The Dynamics of Bax Channel Formation: Influence of Ionic Strength

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SUPPLEMENTARY INFORMATION

MATERIALS AND METHODS

Preparation of cytochrome c

Horse heart cytochrome c (44 mg) was reduced with 11 mg ascorbate in 1 mL of 0.2 M Tris-HCl pH 7.5. The reduced cytochrome c was separated from the ascorbate by gel filtration through a Sepha dex G-10 column equilibrated with 0.2M Tris-HCl pH 7.5.

Isolation of rat liver mitochondria

Fresh mitochondria were purified from rat liver as described (1). The BSA was removed by sedimenting mitochondria in BSA free isolation buffer and the fi nal pellet resuspended in 300 mM mannitol, 5 mM HEPES and 0.1 mM EGTA pH 7.4 (FH-buffer) to a final concentration of 10-20 mg mitochondrial protein/mL. The mitochondrial stock was always kept cold and dilutions were made periodically from the stock for experiments.

Purification of recombinant proteins

Recombinant human Bax was produced as described (2) but the purification procedure was modified. The Bax eluted from the chitin column was dialyzed (12000 MW cutoff) at 4°C for 24 hrs against 3L of 1m M EDTA, 10mM Tris-HCl, pH 8. Then it was dialy zed again for 24 hrs against 5L of 10 mM Tris-HCl pH 8 to remove all remaining DTT. Reco mbinant tBid w as purified as described (3). Bcl-xL was purified as described (4 - 6). All proteins were filter-sterilized (0.2 μ m filter), gl ycerol added (10% fina l), rapidl y flash-frozen with ethanol and dr y-ice, and stored at -80 °C. Prote in concentration was deter mined after dialysis by the BCA method.

Western blot analysis

For all western blotting experiments, three 1mL mitochondrial suspensions (960 μ g/mL) were incubated at 30°C under t he conditions and for the times indicated in the figure legends. The triplicates were centrifuged (10,000 RCF for 5 m in) and each resuspended into 200 μ L of FH buffer. These were pooled together, mitochondria repelleted and subjected to sodium carb onate treat ment as described (7). The pellet was resuspended in 50 μ L of 2% SDS and incubated for 1 hr on ice. Then, the detergent solubilized fraction was separated from the detergent-insoluble fraction by centrifugation and 30 μ L used for PAGE/blotting analysis. VDAC served as loading control. Bax was probed with rabbit anti-Bax (Ab7977: 1:1000) and VDAC was probed with rabbit anti-VDAC anti-serum (1:1,000,000). Bot h the primary antibodies were probed with HRP tagged anti-rabbit secondary antibodies (1:2500 for Bax and 1:5000 for VDAC). Blots were developed with DAB as described in (8).

Cytochrome c accessibility assay

The cy tochrome *c* oxidation or accessi bility assay was perfor med as describ ed previously (6). After appropriate treatment of 96 μ g/mL mitochondria with Bcl-2 family proteins, 100 μ L of this suspension was dispersed in 600 μ L of cy tochrome *c* oxidation assay buffer (equivalent to FH buffer (mannitol + KCl (total 300 mOs), 0.1 mM EGTA, 5 mM HEPES pH 7.5) of appropriate ionic strength supplemented with

5 mM DNP and 5 μ M antimycin A) and quickly assayed for c ytochrome c oxidation by adding 10 μ L of reduced cy tochrome c to a final concentration of ~ 25 µM. Cytochrome c oxidation was measured as decline in absorbance at 5 50 nm. The rate of oxidatio n was calculated from the slope of the linear region of the absorbance chan ge measured for 2 minutes. Maxi mal rate of cytochrome c oxidation (corresponding to 100% perm eabilization) was determ ined by the rate of c ytochrome c oxidation by equivalent amount of hypotonically shocked mitochondria (at least 1 volume of mitochondrial suspension to 50 volumes of double distilled H_2O). Hypotonic lysis breaks the MOM and exposes all the cytochrome oxidase to exogenous cytochrome c. The % oxidation rate was calculated with maximal oxidation rate as 100% and rate of oxidation by control mitochondria (no treatment) as 0 %. Cytochrome oxidase activity was strongly sensitive to ionic strength, so control and maximal oxidation rates were determined for each num erical values of control, Bax-tBid treated and maxi salt concentr ation tested. The actual mal cytochrome c oxidation rates at 10 and 90mM KCl are presented (Fig S1A) for comparison.

Adenylate kinase (AK) assay

The assay was perfor med as described previously (6). This assay measures the release of the 24 kDa protein, adenylate kinase, from the IMS of mitochondria. In this coupled enzyme assay, NADP is reduced to form NADPH which has a strong abs orbance at 340 nm. (The oxidized form NADP does not have this absorbance). Prior to ass aying, 2.5 µL of a mix of Hexokinase/Glucose-6-Phosphate Dehydrogenase (5 units each) was incubated in 350 µL AK buffer (5 mM MgSO₄, 10 mM glucose, 5 mM ADP, 0.2 mM NADP, 50 mM Tris, and pH 7.5) to re move r esidual ATP. After 2-3 m inute incubation, 150 µL of mitochondrial supernatant was added to the AK buffer/ enzyme mix, and absorption was measured for 5 minutes at 340 nm. The initial linear slope of increase in absorption was taken as the rate of AK reaction. Since all other reaction components are in excess, the concentration of AK is the rate-li miting factor. Hence the rate of NADP H formation is a direct measure of am ount of AK released from mitochondria. Maximal rate of NADPH form ation, corresponding to 100% release of AK from mitochondria, was measured from the activit y of su pernatant from equivalent am ount of h ypotonically shocked mitochondria. The actual numerical values of AK activity used as control (0% AK activity or no release of AK), Bax-tBid treated and I ysed (100% AK activ ity or complete release of AK) are presented in Fig S1B for comparison.

Sulfite oxidase (SOX) assay

SOX is a 120 kDa IMS protein and its release from mitochondria was assay ed as described (9). SOX reduces cy tochrome *c* using sulfite as the electron donor. Prior to the assay, a one-to-one solution of 40 mM Na $_2$ SO₃ and 5 0 mg/ml oxidized cytochrom e *c* was made. Then 10 µL of this solution (0.4 mM Na $_2$ SO₃ and 0.5 mg oxidized cy tochrome *c* final) was added to 50 0 µL mitochondrial supernatant and the absorbance was quickly measured at 550 nm for 5 minutes. Maximal release was measured from the activity of supernatant from equivalent amount of hypotonically shocked mitochondria. SOX activity was strongly sensitive to ionic strength. Hence, the supernatant was supplemented with suitable amount of salt by diluti on from stock salt solution to adjust the ionic strength. The actual numerical values of SO X activity used as control (0% SOX activity or no release of SOX), Bax-tBid treated and lysed (100 % SOX activity or complete release of SOX) are presented in Fig S1C for comparison.

Measurement of fumarase activity

Fumarase is a matrix enzyme that reversibly converts malate to fumarate. Fumarase release was measured by a standard procedure (10). Briefly 96 μ g/mL mitochondrial suspension in F H buffer was treated with varying conc entrations of digitonin (from 0.0025% to 0.1% in multiples of 2) for 5 m inutes at room temperature and centrifuged for 5 m inutes at 14000 RCF and supernatant collected. 500 μ L of th e supernatant was mixed with 500 μ L of fumarase assay buffer (50 mM malate, 50 mM Na₂HPO₄ pH 7.3) and change in absorbance measured for 5 m inutes at 250 nm. Maximal fumarase release corresponding to 100% was obtained by treating equivalent amount of mitochondria with 0.2% Triton-X100.

SUPPORTING REFERENCES

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FIGURE S1: Measured rates of enzym atic activity in a set of experiments showing the rates prior to any treatment (control rates), the rate s after tBid/Bax treatment, and the maximal rates after hypotonic shock. The influence of ionic streining the number of these rates is illustrated. By subtracting the control rates and expressing the results as a percentage of the maximal rates we correct for background rates a ind effects of i onic strength. The figure shows the rates of cytochrome c (25 µM) oxidation (A), AK activity as measured by absorbance (OD) change at 340nm (B) and SOX activity as measured by absorbance change at 550nm (C) recorded from 96µg/mL mitochondrial suspensions that were either untreated (c ontrol), or treated with 17nM Bax and 120nM tBid or osm otically lysed (lysed). Incubations (30 m in) were in either 10 or 90mM KCl buffer. * represents P value < 0.05 and ** represents P value < 0.01. n= 3.



FIGURE S2: Control experiments and evidence of ionic strength dependence. A, B: Bax and tBid alone have little or no effect on MOMP. The rates of cytochrom e *c* oxidation (A) or AK release (B) by 96 μ g/mL m itochondrial suspension are expressed as a percent of the m aximal amount. Except as indicated, m itochondria were incubated for 30 m in with 17nM Bax and/or 120nM tBid in the presence of ei ther 10mM (black bars) or 90m M KCl (grey bars) buffer. C. The rates of cytochrom e *c* oxidation (expressed as a % of th e maximal values) induced by a 30 min incubation with 17nM Bax and 120nM tBid in the presence of varying concentrations of either NaCl (solid line) or pot assium lactobion ate (K lac, broke n line). * represents P value < 0.05. n = 3.



Figure S3: MOMP induced by Bax and tBid is in dependent of PTP. Mitochondria were incubated with 35nM Bax and 120nM tBid in the presence or absence of 50 μ M CsA at either 10mM KCl or 90mM KCl for 30 m inutes at 30 °C. Then, the mitochondria from each treatment were assay ed for real-time permeabilization using the cytochrome *c* oxidation assay and the supernatant analyzed for extent of AK release. The statistical tests are between the black bars in each panel. Both are significantly different ** P value < 0.01 and *** P value < 0.001 n = 3.



Figure S4: Comparison of maximal cytochrome c ox idation obtained under swelling and nonswelling conditions. A) At 0.01% digitonin, no activ ity of fumarase, a matrix enzyme, could be detected, but at 0.02% nearly 20% of fumarase was released from 96 µg/mL m itochondrial suspension. B) 96 µg/mL m itochondrial suspension was treated with 0.01% digitonin for 5 minutes either without tBid or after pre-inc ubation of the m itochondrial suspension with 120nM tBid for 30 m inutes at 30 °C at either ionic strength. The n, the m itochondrial suspension was assayed for the rate of cytochrom e c oxidation. The m ean values are the % of cytochrom e c oxidation rate observed with an equivalent am ount of osmotically lysed m itochondria at the corresponding ionic strengths. No statistically significant difference was found between digitonin treated samples (with or without tBid) and osmotically lysed samples. (P values > 0.15). n = 3.



Figure S5: Bax induced MOMP increased at a faster rate at lower ionic strength. Mitochondrial suspensions were treated with 34nM monomeric Bax and 120nM tBid at 30 °C at either 10mM KCl buffer (solid lines) or 90m M KCl buffer (d ashed lines) and the rate of cytochrom e *c* oxidation was determined at different time intervals. * represents P value < 0.05, ** - P value < 0.01. n = 3.



FIGURE S6: Bcl-xL m ediated inhibition of MOMP is sensitive to tBid con centration. Mitochondrial suspensions were incubated w ith 17nM monomeric Bax, either 24nM or 120nM tBid, without Bcl-xL or with 48n M Bcl-xL in 10mM KCl buffer or 90m M KCl buffer for 30 minutes at 30 °C. Then, the extent of AK release was determined for the different treatments. The differences in the inhibitory effects of Bcl-xL at 10 and 90 m M were statistically significant with P<0.0001 at both tBid concentrations. n = 3.



FIGURE S7: Bcl-xL inhibited Bax m ediated MOMP without affecting Bax insertion. Mitochondrial suspensions (960µg mitochondrial protein/mL) were treated with 100nM Bax and 240nM tBid for 10, 20 and 30 m inutes without Bc l-xL or with 100nM Bcl-xL (for 30 m inutes only) in either 10m M KCl buffer or 90m M KCl bu ffer. The western blots were perform ed as described in Experimental Procedures. VDAC served as loading control. Band L corresponds to 10% of the total Bax expected to be present if all the added Bax inserted into the m itochondrial outer membrane.