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M Madesh
Thomas Jefferson University

György Hajnóczky
Thomas Jefferson University

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VDAC-dependent permeabilization of the outer mitochondrial membrane by superoxide induces rapid and massive cytochrome *c* release

Muniswamy Madesh and György Hajnóczky

Department of Pathology, Anatomy and Cell Biology, Thomas Jefferson University, Philadelphia, PA 19107

Enhanced formation of reactive oxygen species (ROS), superoxide ($O_2^{\cdot-}$), and hydrogen peroxide (H_2O_2) may result in either apoptosis or other forms of cell death. Here, we studied the mechanisms underlying activation of the apoptotic machinery by ROS. Exposure of permeabilized HepG2 cells to $O_2^{\cdot-}$ elicited rapid and massive cytochrome *c* release (CCR), whereas H_2O_2 failed to induce any release. Both $O_2^{\cdot-}$ and H_2O_2 promoted activation of the mitochondrial permeability transition pore by Ca^{2+} , but Ca^{2+} -dependent pore opening was not required for $O_2^{\cdot-}$ -induced CCR. Furthermore, $O_2^{\cdot-}$ alone evoked CCR without damage of the inner mitochondrial membrane barrier, as mitochondrial membrane potential was sustained in the presence of extramitochondrial ATP. Strik-

ingly, pretreatment of the cells with drugs or an antibody, which block the voltage-dependent anion channel (VDAC), prevented $O_2^{\cdot-}$ -induced CCR. Furthermore, VDAC-reconstituted liposomes permeated cytochrome *c* after $O_2^{\cdot-}$ exposure, and this release was prevented by VDAC blocker. The proapoptotic protein, Bak, was not detected in HepG2 cells and $O_2^{\cdot-}$ -induced CCR did not depend on Bax translocation to mitochondria. $O_2^{\cdot-}$ -induced CCR was followed by caspase activation and execution of apoptosis. Thus, $O_2^{\cdot-}$ triggers apoptosis via VDAC-dependent permeabilization of the mitochondrial outer membrane without apparent contribution of proapoptotic Bcl-2 family proteins.

Introduction

The superoxide ($O_2^{\cdot-}$)* free radical has come to play a pivotal role in a wide variety of pathophysiological conditions. In an aerobic system, sequential reduction of molecular oxygen leads to the generation of $O_2^{\cdot-}$, which is rapidly converted to H_2O_2 by superoxide dismutase (SOD) (Jacobson, 1996). $O_2^{\cdot-}$ can also rapidly react with another free radical, nitric oxide (NO), yielding more toxic reactive oxygen species (ROS), such as peroxynitrite ($ONOO^-$). In physiological systems, for example, ROS play an important role in signaling

mechanisms, especially in the $TNF\alpha$ -regulated pathways (Hennet et al., 1993). $O_2^{\cdot-}$ is considered one of the primary ROS generated in cells stimulated by $TNF\alpha$ (Hennet et al., 1993). Furthermore, ROS are known to play a part in proliferation and cell death (Buttke and Sandstrom, 1994; Patel et al., 1996; Fiskum, 2000; Yuan and Yankner, 2000). In several models of apoptosis, increased formation of ROS was described as an early event (Petit et al., 1995; Zamzami et al., 1995a). Most of the ROS have the potential to trigger apoptosis (Buttke and Sandstrom, 1994; Lin et al., 1995; Jacobson, 1996; Hildeman et al., 1999). In addition to enhanced ROS formation, impairment of the cellular antioxidant mechanisms may also lead to apoptosis (Rothstein et al., 1994). Decreased antioxidant profiles were found in epithelial cells that undergo apoptosis (Madesh et al., 1999), and antioxidants have been reported to counter apoptotic cell death (Hockenbery et al., 1993; Zamzami et al., 1995a,b).

Central components of the programmed cell death machinery, which have been conserved throughout evolution, include caspase cascade, Apaf-1 complex formation, and heterodimerization of Bcl-2 family proteins. Cellular organelles, particularly mitochondria, play a crucial role by releasing

Address correspondence to György Hajnóczky, Dept. of Pathology, Anatomy, and Cell Biology, Room 253, Thomas Jefferson University, Philadelphia, PA 19107. Tel.: (215) 503-1427. Fax: (215) 923-2218. E-mail: gyorgy.hajnoczky@mail.tju.edu

*Abbreviations used in this paper: BA, bongkreic acid; CCR, cytochrome *c* release; CsA, cyclosporin A; cyto *c*, cytochrome *c*; DIDS, 4,4'-diisothiocyanatostilbene-2,2'-disulfonic acid; GFP, green fluorescent protein; IMM, inner mitochondrial membrane; KP, König's polyanion; NO, nitric oxide; OMM, outer mitochondrial membrane; $ONOO^-$, peroxynitrite; $O_2^{\cdot-}$, superoxide; PI, propidium iodide; PS, phosphatidylserine; PTP, permeability transition pore; ROS, reactive oxygen species; SOD, superoxide dismutase; VDAC, voltage-dependent anion channel; X + XO, xanthine plus xanthine oxidase.

Key words: superoxide anion; mitochondria; VDAC; cytochrome *c*; apoptosis

several apoptotic inducing factors, such as cytochrome *c* (cyto *c*), Smac/DIABLO, apoptosis-inducing factor (AIF), and caspases from the intermembrane space into the cytosolic environment in response to a variety of apoptotic stimuli (Liu et al., 1996; Adams and Cory, 1998; Green and Reed, 1998; Krajewski et al., 1999; Susin et al., 1999a,b; Du et al., 2000; Verhagen et al., 2000). During apoptosis, translocation of cyto *c* to the cytosol determines the assembly of a megaprotein complex (Apaf-1). The Apaf-1 complex initially results in activation of caspase-9 followed by the activation of other caspases. Released Smac/DIABLO interacts with the BIR domain of IAPs (inhibitors of apoptosis) and in turn relieves the inhibition of caspases by IAPs (for review see Green, 2000). Ultimately, caspases chop the cellular proteins resulting in programmed cell death.

Mitochondrial dysfunction is one of the prominent features of ROS-mediated cell death. Mitochondrial depolarization, in association with an increased endogenous production of superoxide anion and subsequent damage to the cardiolipids in the inner mitochondrial membrane, was reported as an early event in dexamethasone-induced apoptosis (Petit et al., 1995; Zamzami et al., 1995a,b). A major pathway leading to mitochondrial damage is based on the amplification of mitochondrial and cytosolic superoxide production in a broad spectrum of inflammatory or ischemia-related conditions. However, the exact mechanism by which ROS, and in particular $O_2^{\cdot-}$ -mediated apoptosis, establish and utilize CCR remains unclear. Recently, several models have been proposed to elucidate how cyto *c* and other apoptotic factors are released from mitochondria during apoptosis (for reviews see Green and Reed, 1998; Desagher and Martinou, 2000; Harris and Thompson, 2000; Korsmeyer et al., 2000; Kroemer and Reed, 2000). One paradigm suggests that apoptotic agents trigger mitochondrial permeability transition pore (PTP) opening that results in swelling of the mitochondrial matrix space, causing rupture of the outer mitochondrial membrane (OMM; Marchetti et al., 1996; Zamzami et al., 1996; Kroemer et al., 1998; Bernardi et al., 1999; Crompton, 1999). PTP opening has also been reported to be involved in initiation of the apoptotic machinery without irreversible damage of the mitochondrial membranes (Szalai et al., 1999). ROS and high intramitochondrial Ca^{2+} may act together to trigger PTP opening, and ROS in interaction with Ca^{2+} have been proposed to utilize PTP opening to evoke CCR and subsequent activation of caspases (for review see Zoratti and Szabo, 1995; Ankarcona et al., 1996; Marzo et al., 1998; Crompton, 1999; Duchon, 2000; Fiskum, 2000; Hajnóczky et al., 2000).

The voltage-dependent anion channel (VDAC, or mitochondrial porin) in the outer membrane and the adenine nucleotide translocator (ANT) in the inner membrane have also been proposed to control release of apoptotic factors without opening of the PTP complex. For example, closure of VDAC was reported to establish a defect in ATP/ADP exchange that results in an inhibition of the F_1F_0 -ATPase and, in turn, an initial hyperpolarization of the inner membrane followed by a loss of the outer membrane integrity and CCR (Vander Heiden et al., 1999).

Other models suggest that release of apoptotic factors occurs via large pores assembled from Bcl-2 family proteins (Bax, Bak) in the mitochondrial outer membrane (for review

see Green and Reed, 1998; Desagher and Martinou, 2000; Korsmeyer et al., 2000). Insertion of Bax to the outer membrane and formation of Bax oligomers may provide channels conducting large proteins (Gross et al., 1998; Basanez et al., 1999; Antonsson et al., 2000; Saito et al., 2000). Translocation of tBid from cytosol to the mitochondria have also been proposed to induce an allosteric activation and oligomerization of Bak, forming a pore that allows transport of cyto *c* (Wei et al., 2000). Bid and Bax may induce cyto *c* efflux through the OMM without affecting the inner mitochondrial membrane (IMM; von Ahsen et al., 2000). Antiapoptotic Bcl-2 family proteins have been shown to regulate mitochondrial ROS formation (Hockenbery et al., 1993; Kane et al., 1993; Gottlieb et al., 2000) and to inhibit ONOO⁻-induced CCR from isolated mitochondria distal to ONOO⁻ formation (Ghafourifar et al., 1999).

Interactions between Bcl-2 family proteins and components of the PTP (ANT, VDAC) have also been implicated in CCR (Marzo et al., 1998; Shimizu et al., 1999). Interaction with Bax has been reported to facilitate CCR via VDAC and this process was inhibited by Bcl-x_L in vitro (Shimizu et al., 1999, 2001). Based on these studies, a number of mechanisms may serve as a potential target of ROS during initiation of the mitochondrial phase of apoptosis. Since mitochondrial ATP production is needed for formation of the apoptosome, it is also necessary to determine whether a damage of the IMM causes impairment in oxidative metabolism before CCR.

Here, we aimed to delineate the pathway superoxide radicals utilize to evoke the mitochondrial and cellular phases of apoptosis. We examined the relationship between ROS formation and CCR, the loss of mitochondrial membrane potential ($\Delta\Psi_m$), caspase activation, and cell death in HepG2 hepatoma cells. We found that $O_2^{\cdot-}$, but not H_2O_2 , induces rapid and massive cyto *c* release from mitochondria. Although $O_2^{\cdot-}$ and Ca^{2+} acted in a coordinated manner in activation of PTP opening, $O_2^{\cdot-}$ -induced CCR was not affected by Ca^{2+} or by PTP blockers. In contrast, VDAC blockers or an anti-VDAC antibody (Ab#25) prevented cyto *c* release induced by $O_2^{\cdot-}$. Furthermore, integrity of the IMM was preserved during $O_2^{\cdot-}$ -mediated CCR. Bcl-2 family proteins did not appear to be essential for the control of $O_2^{\cdot-}$ -mediated CCR. Thus, this study provides evidence for a novel pathway that links $O_2^{\cdot-}$ to VDAC-dependent selective permeabilization of the OMM and CCR, and, ultimately leads to activation of the apoptotic cascade.

Results

$O_2^{\cdot-}$ and H_2O_2 facilitate Ca^{2+} -induced PTP opening, but only $O_2^{\cdot-}$ induces rapid release of cyto *c*

To examine the effect of $O_2^{\cdot-}$ (generated by xanthine plus xanthine oxidase; X + XO) and H_2O_2 exposure on PTP, $\Delta\Psi_m$, and $[Ca^{2+}]_c$ were monitored in permeabilized HepG2 cells incubated in the presence of extramitochondrial ATP supply. Since thapsigargin was added to abrogate endoplasmic reticulum Ca^{2+} uptake, changes in $[Ca^{2+}]_c$ reflected mitochondrial Ca^{2+} uptake and release. We have established a Ca^{2+} pulsing protocol that evoked modest depolarization and rapidly decaying $[Ca^{2+}]_c$ transients in naive cells,

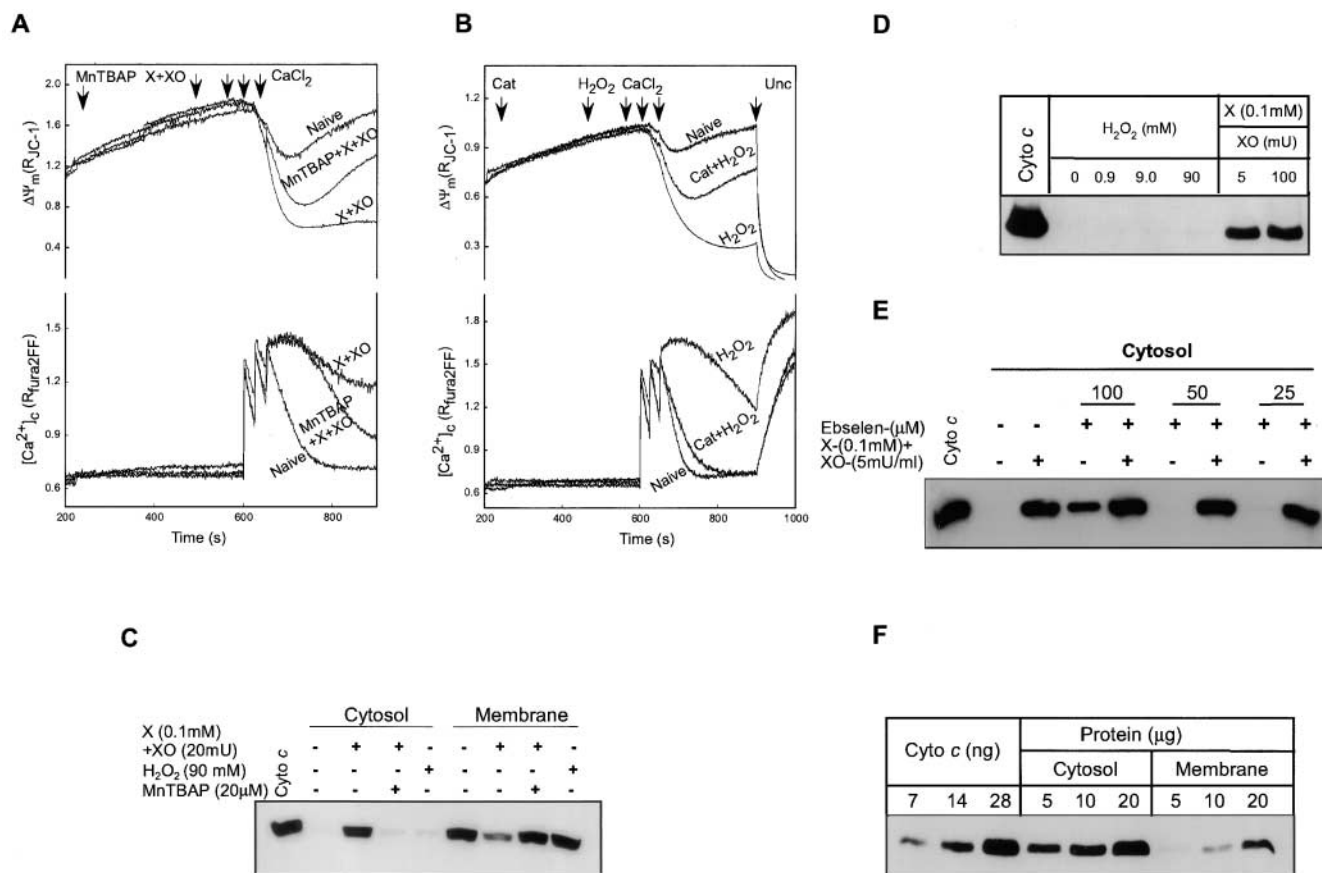


Figure 1. Effect of ROS on Ca²⁺-induced PTP opening and CCR in permeabilized HepG2 cells. (A) O₂⁻-generating system (xanthine [0.1 mM] plus xanthine oxidase [20 mU/ml]) and (B) H₂O₂ (90 mM) augmented Ca²⁺-induced depolarization (three pulses, 30 μM CaCl₂ each) and decreased mitochondrial Ca²⁺ uptake. These effects were inhibited by an O₂⁻ scavenger, MnTBAP (20 μM; 68 ± 4.5% decrease in depolarization and 78 ± 13% decrease in [Ca²⁺]_c rise at 900 s; n = 3), and catalase (Cat; 2500U/ml), respectively. At the end of the measurements, cells were exposed to FCCP (Unc; 1 μM), a protonophore that caused rapid and complete dissipation of ΔΨ_m. (C) Induction of CCR from mitochondria by O₂⁻ but not H₂O₂. Cytosolic and membrane fractions were prepared after fluorometric measurements of ΔΨ_m and [Ca²⁺]_c in permeabilized cells performed as shown in A and B, except that Unc was not added. Released cyto c in the cytosol was visualized by Western blotting with anti-cyto c antibody. (D) Permeabilized cells were incubated with H₂O₂ at three different concentrations (0.9, 9, and 90 mM) or with X + XO for 375 s. (E) Permeabilized cells were preincubated with an ONOO⁻ scavenger, ebselen, at three different concentrations (100, 50, and 25 μM), before the exposure to X + XO. (F) Quantification of O₂⁻-induced CCR from mitochondria. Cells were exposed to X + XO (0.1 mM plus 20 mU/ml) for 375 s. Various amounts of cytosolic and membrane fraction containing cyto c were compared with cyto c standard.

whereas large and extended depolarization and [Ca²⁺]_c elevation in cells treated with agents that promote the gating of PTP by Ca²⁺ (Szalai et al., 1999). As shown in Fig. 1 A, upon addition of X + XO the mitochondrial depolarization evoked by Ca²⁺ pulses was augmented and prolonged. Simultaneously, mitochondria in the X + XO-pretreated cells lost their capacity to keep sequestered the exogenously added Ca²⁺. Enhanced depolarization and impaired Ca²⁺ uptake exhibited by X + XO-treated cells reflected opening of PTP as it was inhibited by cyclosporin A (CsA; see Fig. 3 A). The mitochondrial depolarization and [Ca²⁺]_c rise in X + XO-treated cells were also attenuated and shortened by pretreatment with an O₂⁻ scavenger, superoxide dismutase mimic (Mn[III]tetrakis [4-benzoic acid] porphyrin, MnTBAP, 20 μM for 5 min; Fig. 1 A), suggesting that O₂⁻ formation promoted the activation of PTP by Ca²⁺.

Similar to O₂⁻, H₂O₂ potentiated the Ca²⁺ pulsing-induced depolarization and [Ca²⁺]_c elevation (Fig. 1 B). These effects of H₂O₂ were inhibited by an H₂O₂ scavenger,

catalase (Fig. 1 B) and by CsA (unpublished data; n = 2). This finding suggests that H₂O₂ also promoted Ca²⁺-dependent PTP opening.

At the end of fluorometric measurements, samples were centrifuged and the cytosol was separated from the membrane fraction. In the control, most of the cyto c was detected only in the mitochondrial pellet (Fig. 1 C). In X + XO-pretreated cells (X + XO for 375 s and Ca²⁺ for 300 s), less cyto c was in the pellet and a large amount was observed in the cytosol. Importantly, in the presence of MnTBAP, X + XO did not cause cyto c redistribution from mitochondria to cytosol (Fig. 1 C). Similarly, pretreatment with SOD (500 U/ml) also prevented the CCR from mitochondria (unpublished data, n = 2). In contrast to O₂⁻, H₂O₂ (0.9–90 mM) did not elicit cyto c rise in the cytosol (Fig. 1, C and D). O₂⁻ rapidly reacts with NO, yielding ONOO⁻, which has also been shown to evoke CCR and apoptosis (Lin et al., 1995; Ghafourifar et al., 1999). To evaluate the possibility of involvement of ONOO⁻ in X + XO-dependent CCR, per-

meabilized cells were pretreated with various concentrations of an ONOO⁻ scavenger, ebselen, and then exposed to O₂⁻. O₂⁻-induced CCR was not inhibited in the concurrent presence of ONOO⁻ scavenger (Fig. 1 E). Together, these results demonstrate that O₂⁻ and H₂O₂ promote Ca²⁺-dependent PTP opening, whereas O₂⁻, but not H₂O₂ or the O₂⁻ product, ONOO⁻, selectively induced CCR.

To quantitate O₂⁻-dependent CCR, various amounts of cytosolic and membrane proteins were resolved on SDS-PAGE. The amount of CCR was compared with known quantities of cyto *c* standard. As shown in Fig. 1 F, the vast majority of mitochondrial cyto *c* was released to the cytosol in permeabilized cells exposed to X + XO (5 mU/ml) for 375 s. Thus, the loss of cyto *c* from mitochondria in response to O₂⁻ treatment is rapid and almost complete.

O₂⁻-induced CCR is independent of PTP opening

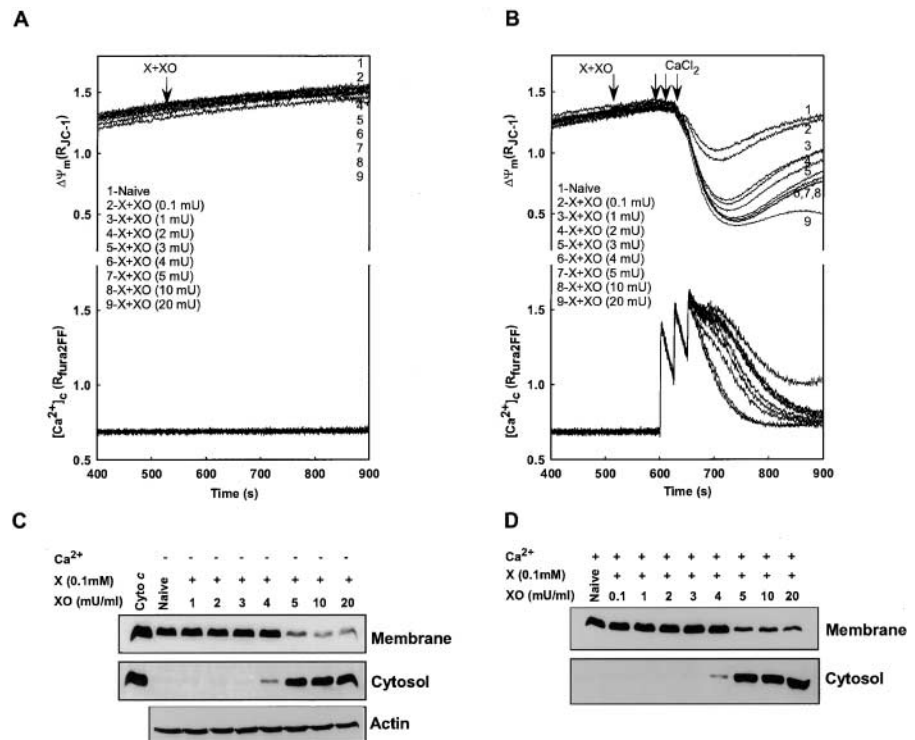
To investigate the mechanism by which O₂⁻ induces CCR, simultaneous measurements of ΔΨ_m and [Ca²⁺]_c were performed at various concentrations of X + XO in the absence and presence Ca²⁺ pulses (Fig. 2). In the absence of Ca²⁺, mitochondrial depolarization was not detectable during 375 s exposure to O₂⁻-generating system (Fig. 2 A). When Ca²⁺ was added to the permeabilized cells, depolarization was seen in a dose-dependent manner (Fig. 2 B). To determine cyto *c* distribution, cytosol and membrane fractions were separated as described above. Interestingly, O₂⁻ was able to induce CCR in a dose-dependent manner, even in the absence of Ca²⁺ addition (Fig. 2 C). CCR was observed at ≥4 mU XO and it was almost complete. Addition of Ca²⁺ did not affect the sensitivity of CCR to X + XO. Collectively, these data show that Ca²⁺ is not required for O₂⁻-dependent CCR.

To determine whether opening of PTP was involved in O₂⁻-induced CCR, permeabilized cells were treated with inhibitors of the inner membrane components of the PTP, CsA (1 μM), or bongkreikic acid (BA; 10 μM) before the addition of X + XO and Ca²⁺ pulses (Fig. 3). As shown in Fig. 3 A, CsA that binds to cyclophilin D in the mitochondrial matrix prevented the Ca²⁺-induced depolarization and sustained [Ca²⁺]_c rise in X + XO-treated permeabilized cells. By contrast, CsA failed to affect the CCR evoked by X + XO both in the presence or absence of Ca²⁺ (Fig. 3 B). Furthermore, BA, a blocker of the adenine nucleotide translocator that spans the IMM, was also ineffective to attenuate O₂⁻-induced CCR (Fig. 3 C). ADP supports the binding and inhibitory effect of BA, but BA also failed to inhibit O₂⁻-induced CCR when the ATP regenerating system was omitted from the medium to prevent extramitochondrial phosphorylation of ADP (Fig. 3 C, right). Together, these results show that opening of PTP is not required for O₂⁻-induced CCR.

Preservation of ΔΨ_m requires extramitochondrial ATP supply in cells exposed to O₂⁻

Following release of most of the mitochondrial cyto *c* (Fig. 2), oxidative metabolism is likely to be impaired. Thus, ΔΨ_m could be maintained in the presence of X + XO in expense of the extramitochondrial ATP supply (Fig. 2 A). To test whether reversed operation of the F₀F₁-ATPase allowed extrusion of H⁺ in mitochondria exposed to O₂⁻, we added an inhibitor of the F₀F₁-ATPase, oligomycin, and monitored the effect of X + XO on ΔΨ_m. Under these conditions X + XO caused rapid and complete depolarization (Fig. 4 A). CsA did not affect the X + XO-induced depolarization in oligomycin-pretreated cells (Fig. 4 A). Notably, pretreatment with oligomycin did not change the amount of CCR induced by O₂⁻ (unpublished data).

Figure 2. Ca²⁺-induced mitochondrial depolarization is not essential for O₂⁻-dependent CCR. Permeabilized cells were treated with X (0.1 mM) and different concentrations of XO (0.1, 1, 2, 3, 4, 5, 10, or 20 mU/ml) before addition of Ca²⁺ pulses (30 μM each) or solvent. (A and B) ΔΨ_m measured simultaneously with [Ca²⁺]_c. (C and D) cyto *c* in the cytosol and membrane fractions. Immunoblotting of actin in the cytosolic fraction was used to evaluate whether O₂⁻ changed the distribution of nonmitochondrial proteins.



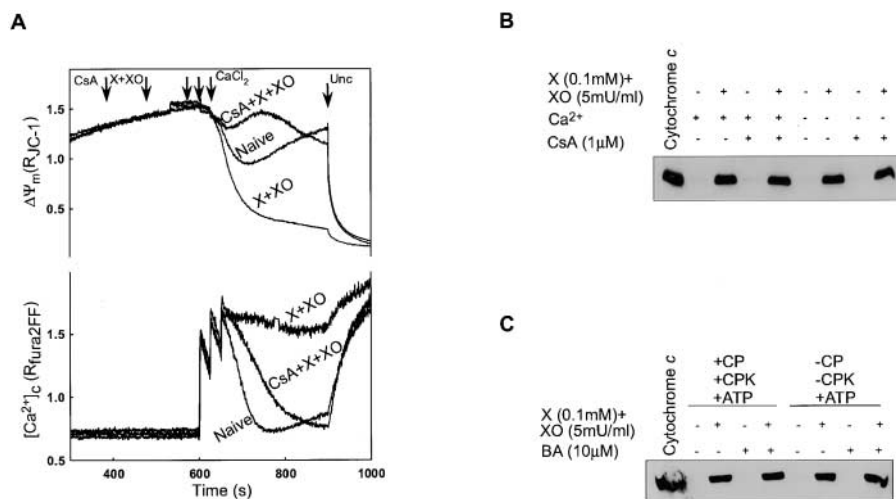


Figure 3. PTP blockers fail to inhibit O₂⁻-induced CCR. Permeabilized cells were pretreated with CsA (1 μM) or BA (10 μM) before the addition of X + XO. (A) Mitochondrial depolarization and [Ca²⁺]_c rise evoked by Ca²⁺ pulses added after the pretreatment with X + XO or CsA + X + XO. (B and C) Released cyto c in the cytosol was visualized by Western blotting. Cytosol samples were prepared after the measurement of ΔΨ_m but FCCP was not added. Permeabilized cells were pretreated with BA (10 μM) in the presence or absence of ATP-regenerating system before the X + XO treatment. Because ADP supports the binding and inhibitory effect of BA, to prevent extramitochondrial phosphorylation of ADP, ATP-regenerating system was omitted from the medium in some experiments.

O₂⁻-induced CCR precedes the ΔΨ_m loss

To clarify the temporal relation between CCR and ΔΨ_m loss, the time-course of CCR and depolarization were studied after exposure to X + XO in oligomycin-pretreated permeabilized cells. As shown in Fig. 5 A, CCR in the cytosol was apparent at 125 s and maximum release was observed at 225 s after X + XO treatment. CCR was associated with a decrease in ΔΨ_m that started >200 s after O₂⁻ addition (Fig. 4 A). Our results suggest that the loss of ΔΨ_m follows the O₂⁻-induced rapid release of cyto c instantaneously.

If depolarization was due to cyto c depletion-induced impairment of ΔΨ_m generation by the oxidative pathway, supply of exogenous cyto c to the bulk cytosolic buffer may sustain ΔΨ_m in oligomycin-treated cells (Waterhouse et al., 2001). As shown in Fig. 4 B, exogenously added cyto c (2, 5, and 10 μM) attenuated O₂⁻-induced ΔΨ_m loss in a dose-dependent fashion. These results show that after O₂⁻-induced CCR, mitochondria can maintain ΔΨ_m only if ATP was provided by an extramitochondrial source or the decrease in the intramitochondrial concentration of cyto c was compensated by added cyto c. Also, these results suggest that the IMM was intact even after O₂⁻-induced CCR.

Inhibitors of VDAC abrogate O₂⁻-induced CCR

Recent studies have reported that Bax, a proapoptotic Bcl-2 family protein may interact with VDAC to trigger CCR. It is also to note that the antiapoptotic protein, Bcl-x_L prevents CCR from mitochondria through the VDAC-Bax complex (Shimizu et al., 1999). The OMM contains a large amount of VDAC, and this protein channel can act as a track for the transport of substances in and out of mitochondria (Colombini et al., 1996). Thus, we next investigated the effect of VDAC blockers, 4,4'-diisothiocyanatostilbene-2,2'-disulfonic acid (DIDS) (Shafir et al., 1998) and König's polyanion (KP), on the O₂⁻-induced CCR. Pretreatment with DIDS (50 μM) for 1 min before the addition of X + XO completely inhibited O₂⁻-induced CCR (Fig. 5 A). Permeabilized cells were also pretreated with different concentrations of KP for 100 s before exposure to X + XO. Treatment with KP also inhibited CCR induced by O₂⁻ (Fig. 5 B). At higher concentrations, the KP caused release of a frac-

tion of cyto c by itself. These results indicated that VDAC is involved in the O₂⁻-induced CCR.

Recently, Shimizu et al. (2001) described an anti-VDAC antibody (Ab#25) that inhibits VDAC activity. We used this antibody to further verify the role of VDAC in CCR

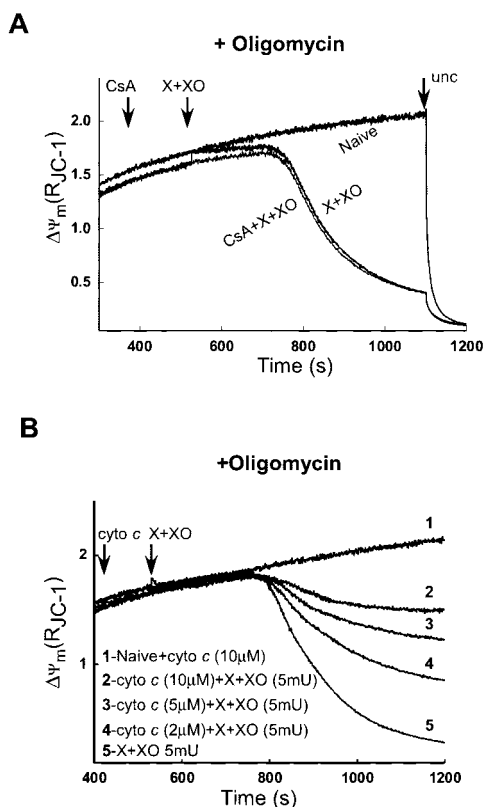


Figure 4. O₂⁻-induced depletion of mitochondrial cyto c yields mitochondrial depolarization in the absence of extramitochondrial ATP supply. (A) Effect of O₂⁻ on ΔΨ_m was monitored in permeabilized cells pretreated with oligomycin (2.5 μg/ml), an inhibitor of the F₀F₁-ATPase before the addition of X + XO or CsA + X + XO. (B) Rescue of mitochondria from O₂⁻-induced depolarization by exogenously added cyto c. Permeabilized cells were preincubated with cyto c (2, 5, or 10 μM) and oligomycin (2.5 μg/ml) before exposure to X + XO.

and $\Delta\Psi_m$ loss of permeabilized cells treated with X + XO. In permeabilized cells pretreated with oligomycin, X + XO showed massive amount of CCR and large $\Delta\Psi_m$ loss. However, cells pretreated with Ab#25 (0.56 $\mu\text{g}/\mu\text{l}$) for 5 min prevented the CCR (Fig. 5 D) and the $\Delta\Psi_m$ loss (Fig. 5 C).

Involvement of proapoptotic Bcl-2 family proteins is not necessary for $\text{O}_2^{\cdot-}$ -induced CCR

The interaction of Bcl-2 family proteins in the mitochondrial membranes can control the permeability to cyto *c*. During apoptosis, translocation of Bax from cytosol to mitochondria facilitates the release of cyto *c* through the OMM (Eskes et al., 1998; Jurgensmeier et al., 1998; Marzo et al., 1998; Narita et al., 1998; Pastorino et al., 1998). Bax has also been reported to interact with VDAC to release cyto *c* (Shimizu et al., 1999). However, in our study, cytosolic Bax did not translocate to the mitochondrial membrane during $\text{O}_2^{\cdot-}$ exposure (Fig. 5 E). By contrast, cells exposed to a proapoptotic agent, staurosporine (1 μM) for 3 h showed significant amount of mitochondrial anchored Bax. Furthermore, washout of the cytosol after cell permeabilization did not prevent $\text{O}_2^{\cdot-}$ -mediated CCR (unpublished data; $n = 3$). These data indicate that $\text{O}_2^{\cdot-}$ -induced CCR through VDAC does not require translocation of Bax to the mitochondria.

Another proapoptotic Bcl-2 family protein, Bak, may substitute for Bax in the CCR (Wei et al., 2001). However, no Bak was detected in the lysate of HepG2 cells using a monoclonal antibody (anti-human Bak, clone TC102; Oncogene Research Products) that recognized Bak in the lysate of H9c2 cardiac myotubes (unpublished data). Thus, Bak does not appear to be necessary for the $\text{O}_2^{\cdot-}$ -induced CCR.

Coupling of caspase activation to $\text{O}_2^{\cdot-}$ -mediated CCR

Mitochondrial permeabilization and CCR into the cytosol are essential events for caspase activation (Liu et al., 1996). We next examined whether the $\text{O}_2^{\cdot-}$ -induced release of cyto *c* and, presumably, other mitochondrial factors, is associated with caspase activation. After a short exposure to $\text{O}_2^{\cdot-}$, cleavage of procaspase-3 was detected, suggesting caspase-3 activation (Fig. 6 A). The VDAC blocker DIDS completely blocked the $\text{O}_2^{\cdot-}$ -mediated procaspase-3 cleavage, whereas CsA or BA did not prevent $\text{O}_2^{\cdot-}$ -mediated caspase activation (Fig. 6 A). One may speculate that the disappearance of procaspase-3 was due to its degradation by a necrosis-stimulating protease. To evaluate formation of the active caspase-3 enzyme, we also performed measurements of the cleavage product of a fluorogenic caspase substrate, DEVD-AMC (Fig. 6 B). In extracts of X + XO-treated permeabilized cells, the cleavage of DEVD-AMC was augmented. This effect was abolished by DIDS, whereas CsA did not attenuate the X + XO-stimulated DEVD-AMC cleavage (Fig. 6 B). We also observed that a caspase inhibitor, Ac-DEVD-CHO (50 μM), did not affect CCR, but completely blocked caspase 3 activation (unpublished data; $n = 2$). Thus, $\text{O}_2^{\cdot-}$ resulted in caspase activation which was closely coupled to VDAC-dependent CCR.

$\text{O}_2^{\cdot-}$ evokes VDAC-dependent CCR in intact cells

To demonstrate whether $\text{O}_2^{\cdot-}$ can also trigger CCR in intact cells, HepG2 cells were treated with X + XO for various time

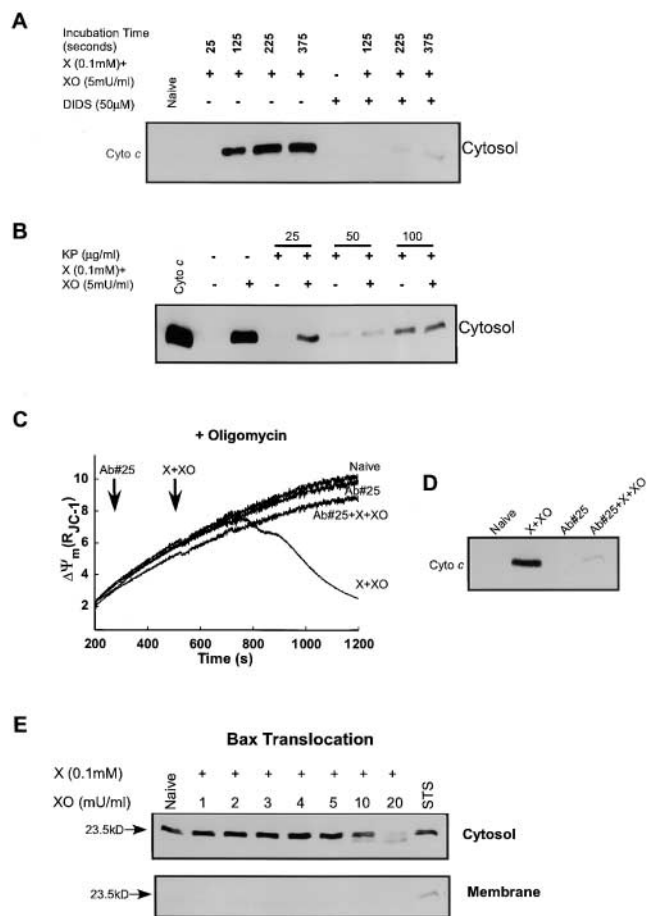


Figure 5. Inhibition of $\text{O}_2^{\cdot-}$ -induced CCR by VDAC blockers and anti-VDAC antibody. Permeabilized cells were exposed to X + XO in the absence of Ca^{2+} pulses. At various times after X + XO exposure, permeabilized cells were centrifuged for the collection of cytosolic fractions. For inhibitory studies, permeabilized cells were pretreated with (A) DIDS, an inhibitor of VDAC, for 50 s before the addition of X + XO, or (B) with indicated concentrations of KP for 100 s before exposure to X + XO. (C) Prevention of $\text{O}_2^{\cdot-}$ -induced $\Delta\Psi_m$ loss by anti-VDAC antibody (Ab#25). Permeabilized cells pretreated with or without Ab#25 (0.56 $\mu\text{g}/\mu\text{l}$) for 5 min, after which X + XO (0.1 mM plus 5 mU/ml) was added and changes of $\Delta\Psi_m$ were monitored for 12 min by spectrofluorimeter. Data presented is representative of one experiment. (D) Inhibition of $\text{O}_2^{\cdot-}$ -induced CCR by anti-VDAC antibody. After the measurement of $\Delta\Psi_m$ shown in C, the cells were centrifuged and cyto *c* was determined in the cytosolic fractions. (E) Translocation of Bax to the mitochondria is absent during exposure to $\text{O}_2^{\cdot-}$ but occurs during staurosporine-mediated CCR. The cytosol and mitochondrial fractions were subjected to 15% SDS-PAGE and transferred to nitrocellulose. Blots were probed with a polyclonal anti-Bax antibody.

periods. The time-course of the exposure to $\text{O}_2^{\cdot-}$ during treatment with X + XO was evaluated fluorimetrically, using a ROS tracer, 5-(and 6)-chloromethyl-2',7'-dichlorodihydrofluorescein diacetate (Molecular Probes). A substantial increase in ROS was noticed after 5 min and a plateau was reached after 60 min treatment with X + XO ($n = 2$). To study the cyto *c* distribution, after treatment with X + XO, cells were harvested, permeabilized and the cytosol was separated from the membrane fraction by centrifugation. Naive cells and cells treated with X + XO for a short duration (15 or 30 min) did not have cyto *c* in the cytosol (Fig. 7 A). Sometimes, we de-

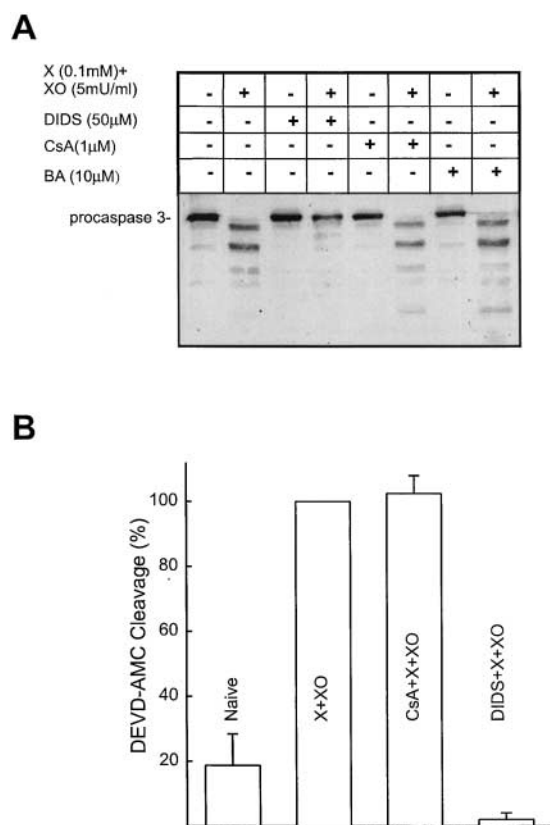


Figure 6. O₂⁻ induces caspase-3 activation; prevention by a VDAC blocker but not PTP inhibitors. (A) Cytosolic fractions (prepared as described above) were resolved on 15% SDS-PAGE followed by transfer to nitrocellulose and the blot was probed with polyclonal anticaspase-3 antibody. (B) Fluorometric assay of DEVD-AMC (12.5 μM) cleavage, in cytosol extracts (incubation for 30 min at 35°C). DEVD-AMC cleavage normalized to the activity obtained in the presence of X + XO (0.1 mM plus 5 mU/ml) is shown as mean ± SEM (*n* = 3).

ected CCR after 45 min but in minute quantities (unpublished data). Treatment with X + XO for 1 h or more caused large release of cyto *c* (Fig. 7 A). We wished to know whether the effects of PTP blockers or VDAC blocker on the O₂⁻-mediated CCR in intact cells were similar to those of in permeabilized cells. The O₂⁻-dependent release of cyto *c* into the cytosol was not detected in DIDS treated cells, whereas CsA or BA did not prevent mitochondrial CCR into the cytosol (Fig. 7 B). Thus, O₂⁻ produced in the bathing medium evoked CCR in intact cells similar to the effect of cytosolic O₂⁻ in permeabilized cells, but the lag time of the CCR was relatively long in intact cells. Several factors may contribute to the delayed onset of the X + XO-induced CCR in intact cells, including the scavenging of ROS in the bathing medium by BSA and in the cytosol by antioxidant mechanisms (e.g., Cu,Zn-SOD). Also O₂⁻ does not readily cross the plasma membrane.

To visualize the effect of O₂⁻ induced CCR in individual cells, we transfected HepG2 cells transiently with cyto *c*-green fluorescent protein (GFP). After transfection, cells were treated with X + XO (0.1 mM and 20 mU/ml) for appropriate time periods. The distribution of cyto *c*-GFP was visualized using a confocal microscope. In normal cells and at 30 min treatment with X + XO, cyto *c*-GFP was mostly

localized in mitochondria (Fig. 7 C). Cyto *c*-GFP appeared in the cytosol and in the nuclear matrix after 1 h or longer treatment with X + XO (Fig. 7 C). Cell to cell differences in the lag time of cyto *c*-GFP redistribution were observed, but once the release started it rapidly involved every mitochondrion. At 3 h of exposure to X + XO, cyto *c*-GFP distribution was homogeneous in most of the cells (Fig. 7 C). The O₂⁻-induced release of cyto *c*-GFP was quantitated based on the fluorescence appearing in the nuclear region of the cells (Fig. 7 D). The onset of cyto *c*-GFP release was similar to that of CCR, whereas the maximal release of cyto *c*-GFP was obtained 5 h after addition of X + XO (Fig. 7 D). Pretreatment with CsA (5 μM) did not inhibit cyto *c*-GFP release from mitochondria after O₂⁻ exposure. However, DIDS prevented the release of cyto *c* (Fig. 7 D). Collectively, these studies demonstrated that increased formation of O₂⁻ is capable of evoking CCR in intact cells and further supported the role of VDAC in this process.

Selective permeabilization of the OMM by truncated Bid has been described to initiate CCR in the absence of major damage of the mitochondrial structure (von Ahlsen et al., 2000). Using transmission electron microscopy, we evaluated the morphology of the mitochondria in intact cells incubated in the absence and presence of X + XO for 1 h (Fig. 7 E). Consistent with previous studies in HepG2 cells (e.g., Lewis et al., 1996), the majority of the mitochondria appeared in close association with the endoplasmic reticulum, was oval shaped, and was packed with cristae (left). Furthermore, 1 h of treatment with X + XO did not cause a major change in the mitochondrial structure (right). Thus, the onset of the CCR was not preceded by X + XO-induced large amplitude swelling of the mitochondria.

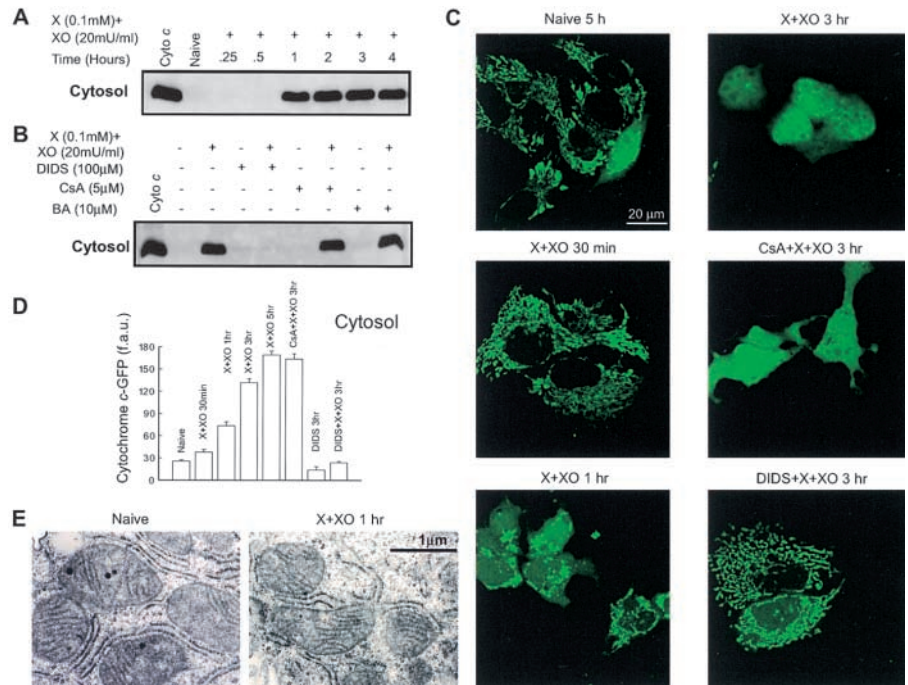
O₂⁻ evokes cyto *c* efflux from liposomes reconstituted with VDAC

To further explore the role of VDAC in O₂⁻-mediated apoptosis, we examined the effect of O₂⁻ on the permeability of VDAC purified (confirmed by Coomassie staining and immunoblotting; Fig. 8 A) from liver mitochondria and incorporated into liposomes. Plain liposomes and VDAC liposomes loaded with cyto *c*-FITC were exposed to O₂⁻ for 7 min, and the distribution of cyto *c*-FITC was visualized using confocal microscopy (Fig. 8 B). Plain, cyto *c*-FITC-loaded liposomes displayed only a small change in fluorescence distribution in response to X + XO (top images; extraliposomal fluorescence: 6.4 ± 0.6 and 11.4 ± 0.3 arbitrary units before and after X + XO addition, respectively; *n* = 3). By contrast, VDAC reconstituted liposomes allowed for a large increase of extraliposomal fluorescence in response to O₂⁻ exposure (middle images; 6.9 ± 0.2 and 33.3 ± 0.7 arbitrary units before and after X + XO addition, respectively; *n* = 3). Involvement of VDAC in the O₂⁻-induced CCR was further tested using DIDS. Pretreatment with DIDS (100 μM) inhibited the X + XO-induced increase in fluorescence between VDAC liposomes (bottom images; 7.6 ± 0.3 and 15.5 ± 0.5 arbitrary units before and after X + XO addition, respectively; *n* = 3).

To quantitate cyto *c*-FITC release from the liposomes, we also centrifuged liposome suspensions and measured the fluorescence in the supernatant and in the pellet using a flu-

Figure 7. $O_2^{\cdot-}$ -induced CCR in intact HepG2 cells; time-course, effect of PTP, and VDAC blockers.

(A) Cells were grown in 25-cm² flasks for 48 r and treated with X + XO for different time periods. After treatment, cells were harvested and permeabilized with 40 μ g/ml digitonin containing ICM at pH 7.2 for 10 min. The cytosol was then separated from the membrane fraction (containing mitochondria) by centrifugation. The cytosolic fractions were resolved on SDS-PAGE and immunoblotted for cyto *c*. (B) Cells pretreated with DIDS, CsA, or BA for 10 min were exposed to X + XO. After 3 h, cells were isolated, permeabilized, centrifuged, and immunoblot was done as described above. (C) Confocal imaging of $O_2^{\cdot-}$ -induced CCR using cyto *c*-GFP. Cyto *c*-GFP-transfected HepG2 cells were treated with X + XO (0.1 mM and 20 mU/ml, respectively) in the absence or presence of inhibitors. Distribution of cyto *c*-GFP was imaged by confocal microscopy. The images were taken after different time periods of X + XO exposure with a 60 \times oil objective. (D) Quantification of cyto *c*-GFP in the cytosol. Since the nucleus is devoid of mitochondria and released cyto *c*-GFP entered the nuclear matrix, the amount of cyto *c*-GFP released into the cytosol was determined by measurements of fluorescence over the nucleus in each cell. f.a.u., fluorescence arbitrary units. To assess the amount of cyto *c*-GFP release after X + XO treatment, we counted \sim 100–200 cells in three independent experiments for each condition. (E) Electron micrographs of control (left) and X + XO-treated (0.1 mM plus 20 mU/ml for 1 h; right) intact cells are shown to illustrate the typical patterns of mitochondrial morphology ($>$ 200 mitochondria were evaluated in each condition).



orimeter. As shown in Fig. 8 C, the fluorescence in the supernatant was small when liposomes were prepared without cyto *c*-FITC loading and reconstitution of VDAC, exposure to X + XO or DIDS did not evoke any changes (bars: 1, 3, 5, 7, and 9). When cyto *c*-FITC-loaded plain liposomes were exposed to X + XO, the fluorescence in the supernatant showed a twofold increase, similar to the data obtained with confocal imaging (bar 4 vs. 2). This increase was due to a direct interaction between the fluorophore and $O_2^{\cdot-}$, because a comparable increase was obtained when FITC was incubated with X + XO (unpublished data). However, the supernatant prepared from suspensions of VDAC liposomes exhibited eightfold increase after exposure to X + XO (bar 8 vs. 6). Comparing the plain and VDAC liposomes, incubation with X + XO caused a much larger increase of the fluorescence in the supernatant (bar 8 vs. 4) and this was accompanied with a much larger decrease of the fluorescence in the pellet prepared from VDAC liposomes (unpublished data, $n = 2$). In addition, X + XO-induced redistribution of the fluorescence was significantly attenuated by pretreatment with DIDS (bar 10 vs. 8). Importantly, Bax was not detected in purified VDAC or in VDAC liposomes, providing further evidence that the effect of $O_2^{\cdot-}$ on VDAC was not mediated by Bax (Fig. 8 A). Together, these studies show that reconstitution of VDAC in liposomes allows for a $O_2^{\cdot-}$ -induced increase in cyto *c* permeability. The lack of $O_2^{\cdot-}$ -dependent efflux of cyto *c*-FITC from plain liposomes suggests that $O_2^{\cdot-}$ does not rupture the lipid domains of the mitochondrial outer membrane. These results further confirmed that $O_2^{\cdot-}$ modulates opening of VDAC and facilitates CCR. The fact that VDAC blockers and anti-VDAC

antibody prevent the CCR in the HepG2 cells and that PTP blockers did not prevent $O_2^{\cdot-}$ -induced CCR suggested that the ability of $O_2^{\cdot-}$ to interact with VDAC is sufficient to induce CCR from mitochondria.

Execution of apoptosis in cells exposed to $O_2^{\cdot-}$

We have demonstrated CCR and caspase activation in cells exposed to $O_2^{\cdot-}$. To determine whether CCR and caspase activation are followed by ordered execution of the apoptotic program, phosphatidylserine (PS) exposure was evaluated by annexin V staining, and membrane permeability was studied by labeling with propidium iodide (Fig. 9). In X + XO-treated cells, imaging of $\Delta\Psi_m$ simultaneously with Alexa 488 annexin showed an early mitochondrial depolarization (Fig. 9 A; decrease in red signal at 1 h). Presumably, extramitochondrial ATP production is relatively small in HepG2 cells and therefore mitochondrial depolarization was closely coupled to cyto *c* release. After 1 h of X + XO treatment, very few cells exhibited annexin labeling of the plasma membrane, whereas at 3 h most of the cells were positive (green signal). Annexin staining of the cell could occur due to permeabilization of the plasma membrane; however, in cyto *c*-GFP-expressing cells, the cyto *c*-GFP released from mitochondria to the cytosol did not leave the cells, suggesting that the plasma membrane barrier was preserved (Fig. 9 B). Also, propidium iodide was initially excluded from the PS-positive cells (Fig. 9 C, see below). Treatment with DIDS (100 μ M) attenuated the $O_2^{\cdot-}$ -induced $\Delta\Psi_m$ loss and annexin staining (unpublished data). These observations suggest that execution of the apoptotic program was initiated by $O_2^{\cdot-}$ through a VDAC-dependent signaling pathway.

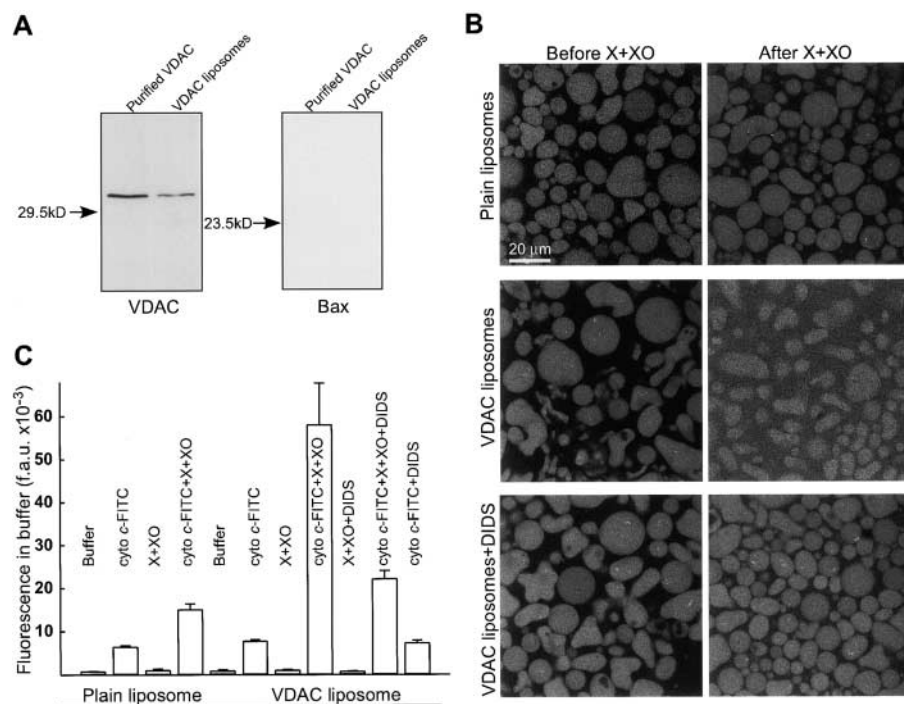


Figure 8. O₂^{•-} directly modulates purified VDAC to facilitate cyto c release. (A) Immunoblotting of purified VDAC and VDAC liposomes for VDAC and Bax. (B) Confocal imaging of FITC–cyto *c* efflux from plain and VDAC liposomes. Cyto *c* labeled with FITC was loaded into both plain and VDAC-incorporated liposomes. Images were taken before and after 7 min of exposure to X + XO (0.1 mM and 20 mU/ml). Furthermore, the effect of O₂^{•-} was evaluated on VDAC liposomes incubated in the presence of DIDS (100 μM). (C) Fluorometry of cyto *c*-FITC release from liposomes. Suspensions of liposomes (4 types; plain, plain loaded with cyto *c*-FITC, VDAC, and VDAC loaded with cyto *c*-FITC) were incubated in the absence or presence of X + XO (0.1 mM and 20 mU/ml) and DIDS (100 μM) for 7 min. Subsequently, the suspensions were centrifuged and FITC fluorescence was determined in the supernatants and liposomal pellets. Cyto *c*-FITC release is shown by the FITC fluorescence measured in the supernatants. We observed that

exposure to X + XO increased the fluorescence of FITC (unpublished data) and this effect could contribute to the increase of FITC in the supernatants of plain as well as VDAC liposomes. However, in response to X + XO, the VDAC liposomes exhibited a several-fold larger increase in the supernatant FITC and a considerably smaller residual pellet FITC than that of in plain liposomes (unpublished data). Data are from two separate experiments; each was performed in duplicates.

Apoptotic cells can be discriminated from necrotic cells based on the fact that apoptotic cells become permeable to propidium iodide (PI) relatively late. To evaluate the contribution of apoptosis and necrosis to cell death, after exposure to the O₂^{•-}-generating system for 1, 3, 5 and 7 h, incubation of the cells with Alexa 488 annexin and PI was performed. Most of the cells treated with O₂^{•-} for 1 h were not annexin V- or PI-positive. At 3 h of O₂^{•-} treatment, cells displayed mostly annexin V staining (i.e., early stage of apoptosis), whereas at 5 and 7 h pretreatment, cells showed both annexin V and PI staining, suggesting the onset of late stage apoptosis (Fig. 9 C). Thus, CCR and mitochondrial depolarization, which occurred at 1–2 h X + XO treatment (Figs. 7 A and 9 A), were followed by phosphatidylserine exposure and permeabilization of the plasma membrane was further delayed. This order of events suggests that the cells went through apoptotic cell death. This conclusion was further supported by the fragmentation of DNA in the cells exposed to O₂^{•-} (Fig. 9 D).

Discussion

This study describes a novel mechanism underlying activation of CCR by O₂^{•-}, but not by H₂O₂ or ONOO⁻. We showed that O₂^{•-}-induced CCR takes place quickly and involves most of the cyto *c* stored in the mitochondria. Although ROS interact with Ca²⁺ in gating of PTP, we have shown that the ROS-induced CCR pathway is independent of the inner membrane components of the pore. Our data suggest that the release of cyto *c* by O₂^{•-} is mainly due to VDAC-dependent selective permeabilization of the mito-

chondrial outer membrane. Ultimately, these alterations lead to activation of caspases and execution of apoptosis.

We examined the molecular machinery used by ROS to trigger the mitochondrial phase of apoptosis. The first important observation was that O₂^{•-} generated by the xanthine/xanthine oxidase system caused rapid release of cyto *c* from mitochondria to the cytosol in intact as well as in permeabilized HepG2 cells. Similar to cyto *c*, Smac/DIABLO was also rapidly released by O₂^{•-} (unpublished data). O₂^{•-}-induced release of cyto *c* and Smac/DIABLO may have a widespread significance in apoptosis, since in a number of pathological conditions, such as ischemia/reperfusion injury, drug insults, and inflammatory responses, large amounts of O₂^{•-} are produced by xanthine oxidase-mediated catabolism of purine nucleotides or by increased electron transport chain activity or by activation of NADPH oxidase particularly in neutrophils (Granger et al., 1986; Babior, 1999). Furthermore, increased formation of O₂^{•-} was described as an early event in several apoptotic paradigms (Petit et al., 1995; Zamzami et al., 1995a).

In the cells, O₂^{•-} can be converted to other ROS (H₂O₂, ONOO⁻), which have been reported to evoke CCR from isolated mitochondria (Yang and Cortopassi, 1998; Ghafourifar et al., 1999). However, in our studies, H₂O₂ (90 mM) did not establish rapid CCR and a ONOO⁻ scavenger, ebselen did not attenuate the O₂^{•-}-induced CCR. Thus, we conclude that O₂^{•-} triggered CCR by itself.

A principal mechanism of the CCR could involve ROS-induced promotion of Ca²⁺-dependent PTP opening. Permanent PTP opening causes CCR via matrix swelling and rupture of the mitochondrial membranes (for review see

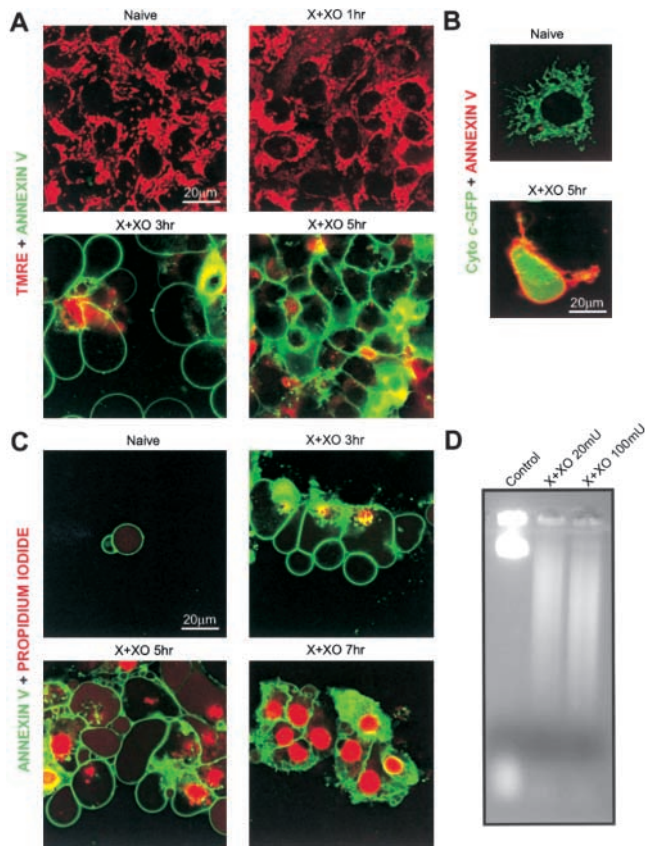


Figure 9. Execution of apoptosis in HepG2 cells exposed to $O_2^{\bullet-}$. Mitochondrial depolarization, PS exposure, disruption of plasma membrane integrity, and DNA fragmentation in cells exposed to $O_2^{\bullet-}$. Cells were treated with X + XO (0.1 mM plus 20 mU/ml) for various time periods in the absence or presence of DIDS (100 μ M). (A) $\Delta\Psi_m$ and PS exposure were evaluated with confocal imaging of annexin V Alexa 488 and TMRE. (B) Distribution of cyto *c*-GFP and annexin staining were visualized. (C) PS exposure and loss of plasma membrane integrity were evaluated with imaging of annexin V Alexa 488 and propidium iodide. (D) DNA fragmentation analysis was performed with cells incubated in the presence or absence of X + XO for 12 h.

Kroemer et al., 1998; Bernardi et al., 1999; Crompton, 1999; Desagher and Martinou, 2000), but reversible PTP opening was also shown to set up partial CCR without irreversible damage of mitochondrial function (Szalai et al., 1999). We also observed that Ca^{2+} -induced PTP opening was amplified in permeabilized cells exposed to $O_2^{\bullet-}$ or H_2O_2 , and simultaneous presence of either $O_2^{\bullet-}$ scavengers SOD or SOD mimic, MnTBAP, or H_2O_2 scavenger catalase attenuated this effect. However, $O_2^{\bullet-}$ -induced CCR was insensitive to inhibitors of PTP (CsA, BA) and was also observed in the absence of Ca^{2+} . Thus, PTP opening is not essential for $O_2^{\bullet-}$ -induced CCR. In addition, we demonstrated that $O_2^{\bullet-}$ -induced CCR does not yield $\Delta\Psi_m$ loss if the mitochondria were provided with exogenous cyto *c* or ATP. These data suggest that integrity of the mitochondrial inner membrane and matrix space was preserved during $O_2^{\bullet-}$ -induced CCR. It is important to note that in intact cells, mitochondrial depolarization was closely coupled to CCR, providing evidence that CCR caused impairment in the function of the electron transport chain and extramito-

chondrial ATP production could not maintain mitochondrial proton extrusion. Thus, $O_2^{\bullet-}$ -induced CCR is mediated by selective permeabilization of the OMM and the $O_2^{\bullet-}$ -induced mitochondrial depolarization was secondary to the loss of mitochondrial cyto *c*.

In regard to the changes in mitochondrial function, the origin of $O_2^{\bullet-}$ is likely to be of great significance. $O_2^{\bullet-}$ produced by the respiratory chain has been reported to cause cardiolipid destruction in the IMM and dissipation of the $\Delta\Psi_m$ (Zamzami et al., 1995a). However, based on our data, $O_2^{\bullet-}$ produced by X + XO in the cytosol or in the extracellular medium elicits selective permeabilization of the OMM and the damage of the IMM may occur after impairment of the OMM barrier. Presumably, slow permeation of $O_2^{\bullet-}$ through the OMM and the ROS scavengers in the mitochondria may attenuate exposure of the IMM to $O_2^{\bullet-}$ produced outside the mitochondria. Protection of the IMM may be particularly effective when a small amount of $O_2^{\bullet-}$ is produced, whereas a robust increase in extramitochondrial $O_2^{\bullet-}$ may cause an early damage of both the OMM and IMM and, in turn, necrosis (Higuchi et al., 1998). Along this line, HepG2 cells are expected to exhibit necrosis when exposed to a high dose of $O_2^{\bullet-}$. Interestingly, 100 mU XO plus 0.1 mM X was not sufficient to cause necrosis in all HepG2 cells, since DNA fragmentation was also noticed in this condition. Notably, cultured cells grown in 20% oxygen are essentially preadapted to conditions of oxidative stress (e.g., Davies, 1999).

Selective permeabilization of mitochondrial outer membrane may be established by pores formed by Bcl-2 family proteins (for review see Green and Reed, 1998; Desagher and Martinou, 2000; Korsmeyer et al., 2000) or by pores produced by the interaction of VDAC and Bax (Shimizu et al., 1999, 2001). Here, we showed that $O_2^{\bullet-}$ -induced CCR was inhibited by VDAC blockers (KP, DIDS) as well as by an antibody that interferes with the function of VDAC (#25; Shimizu et al., 2001). Furthermore, liposomes reconstituted with purified VDAC displayed $O_2^{\bullet-}$ -induced release of FITC-cyto *c*. Together, these data demonstrate that VDAC plays a central role in $O_2^{\bullet-}$ -induced CCR. In the studies of Shimizu et al. (1999), permeation of cyto *c* by VDAC liposomes required addition of Bax. By contrast, in our studies, $O_2^{\bullet-}$ -induced release of cyto *c* from the liposomes did not depend on the presence of Bax. The present study also showed that in normal HepG2 cells, the monomeric Bax form was present in the cytosol and translocated into the mitochondrial membrane upon staurosporine treatment. In contrast, cells exposed to the $O_2^{\bullet-}$ -generating system did not show any translocation of Bax into mitochondria, even at high concentrations of X + XO, but almost complete release of cyto *c* was still observed. On the same line, washout of the cytosol did not attenuate $O_2^{\bullet-}$ -induced CCR in the permeabilized cell experiments. Collectively, these data suggest that $O_2^{\bullet-}$ directly targets VDAC or a factor that is tightly coupled to VDAC in the process that established CCR and Bax does not mediate the effect of $O_2^{\bullet-}$ on VDAC.

In the cytosol, released cyto *c* can bind to Apaf-1 and upstream caspases to form a complex, where they are cleaved into their active forms (Zou et al., 1997). Activated caspase-9

can process inactive procaspase-3 into active caspase-3 and thereby triggers intracellular disintegration (Li et al., 1997). Our studies demonstrated that O₂⁻ triggered caspase-3 activation and similar to O₂⁻-induced CCR, caspase activation was not inhibited by PTP blockers (CsA or BA), whereas it was prevented by DIDS. In intact cells, O₂⁻-induced CCR was followed by mitochondrial depolarization, PS exposure and finally the cells become propidium iodide positive. This sequence of events suggests that O₂⁻-induced execution of the apoptotic death program. The apoptotic process exhibits rapid progression and takes place in most of the cells exposed to O₂⁻, suggesting that O₂⁻ may play a fundamental role in apoptosis caused by enhanced ROS production.

Materials and methods

Cell culture and transfections

HepG2, human hepatocellular carcinoma cells were grown in minimum essential medium (MEM; GIBCO-BRL) supplemented with 10% fetal bovine serum, 100 U/ml penicillin and 100 µg/ml streptomycin, and 2 mM glutamine at 37°C in 5% CO₂. For imaging experiments, cells were plated onto poly-D-lysine-treated glass coverslips (2–3 × 10⁴/cm²) and were grown for 72 h. For cell suspension studies, cells were cultured for 4–5 d in 75-cm² flasks, and were harvested by brief treatment with trypsin-EDTA. For transient transfections, cells were transfected using plasmid DNA (cyto c-GFP; Heiskanen et al., 1999) keeping total DNA at 2 µg/ml and cationic lipid LipofectA-MINE 2000 according to the manufacturer's instructions. After 5 h, the cells were placed in a normal growth medium. Imaging was performed 48 h after transfection.

Measurement of ΔΨ_m and [Ca²⁺]_i in suspension of permeabilized cells

Cells were washed in ice cold Ca²⁺-free extracellular buffer containing 120 mM NaCl, 5 mM KCl, 1 mM KH₂PO₄, 0.2 mM MgCl₂, 0.1 mM EGTA, 20 mM Hepes-NaOH, pH 7.4. An equal amount of cells (9–10 × 10⁶ cells, 7 mg/ml protein) were resuspended and permeabilized with 40 µg/ml digitonin in 1.8 ml of intracellular medium (ICM) composed of 120 mM KCl, 10 mM NaCl, 1 mM KH₂PO₄, 20 mM Hepes-Tris, pH 7.2, supplemented with 1 µg/ml of each of antipain, leupeptin, and pepstatin. ICM was passed through a Chelex column before addition of protease inhibitors to lower the ambient Ca²⁺. All the measurements were performed in the presence of 2 mM MgATP, 2 mM succinate, and ATP-regenerating system composed of 5 mM phosphocreatine and 5 U/ml creatine kinase.

For the simultaneous measurements of ΔΨ_m and mitochondrial Ca²⁺ uptake, the permeabilized cells were supplemented with 800 nM JC-1 and 0.5 µM fura2FF/FA. Fluorescence was monitored in a multiwavelength excitation dual wavelength emission fluorimeter (Delta RAM, PTI) using 340 and 380 nm excitation and 535 nm emission for fura2FF, whereas 490 nm excitation/535 nm emission and 570 nm excitation/595 nm emission were used for JC-1. Cytosolic Ca²⁺ is shown as the excitation ratio (340 nm/380 nm) of fura2FF/FA fluorescence, whereas ΔΨ_m is shown as the ratio of the fluorescence of J-aggregate (570 nm excitation/595 nm emission) and monomer (490 nm excitation/535 nm emission) forms of JC-1 (Szalai et al., 1999). All the experiments were performed at 35°C with constant stirring.

Analysis of cyto c distribution and caspase-3 activation by immunoblotting

At the end of the ΔΨ_m and [Ca²⁺]_i measurements in permeabilized cell suspensions, cells were immediately centrifuged at 14,000 g for 10 min and the supernatant collected as cytosol. The amount of protein present in the cytosol was measured using the Bradford method.

In studies with intact HepG2 cells, cells were grown in 25-cm² flasks for 48 h and were later treated with either xanthine 0.1 mM plus xanthine 20 mU/ml or in combination with DIDS (100 µM), CsA (5 µM), and BA (10 µM). At appropriate time periods, cells were harvested and washed in PBS. Cells were then resuspended in and permeabilized with 40 µg/ml digitonin in 1.0 ml of ICM supplemented with 1 µg/ml of each of antipain, leupeptin, and pepstatin for 15 min. The cytosol (14,000 g supernatant) and mitochondria (pellet) were stored at -80°C for Western blot analysis.

For Western blot experiments, proteins (20–25 µg) were loaded onto each lane of a 15% SDS-PAGE and electrophoretically transferred to nitro-

cellulose filters. Filters were blocked with blocking buffer (Pierce Chemical Co.) overnight, followed by incubation with anti-cyto c (7H8.2C12; BD PharMingen) and anticaspase-3 antibody (CPP32, YAMA, APOPAIN; BD PharMingen) at a dilution of 1:500 and 1:1,000, respectively. After incubation with the primary antibody, bound antibodies were visualized using horseradish peroxidase-coupled secondary antibody and Dura Signal chemiluminescence-developing kit (Pierce Chemical Co.).

Fluorimetric measurement of caspase activity

DEVD-AMC cleavage was measured in cell extracts as described previously (Szalai et al., 1999).

DNA fragmentation analysis

Cells grown in 100-mm culture dishes (3 × 10⁷ cells) were collected in their culture medium, pelleted at 125 g for 5 min, and pellets were washed twice with ice-cold PBS. Pellets were digested with 400 µl of digestion buffer (100 mM NaCl, 100 mM Tris-Cl, pH 8.0, 25 mM EDTA, pH 8.0, 0.5% SDS, 20 µg/ml DNase-free RNase final concentration) for 60 min at 37°C. These homogenates were further incubated with proteinase K (0.1 mg/ml) for 15 min at 50°C. Supernatants were collected to a new tube, in which they were extracted with phenol:chloroform:isoamyl alcohol (25:24:1). DNA was precipitated with 2 vol of cold ethanol and DNA was recovered by centrifugation at 500 g for 2 min. Precipitated DNA was washed once with 70% ethanol and resuspended in 1 mM EDTA, 10 mM Tris-HCl, pH 8.0. Isolated genomic DNA was resolved on a 1.5% agarose gel and visualized by ethidium bromide staining.

Purification and incorporation of VDAC into liposomes and confocal imaging of cyto c translocation

Mitochondria isolated from rat liver were used for purification of VDAC as described previously (de Pinto et al., 1987). The purity of VDAC was assessed by SDS-PAGE electrophoresis and confirmed by Western blotting (antiporin 31HL antibody Calbiochem). Liposomes were prepared by sonicating 500 mg of phospholipid mixture (soybean Type II-S) in 10 ml of buffer containing 50 mM KCl, 20 mM KH₂PO₄, 20 mM Hepes and 1 mM EDTA, pH 7.0. After sonication, incorporation of purified VDAC and loading of cyto c-FITC (cyto c-FITC was purified as described in the labeling kit protocol [Molecular Probes] and purity was assessed by phosphor imager and Western blotting) was performed by subjecting these vesicles for two freeze-thaw cycles. For imaging of cyto c efflux, fluorescence of cyto c-FITC was monitored using a confocal system (BioRad-1024/2P, excitation 488 nm) fitted to an Olympus IX70 inverted microscope (60× objective). Images were taken from plain and VDAC incorporated liposomes before and 7 min after addition of X (100 µM) and XO (20 mU/ml). For quantitative analysis, liposomes were centrifuged and FITC was released to the supernatant and residual FITC in the liposomal fraction were determined fluorometrically (490 nm excitation/510 nm emission).

Electron microscopy

Naive and X + XO (0.1 mM plus 20 mU/ml) -treated cells were fixed in 2% glutaraldehyde, 1% tannic acid, 0.1 M sodium cacodylate at pH 7.4 for 2 h at room temperature. Processing, embedding of the fixed cells, cutting, staining, and examination of the sections (80-nm thick) were performed as described recently (Pacher et al., 2000).

Confocal imaging analysis of CCR and apoptotic markers in intact individual cells

For imaging of cyto c distribution, cells transfected with cyto c-GFP construct were treated with either X (0.1 mM) plus XO (20 mU) or in combination with DIDS (100 µM) or CsA (1 µM). After 3 h, cells were washed twice with extracellular medium (2.0% BSA/ECM) consisting of 121 mM NaCl, 5 mM NaHCO₃, 10 mM NaHepes, 4.7 mM KCl, 1.2 mM KH₂PO₄, 1.2 mM MgSO₄, 2 mM CaCl₂, 10 mM glucose, and 2% BSA, pH 7.4. The coverslips were transferred to the microscope stage and cyto c-GFP distribution was monitored with confocal imaging (488 nm excitation).

For imaging of phosphatidylserine exposure and ΔΨ_m, HepG2 cells were exposed to X + XO for various time periods in the absence or presence of DIDS (100 µM), then incubated with annexin V Alexa 488 for 15 min using 1 × binding buffer (Molecular Probes). During the annexin incubation period, TMRE (50 nM) was also loaded (10 min) for the measurement of ΔΨ_m or propidium iodide (0.5 µg/ml) to evaluate the plasma membrane integrity.

Fluorometry recordings and confocal images are representative of three to six independent experiments. The data combined from separate experiments are shown as mean ± SE. Significance of differences from the relevant controls was calculated by Student's *t* test.

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