

Overall, our results strongly suggest that BAX-mediated OMM permeabilization in brain mitochondria can be modulated by the mPT and by SH-redox state. Correspondingly, induction of the mPT, increased ROS generation, and oxidation of critical SH-groups could significantly augment BAX-mediated permeabilization of the OMM and thus promote neuronal apoptosis in various neurodegenerative diseases, stroke, and traumatic brain injury.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Abbreviations used are

tBID	truncated BID
mPT	mitochondrial permeability transition
OMM	outer mitochondrial membrane
ER	endoplasmic reticulum
CsA	cyclosporin A
COX IV	cytochrome oxidase subunit IV
CHAPS	3-[(3-cholamidopropyl) dimethylammonio]-1-propanesulfonate
NP-40	Nonidet P-40, [octylphenoxy] polyethoxyethanol
OG	octyl glucoside
Tr X-100	Triton X-100
SDS-PAGE	sodium dodecyl sulfate-polyacrylamide gel electrophoresis
EGS	ethylene glycol bis(succinimidyl succinate)
DSS	disuccinimidyl suberate
BMH	bismaleimido-hexane

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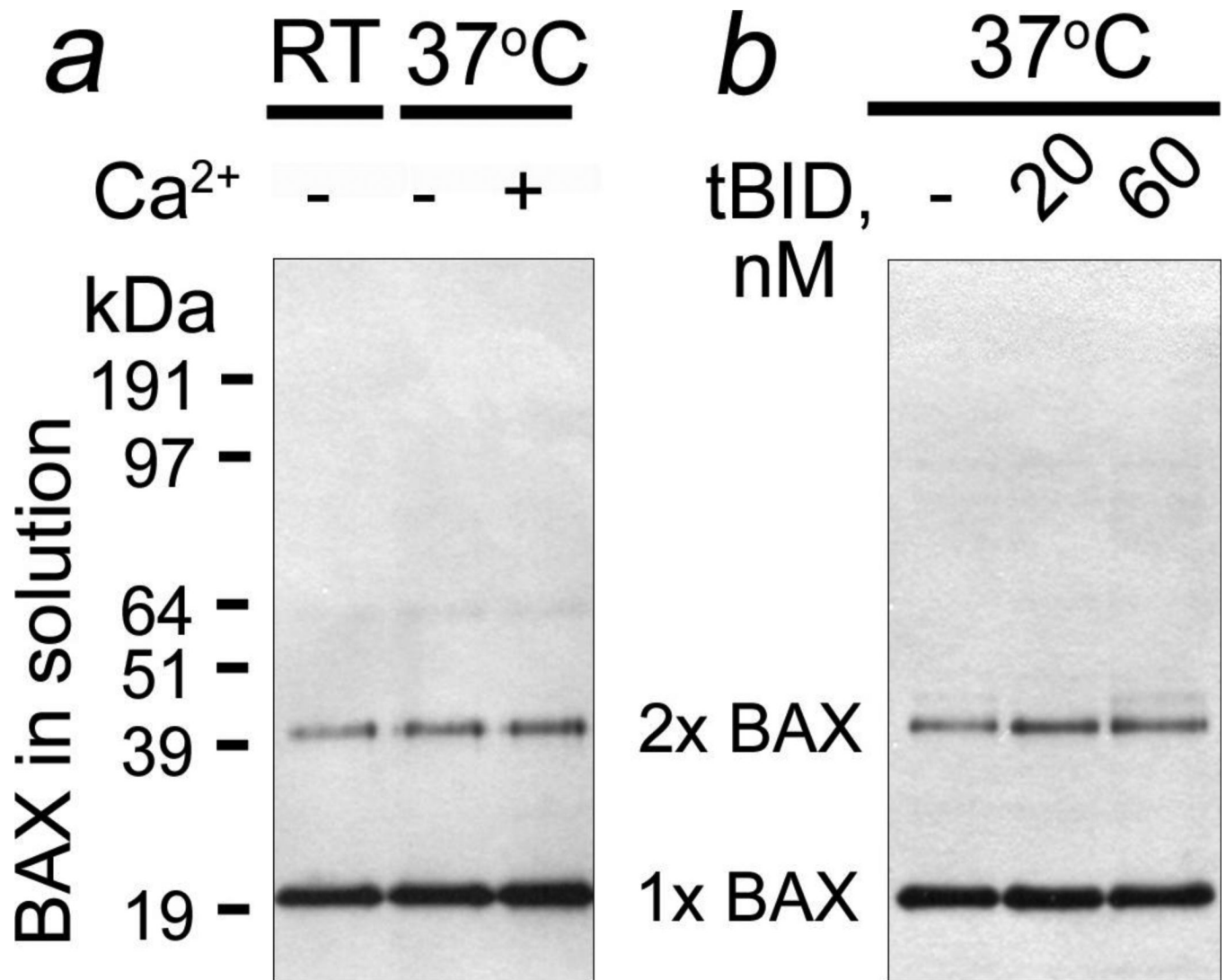


Figure 1. Ca²⁺ and tBID failed to induce BAX oligomerization in the solution without mitochondria
 In *a*, 50 nM BAX was incubated in the standard mitochondrial incubation medium with or without 250 μM Ca²⁺ for 30 minutes at room temperature (RT, 23°C) or 37°C as indicated. In *b*, 50 nM BAX was incubated with 20 or 60 nM tBID for 30 minutes at 37°C in the standard incubation medium. In these and other experiments with BAX, western blotting followed non-reducing SDS-PAGE.

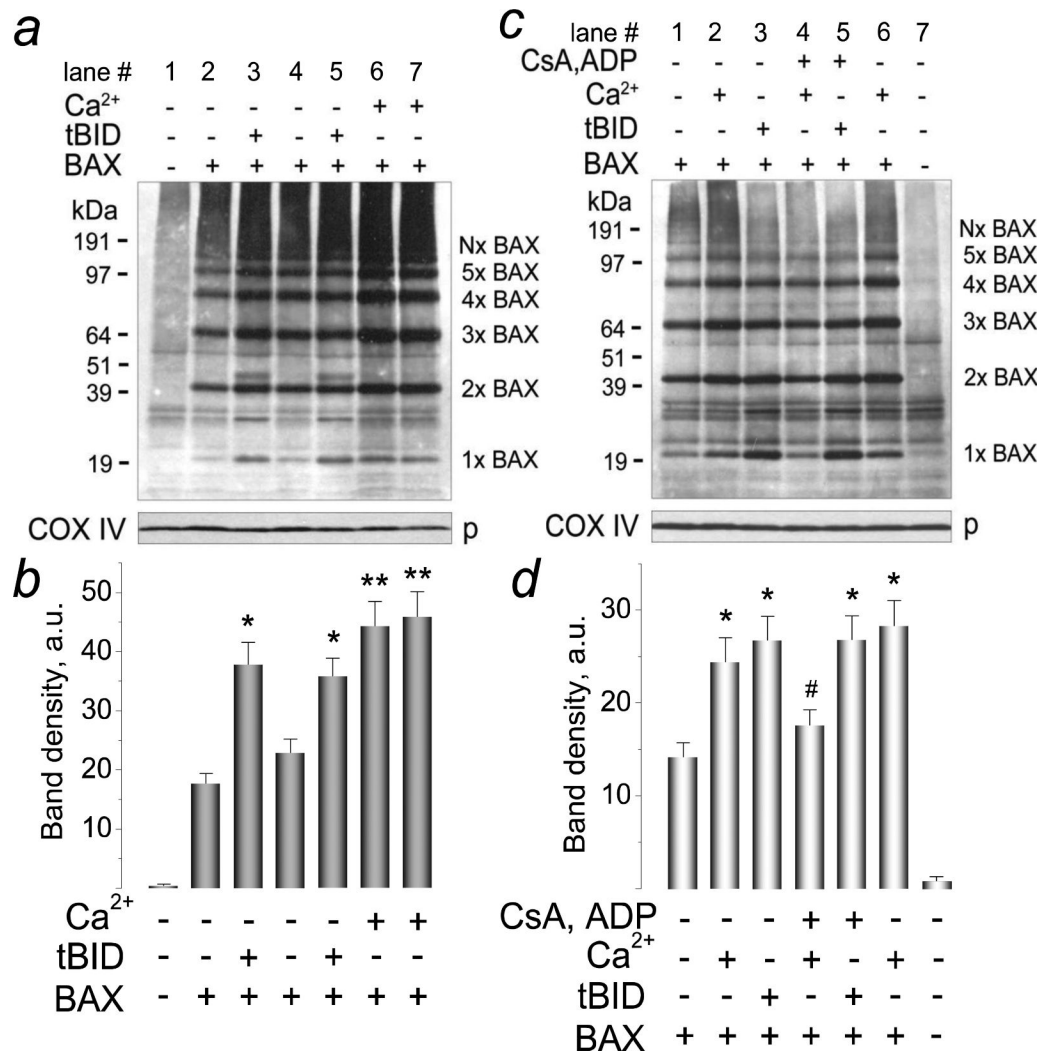


Figure 2. Recombinant BAX self-integrated and self-oligomerized in the outer mitochondrial membrane of isolated brain mitochondria. Ca²⁺ and tBID stimulated BAX insertion and oligomerization. Cyclosporin A (CsA) and ADP antagonized the effect of Ca²⁺ but did not influence the effect of tBID

In **a**, alkali-resistant BAX insertion and oligomerization with and without Ca²⁺ or tBID. Mitochondria were treated for 30 minutes at 37°C with 50 nM BAX, with 1.4 μmol Ca²⁺/mg protein, or with 20 nM tBID as indicated. Following alkali treatment (30 minutes on ice in 0.1 M Na₂CO₃ at pH 11.5), mitochondrial membranes were solubilized with 1% CHAPS. In **c**, alkali-resistant BAX insertion and oligomerization with and without Ca²⁺ or tBID and the effect of CsA plus ADP. Mitochondria were treated for 30 minutes at 37°C with 50 nM BAX, with 1.4 μmol Ca²⁺/mg protein, with 20 nM tBID, and with a combination of 1 μM CsA and 100 μM ADP (in the presence of 1 μM oligomycin to prevent ADP phosphorylation) as indicated. Following alkali treatment, mitochondrial membranes were solubilized with 1% Nonidet P-40. **p**, pellets. In **b** and **d**, statistical analyses of BAX insertion into the OMM were performed using one-way ANOVA followed by Bonferroni's *post-hoc* test. The graphs show the averaged sum of 1×BAX–5×BAX band densities under corresponding experimental conditions. Here and in all other similar experiments, the densitometry of individual bands was performed with NIH ImageJ 1.40g software (<http://rsb.info.nih.gov/ij/>). Data are mean±SEM, N=3. In **b**, **p*<0.05 comparing band densities with BAX versus BAX plus tBID; ***p*<0.01

comparing band densities with BAX versus BAX plus Ca^{2+} . In **d**, $*p < 0.01$ comparing band densities with BAX versus BAX plus tBID or BAX plus Ca^{2+} ; $\#p < 0.05$ comparing band densities with BAX plus Ca^{2+} versus BAX plus Ca^{2+} with CsA and ADP. These and all other experiments with BAX were performed without cross-linkers.

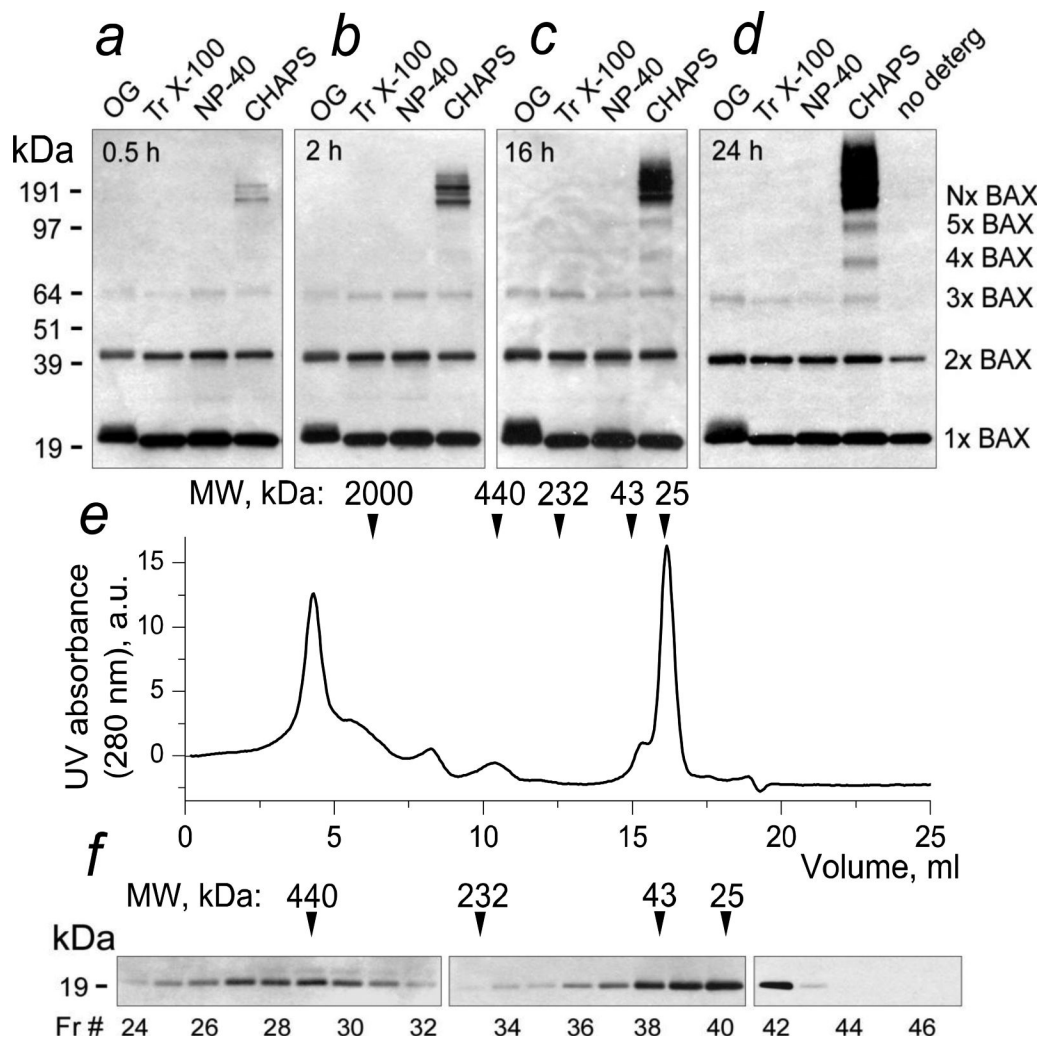


Figure 3. CHAPS but not Nonidet P-40, Triton X-100, or octyl glucoside produced oligomerization of BAX in the solution without mitochondria. Analytical gel-filtration confirmed BAX oligomerization by CHAPS

In *a-d*, BAX (50 nM) was incubated for 0.5-24 hours in the incubation medium at 4°C either with 1% octyl glucoside (OG), or 1% Triton X-100, or 1% Nonidet P-40, or 1% CHAPS, or without detergent. In all cases, the amount of protein loaded on the gel was the same (100 ng/lane). In *e* and *f*, BAX (500 nM) was incubated with 1% CHAPS for 24 hours at 4°C. Larger BAX concentration was used to compensate dilution during gel-filtration. Then, BAX incubated with the detergent was passed through a Superdex-200 HR 10/30 column equilibrated with 125 mM KCl, 10 mM HEPES, pH 7.4, and 1% CHAPS. In *e*, a UV absorbance (280 nm) profile of the gel-filtration eluate. **a.u.**, arbitrary units. In *f*, western blotting analysis of the collected fractions. Fr, fraction.

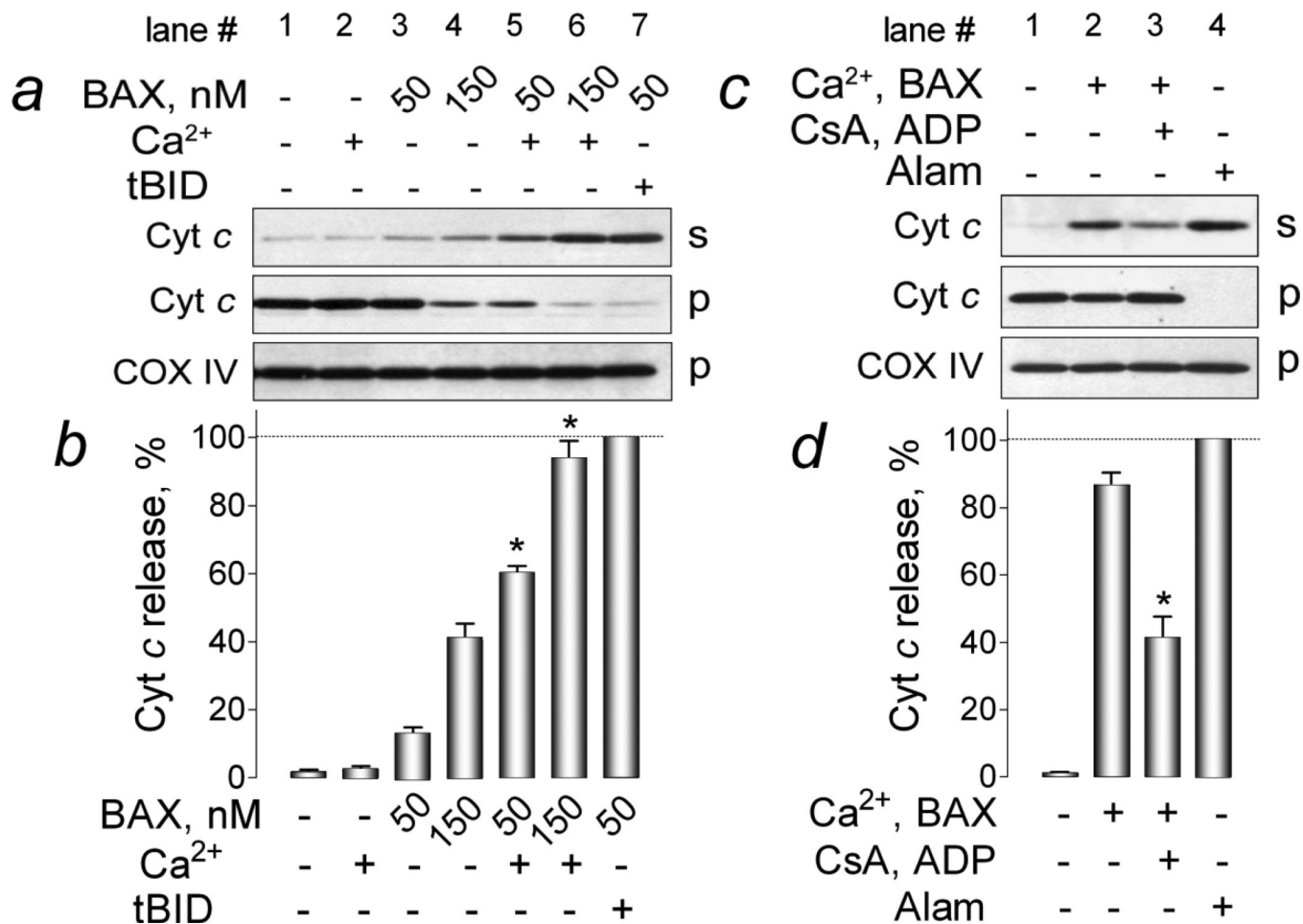


Figure 4. Ca²⁺ amplified cytochrome *c* release induced by recombinant monomeric BAX. Cyclosporin A and ADP, inhibitors of the mitochondrial permeability transition, attenuated the Ca²⁺ effect

In **a**, the effect of Ca²⁺ and tBID on cytochrome *c* (Cyt *c*) release induced by BAX. Where indicated, 250 μM Ca²⁺ (1.4 μmol Ca²⁺ per mg of mitochondrial protein) or 20 nM tBID was added. Here and in other experiments, COX IV was used as a loading control. In **b**, densitometry analysis of western blots performed with ImageJ software. The densitometry results are shown as percentage of Cyt *c* release. The Cyt *c* release induced by 50 nM BAX plus 20 nM tBID was taken as 100%. Data are mean ± SEM, **p* < 0.01 between Cyt *c* release induced by 50 or 150 nM BAX alone or in the presence of 1.4 μmol Ca²⁺/mg protein, N=3. In **c**, a combination of cyclosporin A (CsA, 1 μM) and ADP (100 μM) (here and in other similar experiments, ADP was added in the presence of 1 μM oligomycin, an inhibitor of ATP synthase, to prevent ADP phosphorylation) decreased Cyt *c* release induced by 50 nM BAX in the presence of 1.4 μmol Ca²⁺/mg protein. In these experiments, alamethicin (Alam, 30 μg/ml) was used to produce the maximal Cyt *c* release. In **d**, densitometry analysis performed with ImageJ software. The densitometry results are shown as percentage of Cyt *c* release. The Cyt *c* release induced by 30 μg/ml alamethicin was taken as 100%. Data are mean ± SEM, **p* < 0.01 between Cyt *c* release induced by BAX plus Ca²⁺ in the absence or in the presence of CsA and ADP, N=4. In **a** and **c**, s, supernatants; p, pellets.

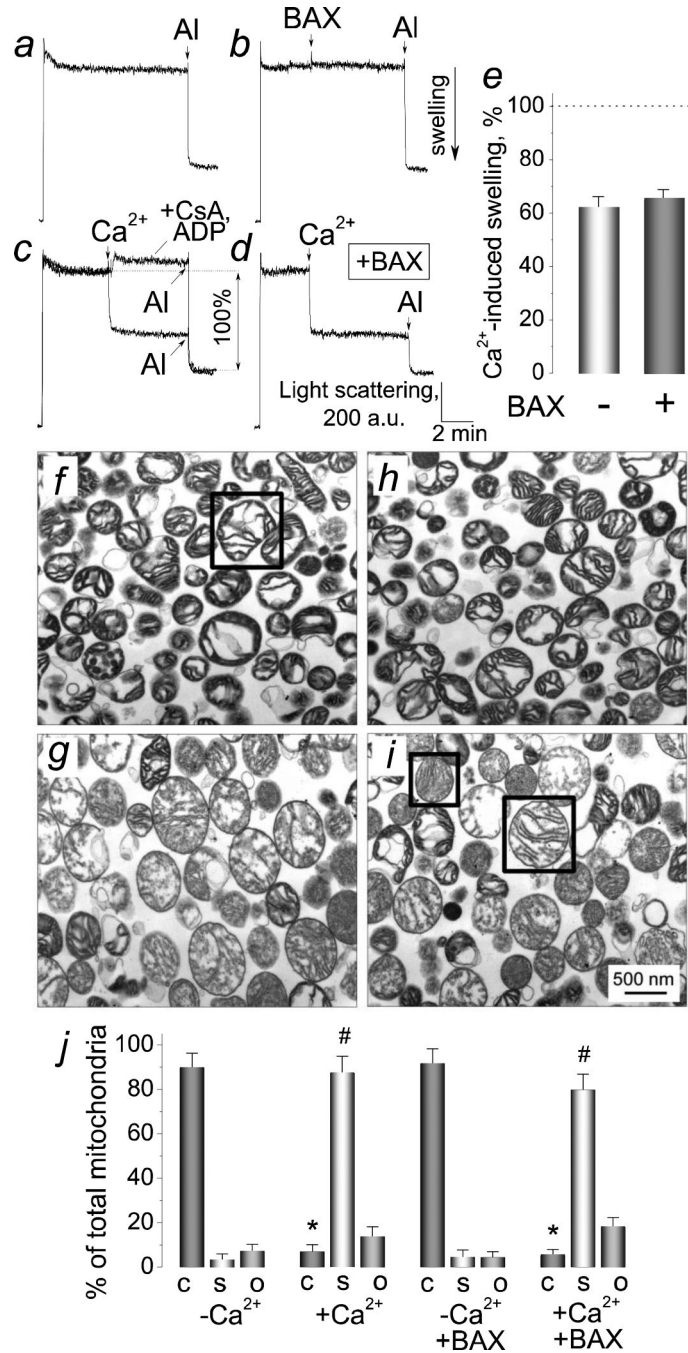


Figure 5. Recombinant monomeric BAX did not cause mitochondrial swelling and failed to augment Ca²⁺-induced mitochondrial swelling

In *a-d*, 90° light scattering traces indicative of changes in mitochondrial volume obtained with or without Ca²⁺ and BAX. Here, 30µg/ml alamethicin was added as indicated to cause maximal mitochondrial swelling. In *c-d*, 1.4µmol Ca²⁺/mg protein was added to mitochondria where indicated. In *b*, 50 nM BAX was added as indicated. In *d*, 50 nM BAX was added to the cuvette prior to mitochondria. In *e*, the extent of Ca²⁺-induced mitochondrial swelling with or without 50 nM BAX calculated as a percentage of maximal, alamethicin-induced swelling taken as 100% (*c*). In *f-i*, electron micrographs of isolated brain mitochondria incubated with or without Ca²⁺ and BAX. In *f* and *g*, mitochondria were incubated without BAX, but were treated with

a vehicle (3 μ l of the dialysis buffer for BAX (25 mM HEPES-NaOH, pH 7.5, 0.2 mM dithiothreitol, 30% glycerol (v/v)) in 300 μ l of incubation medium). In **h** and **i**, mitochondria were incubated for 30 minutes at 37°C in the medium supplemented with 50 nM BAX. In **g** and **i**, the incubation medium was supplemented with 1.4 μ mol Ca²⁺/mg protein. In **j**, morphometric analysis of mitochondria incubated under the indicated conditions. Total mitochondrial population was categorized into three groups according to their morphology: condensed (C), swollen (S), and orthodox (O). The representative example of condensed morphology is shown in the box in *panel f*; swollen (small box) and orthodox (large box) morphology in *panel i*. * p <0.001, comparing amount of condensed mitochondria with and without Ca²⁺; # p <0.001, comparing amount of swollen mitochondria with and without Ca²⁺. Data are mean \pm SEM, N=3.

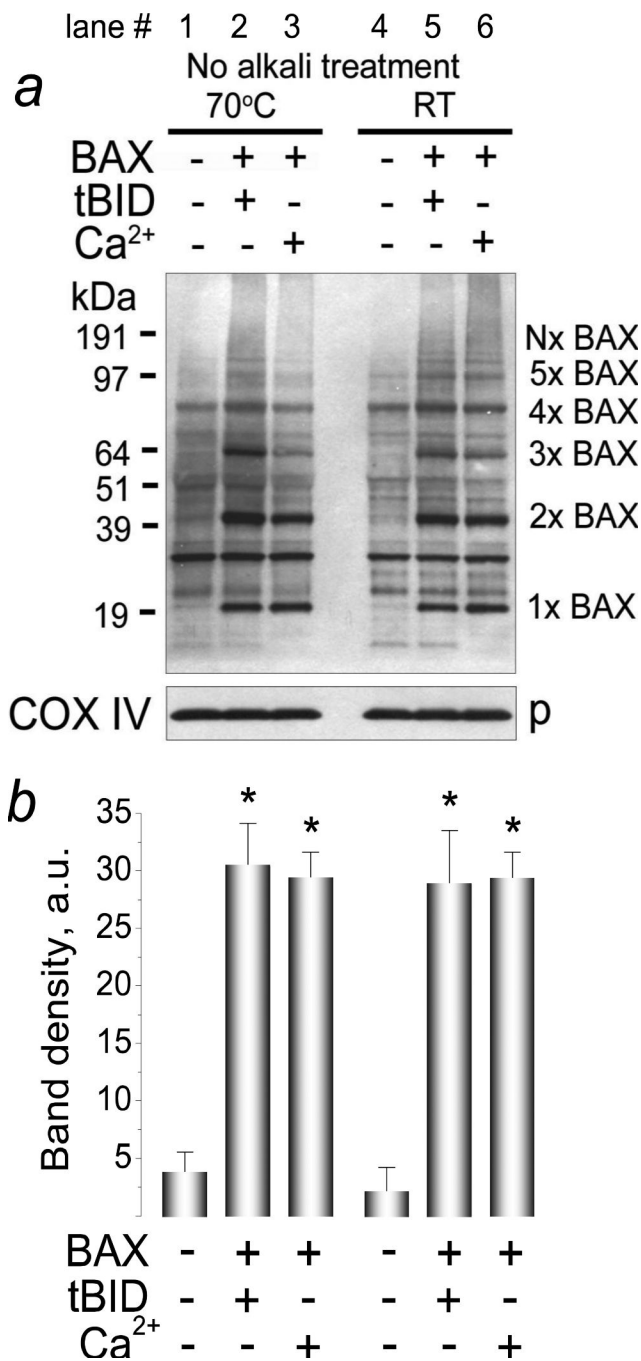


Figure 6. Alkali treatment of mitochondria and heating the samples before SDS-PAGE were not essential for detecting BAX oligomers

The experiments were performed as described in the legend to Figure 2 but without alkali treatment of mitochondria and without heating the mitochondrial samples before SDS-PAGE. In **a**, mitochondria were incubated for 30 minutes at 37°C with 50 nM BAX in combination with 20 nM tBID or 1.4 μmol Ca²⁺/mg protein as indicated. Mitochondrial pellets were solubilized with 1% Nonidet P-40. **p**, pellets. In **b**, statistical analysis of BAX insertion/oligomerization in the OMM was performed using one-way ANOVA followed by Bonferroni's *post-hoc* test. The graph shows the averaged sum of 1×BAX–5×BAX band densities under

corresponding experimental conditions. Data are mean \pm SEM, N=3. In **b**, * p <0.01 comparing band densities with or without BAX plus tBID or BAX plus Ca²⁺.

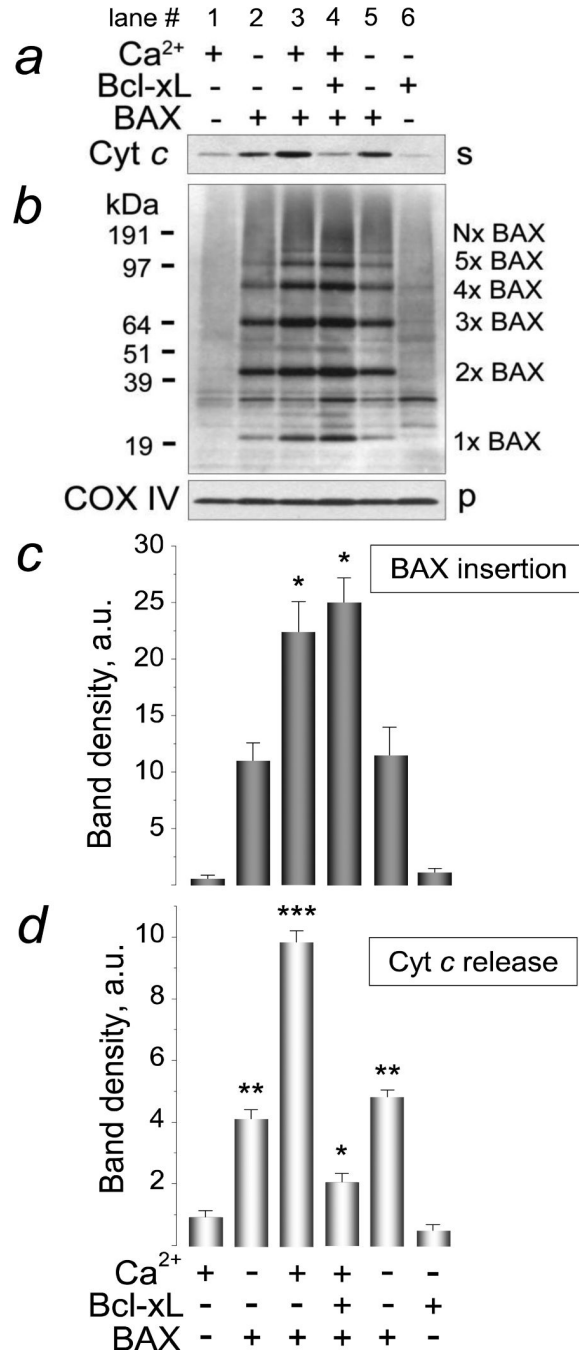


Figure 7. Recombinant Bcl-xL inhibited cytochrome *c* release but failed to attenuate BAX insertion and oligomerization in the outer mitochondrial membrane augmented by Ca²⁺

Mitochondria were treated for 30 minutes at 37°C with 150 nM BAX, 1.4 μmol Ca²⁺/mg protein, and with 300 nM Bcl-xL as indicated. After 30 minutes of incubation, mitochondria were pelleted by centrifugation and cytochrome *c* (Cyt *c*) release was evaluated in the supernatants (**a**). In **b**, alkali-resistant BAX insertion was assessed in mitochondrial membranes solubilized with 1% Nonidet P-40. **s**, supernatants, **p**, pellets. In **c**, the statistical analysis of BAX insertion into the OMM was performed using one-way ANOVA followed by Bonferroni's *post-hoc* test. The graph shows the averaged sum of 1×BAX–5×BAX band densities under corresponding experimental conditions. Data are mean±SEM, N=3. **p*<0.01 comparing band

densities with BAX versus BAX plus Ca^{2+} or BAX plus Ca^{2+} and Bcl-xL. In *d*, the densitometry analysis of Cyt *c* release. Data are mean \pm SEM, * p <0.05, ** p <0.01, *** p <0.001 versus Cyt *c* release in the presence of Bcl-xL alone.

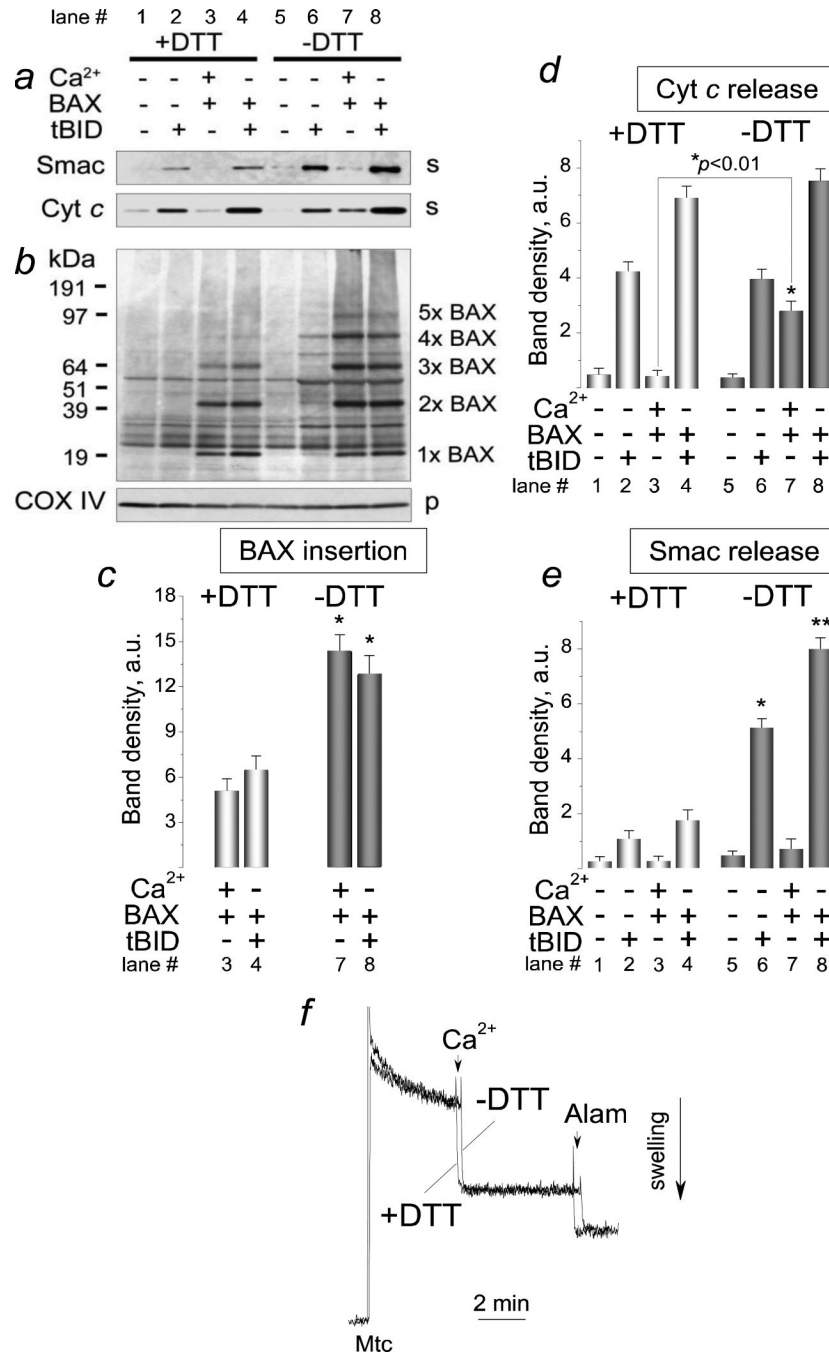


Figure 8. Dithiothreitol attenuated BAX insertion/oligomerization and outer mitochondrial membrane permeabilization but failed to inhibit mitochondrial swelling induced by Ca²⁺ Mitochondria were treated for 30 minutes at 37°C with 50 nM BAX, with 1.4 μmol Ca²⁺/mg protein, and with 20 nM tBID as indicated. Where indicated, the incubation medium was supplemented with 20 mM dithiothreitol (DTT). Then, mitochondria were pelleted by centrifugation and cytochrome *c* (Cyt *c*) and Smac/DIABLO (Smac) releases were evaluated in the supernatants (**a**). In **b**, alkali-resistant BAX insertion was assessed in mitochondrial membranes solubilized with 1% Nonidet P-40. **s**, supernatants, **p**, pellets. In **c**, statistical analysis of BAX insertion into the OMM was performed using one-way ANOVA followed by Bonferroni's *post-hoc* test. The graph shows the averaged sum of 1×BAX–5×BAX band

densities under corresponding experimental conditions. Data are mean \pm SEM, N=3. * p <0.01 comparing BAX band densities with or without 20 mM DTT. In **d**, the densitometry analysis of Cyt *c* release. In **e**, the densitometry analysis of Smac/DIABLO (Smac) release. Data are mean \pm SEM, N=3, * p <0.01, ** p <0.001 comparing Smac/DIABLO release in the presence or absence of 20 mM DTT. In **f**, mitochondrial swelling followed with a 90° light scattering assay. Light scattering traces obtained with and without dithiothreitol (DTT, 20 mM) are overlapped for comparison. Where indicated, 1.4 μ mol Ca²⁺/mg protein, and 30 μ g/ml alamethicin (Alam) were added to mitochondria