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FURTHER CHARACTERIZATION OF THE EVENTS INVOLVED IN MITOCHONDRIAL Ca²⁺ RELEASE AND PORE FORMATION BY PROOXIDANTS

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Abstract—Addition of the prooxidant 3,5-dimethyl-N-acetyl-p-benzoquinone imine (3,5(Me)₂NAPQI) to Ca²⁺-loaded mitochondria caused a rapid and extensive release of the sequestered Ca²⁺. Ca²⁺ release was accompanied by irreversible NAD(P)H oxidation and was followed by the release of adenine and pyridine nucleotides into the extramitochondrial medium; this is evidence of the opening of the pore in the inner mitochondrial membrane. Preincubation of the mitochondria with ADP, cyclosporin A (CSA), m-iodobenzylguanidine (MIBG) or Mg^{2+} inhibited the prooxidant-induced Ca^{2+} release and prevented pore-opening. When mitochondria were preincubated with ruthenium red, Ca^{2+} release was only minimally stimulated by 3,5(Me)2NAPQI. However, increasing the concentration of the prooxidant caused release of an increasing fraction of the sequestered Ca²⁺. Alternatively, increasing the intramitochondrial Ca²⁺ load resulted in a lowering of the concentration of 3,5(Me)₂NAPQI required for near complete Ca²⁺ release to occur. In the presence of ruthenium red, 3,5(Me)₂NAPQI-induced Ca²⁺ release was accompanied by irreversible pyridine nucleotide oxidation and followed by the release of nucleotides into the extramitochondrial medium, events which were prevented on preincubation with CSA. Similarly, the addition of CSA, ADP or MIBG during 3,5(Me)₂NAPQI-induced Ca²⁺ release arrested further Ca2+ release. In addition to their inhibitory effect on the 3,5(Me)2NAPQI-induced Ca²⁺ release, CSA, ADP or MIBG also decreased the rate of the basal, ruthenium red-induced mitochondrial Ca2+ release by 45-70%. It is proposed that the basal, ruthenium red-induced and the prooxidant-induced mitochondrial Ca²⁺ release occur through a common component that is sensitive to inhibition by CSA, ADP and MIBG and that is involved in mitochondrial pore formation. Furthermore, $3.5(Me)_2NAPQI$ -induced pore opening does not involve Ca^{2+} -cycling, but rather involves a site(s) that is (are) synergistically activated by Ca^{2+} and the prooxidant.

Key words: mitochondria; calcium; oxidative stress; pore; cyclosporin A; pyridine nucleotides

Intramitochondrial Ca²⁺ homeostasis is controlled by separate Ca²⁺-uptake and release pathways. Uptake of Ca²⁺ into the mitochondrial matrix occurs through a ruthenium red-sensitive uniporter and is driven by the electrochemical transmembrane potential generated during respiration [1]. Release of Ca²⁺ from mitochondria may involve several distinct pathways [2]. Thus, under physiological conditions efflux of Ca²⁺ occurs in exchange for Na⁺ or with H⁺, the relative contribution of H⁺ versus Na⁺ depending on the tissue of origin of mitochondria. However, evidence has been presented to suggest that under pathological conditions such as oxidative stress from exposure to prooxidants or during the reoxygenation of hypoxic tissue, excessive

The aim of the present study was to further investigate the early and critical event(s) triggering prooxidant-induced Ca²⁺ release. We show here that Ca²⁺ release induced by the prooxidant

mitochondrial Ca2+ uptake and subsequent Ca2+ loss occur [3-5]. Ca2+ release may then be a nonspecific event, possibly due to opening of a pore of about 20 Å diameter in the inner mitochondrial membrane [6-11]. Pore opening has been intensively investigated in isolated mitochondria, and been shown to require relatively high intramitochondrial Ca2+ together with an inducing agent, such as phosphate or the prooxidant tBuOOH† [6-11]. This pore is probably identical with the permeability transition pore observed in mitochondrial swelling experiments [12-18], and has also been detected in patch clamp experiments [19]. Pore opening is fully reversed on Ca²⁺ removal [8, 10] and is potently inhibited by the immunosuppressant cyclosporin A [9, 15, 16].

Recently, however, evidence has been presented to suggest that the prooxidant tBuOOH can also activate a selective pathway of Ca²⁺ efflux from mitochondria without pore opening if Ca²⁺ reuptake is prevented by ruthenium red or EGTA. Thus, in the case of tBuOOH, pore opening appears to be an event that is secondary to specific Ca²⁺ release and is caused by Ca²⁺-cycling [20].

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[†] Abbreviations: tBuOOH, tert-butyl hydroperoxide; Arsenazo III, 2,2'-(1,8-dihydroxy-3,6-disulphonaphthalene-2,7-bisazo)-bis-(benzene arsonic acid); Hepes, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid; Me₂SO, dimethylsulfoxide; TCA, trichloroacetic acid; MIBG, meta-iodobenzylguanidine; 3,5(Me)₂NAPQI, 3,5-dimethyl-N-acetyl-p-benzoquinone imine; 3,5(Me)₂ NAPQIH₂, 3,5-dimethyl-N-acetyl-p-aminophenol; FCCP, carbonyl cyanide-p-trifluoromethoxy-phenylhydrazone; ruthenium red, (Ru₃O₂(NH₃)₁₄)Cl₆·4H₂O.

3,5(Me)₂NAPQI, either in the absence or in the presence of ruthenium red, was inhibited by CSA, ADP, Mg²⁺ or MIBG. 3,5(Me)₂NAPQI-induced Ca²⁺ release was accompanied by pore opening, and pore opening was prevented by the same agents. CSA, ADP or MIBG also inhibited the basal, ruthenium red-induced Ca²⁺ release. We suggest that Ca²⁺ release observed under these different experimental conditions involves a common component that represents a structure involved in the mitochondrial pore. We furthermore show that 3,5(Me)₂NAPQI-induced pore opening does not require Ca²⁺-cycling, but is synergistically triggered by the intramitochondrial Ca²⁺ load and the oxidant-concentration.

MATERIALS AND METHODS

Chemicals. 3,5(Me)₂NAPQI was obtained from Dalton Chemical Laboratories Inc. (Toronto, Ontario, Canada). MIBG and CSA were gifts from Dr L. Smets (The Netherlands Cancer Institute, Amsterdam, The Netherlands) and Sandoz (Basel, Switzerland), respectively. ADP, glucose-6-phosphate dehydrogenase (EC 1.1.1.49) and glucose-6-phosphate were obtained from Boehringer (Mannheim, Germany). Arsenazo III (98% pure) and ruthenium red were purchased from the Sigma Chemical Co. (St Louis, MO, U.S.A.). Cellulosenitrate filters (0.45 μm pore size) were obtained from Sartorius (Göttingen, Germany). All other chemicals were of the highest purity commercially available.

Isolation of mitochondria. Liver mitochondria were isolated from male Wistar rats (ALAB AB, Sollentuna, Sweden; 200–250 g, fed ad libitum) as described previously [21]. The protein content of the final mitochondrial suspension was determined by the biuret method [22] using crystallized and lyophilized bovine serum albumin as standard.

Mitochondrial incubations. Unless stated otherwise, all incubations were performed in MSH buffer (mannitol (210 mM), sucrose (70 mM) and Hepes (3 mM), pH 7.1), at 25°. Mitochondria (1 mg protein/mL) were energized with succinate (5 mM) in the presence of rotenone (3 μM) for 2.5 min, before loading with Ca²+ (15 nmol/mg protein) for an additional 1.5 min. ADP, CSA, Mg²+ or MIBG was added 1 min after Ca²+, or subsequent to 3,5(Me)₂NAPQI as described in the text. Ruthenium red was added 2 min after Ca²+, and 3,5(Me)₂NAPQI was added (from concentrated stocks in dry Me₂SO prepared fresh every day) as described in the text. The final concentration of Me₂SO in the incubation mixtures never exceeded 0.3%.

Mitochondrial Ca²⁺ fluxes. Mitochondrial Ca²⁺ fluxes were measured by monitoring the change in absorbance of the metallochromophoric dye Arsenazo III (40 μ M) using the wavelength pair 654–685 nm. The change in absorbance of 40 μ M Arsenazo III linearly reflected the Ca²⁺ concentration up to a concentration of around 20 μ M of Ca²⁺.

Analysis of total and extramitochondrial pyridine nucleotides and adenine nucleotides. The mitochondrial pyridine nucleotide redox state was determined spectrophotometrically in a Shimadzu UV-3000 spectrophotometer using the wavelength

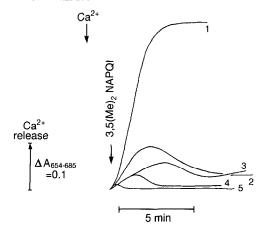


Fig. 1. Inhibitory effect of CSA, ADP, Mg²⁺ or MIBG on quinone imine-induced mitochondrial Ca²⁺ release. Mitochondria were loaded with Ca²⁺ and preincubated in the absence (trace 1) or presence of ADP (trace 2) (150 nmol/mg protein), MIBG (trace 3) (60 nmol/mg protein), Mg²⁺ (trace 4) (60 nmol/mg protein), or CSA (trace 5) (1.5 nmol/mg protein). The stimulation of Ca²⁺ release by 3,5(Me)₂NAPOI (90 nmol/mg protein, a concentration that resulted in near complete release of all sequestered Ca²⁺), was measured as described in Materials and Methods. For clarity, trace 4 was corrected for the spectral interference produced by the Arsenazo III–Mg²⁺ complex. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

pair 340-375 nm. In addition, the total and extramitochondrial adenine and oxidized pyridine nucleotide content was assayed by HPLC [23] as follows. Mitochondria (1 mg protein/mL) were energized and loaded with Ca2+ as described above. At selected time points, duplicate aliquots of 0.5 mL were then withdrawn from the mitochondrial incubations. One aliquot was directly added to $80 \mu L$ of 70% (v/v) PCA to determine total mitochondrial nucleotides. The other aliquot was filtered over a nitrocellulose filter into $80 \,\mu\text{L}$ of 70% (v/v) PCA to assay for released mitochondrial nucleotides. The samples were then processed for the analysis of pyridine nucleotides and adenine nucleotides as described in Ref. [23]. The nucleotides were separated using a CT-sil C₁₈ column (5 μ m, 250 × 4.6 mm) (Chrom Tech AB, Norsborg, Sweden) as reported previously [24].

In addition, extramitochondrial NADP⁺ was measured spectrophotometrically using an enzymatic assay without prior separation of the mitochondria from the incubation medium. In this case, mitochondria (2 mg protein/mL) were loaded with Ca²⁺ (15 nmol/mg protein) and 3,5(Me)₂NAPQI-induced oxidation of NAD(P)⁺ was allowed to proceed for 1.5 min before addition of CSA (1.5 nmol/mg) to cause pore closure. After NAD(P)⁺ re-reduction in the presence of CSA was complete, glucose-6-phosphate (250 µM) and glucose-6-phosphate dehydrogenase (0.4 U/mL) were added to the mitochondrial incubation to reduce accessible (extramitochondrial) NADP⁺.

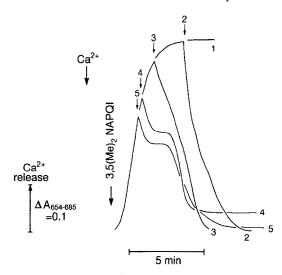


Fig. 2. CSA, ADP, $\mathrm{Mg^{2^+}}$ or MIBG induces reuptake of $\mathrm{Ca^{2^+}}$ by mitochondria after exposure to 3,5(Me)₂NAPQI. Mitochondria were loaded with $\mathrm{Ca^{2^+}}$ as described in Materials and Methods and release of all sequestered $\mathrm{Ca^{2^+}}$ was induced by the addition of 3,5(Me)₂NAPQI (90 nmol/mg protein) (trace 1). CSA (trace 2) (1.5 nmol/mg protein), ADP (trace 3) (150 nmol/mg protein), Mg²⁺ (trace 4) (60 nmol/mg protein) and MIBG (trace 5) (60 nmol/mg protein) were added where indicated by the arrows. $\mathrm{Ca^{2^+}}$ release and reuptake was measured as described in Materials and Methods. Note that the change in absorbance of Arsenazo III linearly reflected the $\mathrm{Ca^{2^+}}$ concentration up to around $20\,\mu\mathrm{M}$. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

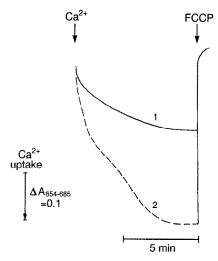


Fig. 3. Uptake of Ca^{2+} by control and 3,5(Me)₂NAPQI-treated mitochondria. Mitochondria (1 mg protein/mL) were preincubated with CSA (1.5 nmol/mg) for 1 min and were subsequently diluted to a final protein concentration of 0.2 mg/mL by addition of a corresponding volume of MSH-buffer containing 15 μ M Ca^{2+} and 40 μ M Arsenazo III (final concentrations). Ca^{2+} uptake was measured at 685–654 nm, and a typical rate is shown in trace 1. Trace 2 shows the rate of CSA-induced reuptake of Ca^{2+} (after near complete release) by 3,5(Me)₂NAPQI-treated mitochondria (1 mg protein/mL) using the same batch of mitochondria. FCCP (0.5 μ M) was added as indicated by the right arrow. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

RESULTS

Mitochondrial Ca²⁺ release in the absence of ruthenium red and induction of Ca²⁺ reuptake by CSA, ADP, Mg²⁺ or MIBG

When Ca2+-loaded mitochondria were incubated in the absence of ruthenium red, the addition of 3,5(Me)₂NAPQI (90 nmol/mg protein) induced a rapid release of all the sequestered Ca²⁺. Ca²⁺ release was inhibited by preincubating the mitochondria with either CSA, ADP, Mg²⁺ or MIBG (Fig. 1). In addition to their inhibitory effect on Ca²⁺ release when added before 3,5(Me)2NAPQI, these agents also induced complete reuptake of all the released Ca²⁺ by the mitochondria when added after exposure to 3,5(Me)₂NAPQI (Fig. 2). CSA or ADP stimulated the reuptake of all the released Ca²⁺, even after near complete discharge of the entire Ca²⁺ load from mitochondria (Fig. 2). By contrast, Mg²⁺ and MIBG induced total reuptake only when added before maximal Ca²⁺ release had occurred (Fig. 2). The observed reuptake suggests that the Ca²⁺ release pathway was operative in a reversible manner, and that ADP or CSA could induce its closure.

The criterion of reuptake as indicative of reversible opening of a Ca^{2+} release pathway has been questioned [2]. Since mitochondria can sequester large amounts of Ca^{2+} , the sequestration of all the released Ca^{2+} by a small subpopulation of mitochondria

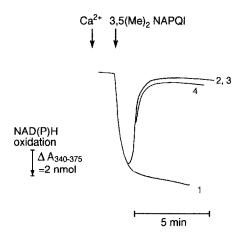


Fig. 4. Effect of CSA, ADP, or Mg²⁺ on 3,5(Me)₂NAPQI-induced mitochondrial pyridine nucleotide oxidation. Mitochondria were loaded with Ca²⁺ and preincubated in the absence (trace 1) or presence of CSA (trace 2) (1.5 nmol/mg protein), Mg²⁺ (trace 3) (60 nmol/mg protein) or ADP (trace 4) (150 nmol/mg protein) before the addition of 3,5(Me)₂NAPQI (90 nmol/mg protein). Pyridine nucleotide oxidation was measured as described in Materials and Methods. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

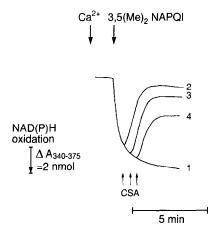


Fig. 5. Time-dependence of CSA-induced re-reduction of mitochondrial pyridine nucleotides when added after 3,5(Me)₂NAPQI. Mitochondria were loaded with Ca²⁺ as described under Materials and Methods, and exposed to 3,5(Me)₂NAPQI (90 nmol/mg protein) (trace 1). CSA (1.5 nmol/mg protein) was added 0.5 min (trace 2), 1 min (trace 3) or 1.5 min (trace 4) after 3,5(Me)₂NAPQI as indicated by the arrows. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

whose Ca²⁺ release pathway has not been activated has been suggested to be indistinguishable from the reuptake by the whole population on closure of the release pathway [2]. Therefore we investigated whether a small subpopulation of control mitochondria could accumulate Ca2+ at a rate comparable to that observed with 3,5(Me)₂NAPQI-treated mitochondria. Using a fraction that accounted for 20% of the total population, accumulation of Ca²⁺ in the presence of CSA was much slower than CSAinduced Ca²⁺ reuptake by 3,5(Me)₂NAPQI-treated mitochondria (Fig. 3). Virtually identical results were obtained with ADP (data not shown). Around 40-50\% of the total mitochondrial population were required to achieve a rate of Ca2+ uptake that was comparable to the rate of the CSA- or ADPinduced Ca²⁺ reuptake by 3,5(Me)₂NAPQI-treated mitochondria (data not shown). It is unlikely that, after near complete release of Ca2+ from the whole mitochondrial population, in 40-50% of the mitochondria the Ca²⁺ release pathway would not have been activated. Consequently, the CSA- or ADPinduced reuptake of Ca²⁺ cannot be explained solely by the heterogeneity of the mitochondrial suspension. In contrast with CSA and ADP, Mg2+ or MIBG could induce reuptake of Ca2+ only when added before near complete Ca2+ release had occurred, and in this case both explanations, i.e. inactivation of the release pathway and reuptake of Ca²⁺ by a subpopulation need to be considered.

Mitochondrial NAD(P)H oxidation and release of mitochondrial nucleotides

In agreement with previous results [24, 25], 3,5(Me)₂NAPQI-induced mitochondrial Ca²⁺ release was accompanied by an irreversible oxidation

of mitochondrial NAD(P)H. When mitochondria were preincubated with CSA, MIBG, ADP or Mg²⁺ before the addition of 3,5(Me)₂NAPQI, NAD(P)H oxidation was fully reversible (Fig. 4). By contrast, when ADP (data not shown) or CSA was added during Ca²⁺ release, a time-dependent decrease of the extent of NAD(P)+ re-reduction occurred (Fig. 5). HPLC analysis of the incubation medium revealed a time-dependent appearance of NAD⁺ and NADP⁺ in the extramitochondrial medium (Table 1). Only 40-50% of total NAD⁺ and NADP⁺ were detectable in the extramitochondrial medium at a time of near complete Ca²⁺ release (95% of total Ca²⁺ load). When mitochondria were preincubated with CSA before the addition of 3,5(Me)₂NAPQI, the release of NAD+ and NADP+ into the medium was prevented (data not shown). Thus, the levels of pyridine nucleotides retained within the mitochondria, as assessed by HPLC analysis (Table 1), closely corresponded to the degree of NAD(P)+ re-reduction observed by spectrophotometric analysis (Fig. 5).

For the measurements of extramitochondrial NAD(P)+ by HPLC, the separation of the mitochondria from the incubation medium by filtration under vacuum was required. To exclude the possibility that the filtration process might have ruptured some of the mitochondria, thereby causing a release of nucleotides into the filtrate, we also tested the extramitochondrial presence of NADP⁺ using a non-disruptive enzymatic method. The addition of glucose-6-phosphate dehydrogenase together with glucose-6-phosphate to Ca²⁺-loaded control mitochondria did not alter the absorbance at 375-340 nm (data not shown). However, when enzyme plus substrate was added to mitochondria undergoing 3,5(Me)₂NAPQI-induced Ca²⁺ release and pyridine nucleotide oxidation, an increase in the absorbance at 340-375 nm was observed (Fig. 6). Thus, mitochondrial NADP+ was accessible to externally added enzymes, confirming the data obtained by HPLC analysis of the filtrate. Estimation of extramitochondrial NAD+ by enzymatic (sorbitol dehydrogenase catalyzed) reduction proved difficult due to interference by an endogenous mitochondrial NADH oxidizing activity (data not shown).

In addition to pyridine nucleotides, adenine nucleotides were also present in the mitochondrial filtrate (Table 1). The relative amount of individual nucleotides detected in the filtrate varied to a considerable degree, and was highest for AMP, intermediate for ADP and lowest for ATP (Table 1) (71%, 44% and 15%, respectively, at a time when 95% of total Ca²⁺ was released).

Taken together our results show that the Ca²⁺ release pathway was activated in a reversible manner by 3,5(Me)₂NAPQI and that Ca²⁺ release was accompanied by the release of adenine and pyridine nucleotides from mitochondria. Release of all the nucleotides was prevented by CSA and is therefore concluded to occur through the mitochondrial pore described by Crompton's group [6–11]. It should be noted that despite pore opening, extramitochondrial ATP levels remained low. This explains why mitochondrial ATP was recently found inaccessible to exogenously added hexokinase after NAPQI-mediated Ca²⁺ release was near complete [24]. Conse-

Time after addition of 3,5(Me) ₂ NAPQI min		Ca ²⁺	ATP	ADP AMP NAD ⁺ nmol/mg protein (% of total)			NADP ⁺
0	total		4.03	2.89	3.45	0.93	0.25
	released	(0)	1.05 (26)	0.42 (15)	0.51 (15)	< 0.2	< 0.2
2	total		5.15	2.73	3.78	4.28	5.84
	released	(95)	0.66 (13)	1.20 (44)	2.68 (71)	2.04 (48)	2.54 (43)
7	total	. ,	2.78	2.69	3.75	3.87	4.25
	released	(100)	0.61 (22)	0.94 (35)	2.78 (74)	2.25 (58)	3.19 (75)

Table 1. Loss of nucleotides from mitochondria during 3,5(Me)₂NAPQI-induced Ca²⁺ release

Mitochondria were preincubated with succinate (5 mM) and rotenone (3 μ M) for 2.5 min, then loaded with Ca²⁺ (15 nmol/mg protein) for 1.5 min before the addition of 3,5(Me)₂NAPQI (90 nmol/mg protein) (zero time). Duplicate samples were taken at the indicated time points and analysed for total and released nucleotides by HPLC as described in Materials and Methods. Mitochondrial Ca²⁺ release was measured using Arsenazo III. Values are means of duplicate experiments conducted on separate mitochondrial preparations.

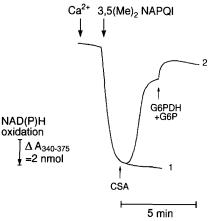


Fig. 6. Presence of oxidized pyridine nucleotides in the extramitochondrial medium. Mitochondria (2 mg/mL) were loaded with Ca²⁺ (15 nmol/mg) as described under Materials and Methods, and exposed to 3,5(Me)₂NAPQI (90 nmol/mg protein) (trace 1). For trace 2, CSA (1.5 nmol/mg protein) was added 1 min after 3,5(Me)₂NAPQI and intramitochondrial re-reduction of NAD(P)⁺ was allowed to proceed to completeness. Subsequently, glucose-6-phosphate (G6P, 250 μM) plus glucose-6-phosphate dehydrogenase (G6PDH, 0.4 U/mL) were added to the mitochondrial incubation as indicated by the arrow. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

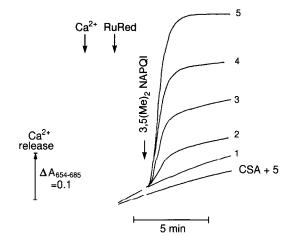


Fig. 7. Concentration-dependence of 3,5(Me)₂NAPQIinduced mitochondrial Ca2+ release in the presence of ruthenium red. Ca2+-loaded mitochondria were incubated for 2 min in the presence of ruthenium red (2 nmol/mg protein) before 3,5(Me)2NAPQI was added to the incubation, and Ca2+ release measured as described in Materials and Methods. The concentrations of 3,5(Me)₂NAPQI (nmol/mg protein) were 90 (trace 2), 97 (trace 3), 104 (trace 4), and 110 (trace 5), where trace 5 represents near complete release of all the sequestered Ca2-Where indicated, 110 nmol/mg protein of 3,5(Me)₂NAPQI was added to mitochondria preincubated with CSA (1.5 nmol/mg protein) before addition of ruthenium red (trace CSA + 5). Trace 1 is in the absence of any 3,5(Me)₂NAPQI. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

quently, extramitochondrial ATP is not an adequate parameter for the estimation of mitochondrial pore formation.

Ca²⁺ release induced by 3,5(Me)₂NAPQI in the presence of ruthenium red

The addition of ruthenium red, an inhibitor of the Ca²⁺ uniporter [26–28] to Ca²⁺-loaded control mitochondria produced a slow release of the accumulated Ca²⁺ (basal, ruthenium red-induced release). The subsequent exposure of the mitochondria to a

concentration of 3,5(Me)₂NAPQI (90 nmol/mg protein) that induced rapid release of Ca²⁺ in the absence of ruthenium red (cf. Fig. 1) only slightly enhanced the rate of the basal, ruthenium redinduced Ca²⁺ release (Fig. 7). When increasing the concentration of 3,5(Me)₂NAPQI (97, 104 or 110 nmol/mg protein), mitochondria released an

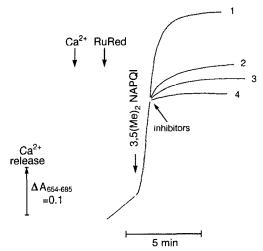


Fig. 8. Effect of CSA, ADP or MIBG on 3,5(Me)₂NAPQI-induced mitochondrial Ca²⁺ release in the presence of ruthenium red. Ca²⁺-loaded mitochondria were incubated for 2 min in the presence of ruthenium red (2 nmol/mg protein) and release of all sequestered Ca²⁺ was induced by addition of 3,5(Me)₂NAPOI (110 nmol/mg protein) (trace 1). Where indicated, MIBG (60 nmol/mg protein) (trace 2), ADP (125 nmol/mg protein) (trace 3), or CSA (1.5 nmol/mg protein) (trace 4) was added during Ca²⁺ release. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

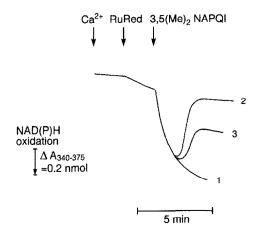


Fig. 9. 3,5(Me)₂NAPQI-induced pyridine nucleotide oxidation in the presence of ruthenium red. Ca²⁺-loaded mitochondria were incubated for 2 min in the absence (trace 1) or presence of ruthenium red (2 nmol/mg protein) (traces 2 and 3) before 3,5(Me)₂NAPQI (90 nmol/mg protein (traces 1 and 2) or 100 nmol/mg protein (trace 3)) was added to the incubation. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

increasing fraction of the total Ca^{2+} , after which the rate of Ca^{2+} release returned to the basal, ruthenium red-induced rate (Fig. 7). The addition of CSA (Fig. 7) or Mg^{2+} (data not shown) before a high concentration of $3.5(Me)_2NAPQI$ (110 nmol/mg protein) inhibited the prooxidant-stimulated Ca^{2+} release also in the presence of ruthenium red. Addition of CSA, MIBG or ADP, but not Mg^{2+} during $3.5(Me)_2NAPQI$ -induced Ca^{2+} release immediately prevented any further Ca^{2+} release (Fig. 8).

Mitochondrial NAD(P)H oxidation and nucleotide release in the presence of ruthenium red

As was observed in the absence of ruthenium red, 3,5(Me)₂NAPQI-induced Ca²⁺ release in the presence of ruthenium red was associated with oxidation of NAD(P)H. NAD(P)H oxidation was either partially reversible or irreversible, depending on whether the concentration of 3,5(Me), NAPQI used caused partial or near complete release of sequestered Ca²⁺ (Fig. 9). HPLC analysis of the filtrate revealed that also in the presence of ruthenium red, Ca2+ release was followed by a time-dependent release of adenine and pyridine nucleotides into the extramitochondrial medium (Table 2). Recovery of nucleotides in the filtrate varied for individual nucleotides and was highest for AMP, lowest for ATP and intermediate for ADP, NAD+ and NADP+ (Table 2). This pattern was similar to that observed in the absence of ruthenium red. Consequently, 3,5(Me)₂NAPQI-induced mitochondrial Ca²⁺ release was followed by pore opening also in the presence of ruthenium red.

Role of extramitochondrial versus intramitochondrial Ca^{2+} in the induction of Ca^{2+} release

Ruthenium red may not completely block Ca2+ uptake [20], leaving open the possibility of a limited degree of Ca²⁺-cycling. Therefore, the possibility of minimal Ca²⁺-cycling contributing to pore opening was tested by increasing the extramitochondrial Ca²⁺ concentration. To this aim, an additional 7.5 nmol of Ca²⁺/mg mitochondrial protein were added to Ca2+-loaded mitochondria (15 nmol Ca2+/mg protein) after blocking the uniporter with ruthenium red. The increased extramitochondrial Ca2+ concentration did not affect Ca2+ release induced by 3,5(Me)₂NAPQI (97 nmol/mg protein, a concentration which caused release of about 50% of the sequestered Ca²⁺) (Fig. 10). Instead, when the same amount of Ca²⁺ (i.e. 7.5 nmol/mg protein) was added and allowed to be completely sequestered by mitochondria (preloaded with 15 nmol Ca²⁺/mg protein) before the addition of ruthenium red, 97 nmol of 3,5(Me)₂NAPQI/mg protein induced near complete release of all the sequestered Ca2+ (i.e. 22.5 nmol Ca²⁺/mg protein) (Fig. 10). Again, Ca²⁺ release was prevented and NAD(P)H oxidation was reversible on preincubation with CSA (data not shown).

Effect of CSA, ADP or MIBG on the basal, ruthenium red-induced Ca²⁺ release

As shown in Fig. 11, CSA, MIBG and ADP also decreased the rate of the basal, ruthenium red-

Time after addition of 3,5(Me) ₂ NAPQI min		Ca ²⁺	ATP 5.53	ADP AMP NAD+ nmol/mg protein (% of total)			NADP+
0	total			2.66	3.92	0.35	<0.2
	released	(0)	1.05 (19)	0.40(15)	0.41(11)	< 0.2	< 0.2
2	total		4.74	3.43	3.99	3.50	4.38
	released	(95)	0.79(17)	1.25 (36)	1.86 (47)	1.86 (53)	1.50 (34)
7	total	` ′	4.75	2.57 `	3.35	3.58	3.51
	released	(100)	0.92 (19)	1.25 (49)	2.27 (68)	2.16 (58)	2.50 (60)

Table 2. Loss of nucleotides from mitochondria during 3,5(Me)₂NAPQI-induced Ca²⁺ release in the presence of ruthenium red

Mitochondria were preincubated with succinate (5 mM) and rotenone (3 μ M) for 2.5 min, then loaded with Ca²⁺ (15 nmol/mg protein) for 2 min before the addition of ruthenium red (2 nmol/mg protein). After an additional 2 min, 3,5(Me)₂NAPQI (110 nmol/mg protein) was added (zero time). Duplicate samples were taken at the indicated time points and analyzed for total and released nucleotides by HPLC as described in Materials and Methods. Mitochondrial Ca²⁺ release was measured using arsenazo III. Values are means of duplicate experiments conducted on separate mitochondrial preparations.

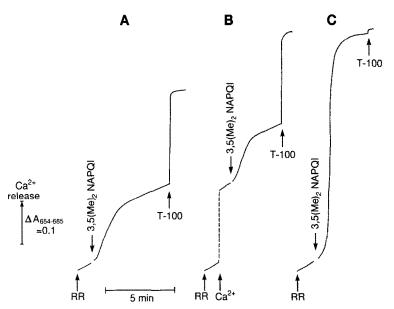


Fig. 10. Effect of extra- versus intramitochondrial Ca²⁺ on 3,5(Me)₂NAPQI-induced Ca²⁺ release in the presence of ruthenium red. Ca²⁺-loaded mitochondria were incubated for 2 min in the presence of ruthenium red (2 nmol/mg protein) before 3,5(Me)₂NAPQI (97 nmol/mg protein) was added to the incubation medium (A). In (B) and (C) an additional 7.5 nmol of Ca²⁺/mg protein were added to the incubation after (B) or before (C) uptake was inhibited by ruthenium red. Traces are representative of three independent experiments conducted on separate mitochondrial preparations. Abbreviation used: T-100, Triton X-100.

induced Ca²⁺ release by around 45% (ADP, MIBG) to 70% (CSA); however, neither of these reagents could prevent its release completely. The results suggest that the basal, ruthenium red-induced mitochondrial Ca²⁺ release involves at least two components, one of which is sensitive to the same agents that inhibit the prooxidant-stimulated Ca²⁺ release.

DISCUSSION

In this study we have shown that opening of the

Ca²⁺-dependent mitochondrial pore [6–11] occurred during the stimulation of Ca²⁺ release from mitochondria by the prooxidant 3,5(Me)₂NAPQI. This was evidenced by (i) the release of mitochondrial adenine and pyridine nucleotides and (ii) the reversibility of opening induced by ADP and CSA, in agreement with other reports [29, 30]. In addition to CSA or ADP, MIBG and Mg²⁺ were found to be inhibitors of both Ca²⁺ release and pore opening.

In the absence of ruthenium red, Ca²⁺ release markedly preceded the appearance of mitochondrial nucleotides in the extramitochondrial medium. This

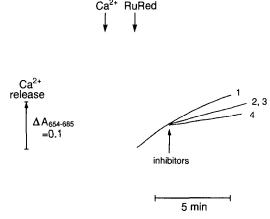


Fig. 11. Effect of CSA, ADP or MIBG on the basal, ruthenium red-induced mitochondrial Ca²⁺ release. Ca²⁺-loaded mitochondria were incubated for 2 min in the presence of ruthenium red (2 nmol/mg protein) (trace 1). Where indicated, MIBG (60 nmol/mg protein) (trace 2), ADP (150 nmol/mg protein) (trace 3), or CSA (1.5 nmol/mg protein) (trace 4) was added. Trace 1 is in the absence of any inhibitor. Traces are representative of three independent experiments conducted on separate mitochondrial preparations.

difference in kinetics could be due to different rates of release through the mitochondrial pore [8] or. alternatively, activation of an additional Ca²⁺selective pathway by the prooxidant. Indeed, the existence of such a Ca²⁺-selective pathway that can be activated by prooxidants has been suggested by many reports [31–37], although the participation of the Ca²⁺-dependent pore could not be ruled out unequivocally. However, Schlegel et al. have recently shown that tBuOOH activates a Ca²⁺-selective release pathway without pore opening under experimental conditions where Ca2+ reuptake through the uniporter was prevented by ruthenium red or EGTA [20]. In agreement with that report, pore opening was not induced in the presence of ruthenium red when using a concentration of 3,5(Me)₂NAPQI that caused pore opening in the absence of ruthenium red. However, in contrast with tBuOOH, Ca²⁺ release was only minimally stimulated by this concentration of 3,5(Me)₂NAPQI when ruthenium red was present. When the concentration of the prooxidant was increased sufficiently to cause release of Ca²⁺ in the presence of ruthenium red, nucleotides were also lost from the mitochondria. As was observed in the absence of ruthenium red, ADP, CSA, Mg2+ and MIBG were inhibitory on both Ca2+ and nucleotide release. Thus, in the case of the prooxidant 3,5(Me)2NAPQI, blocking the uniporter by ruthenium red did not allow a clear dissociation between a Ca²⁺-selective efflux pathway and mitochondrial pore opening.

Ruthenium red may not completely block Ca²⁺ uptake [20], leaving open the possibility of a limited degree of Ca²⁺-cycling. If this low level of Ca²⁺-cycling were the critical event triggering pore opening by 3,5(Me)₂NAPQI in the presence of ruthenium

red, increasing the extramitochondrial Ca²⁺ concentration would be expected to stimulate 3,5(Me)₂NAPQI-induced pore opening and Ca²⁺ release. However, this was not the case. On the other hand, an increase in the intramitochondrial Ca²⁺ load was found to promote the Ca²⁺ release. Consequently, the results from this study do not support a role for the process of Ca²⁺-cycling in pore opening by 3,5(Me)₂NAPQI, but implicate the intramitochondrial Ca²⁺ load as a critical factor in triggering pore opening.

In the presence of ruthenium red, mitochondria released a concentration-dependent fraction of all sequestered Ca²⁺ upon addition of 3,5(Me)₂NAPQI. The latter finding clearly shows that the mitochondrial suspension does not react homogeneously to 3,5(Me)₂NAPQI, but rather that mitochondrial subpopulations of different sensitivity of Ca²⁺ release towards the concentration of prooxidant exist, as originally suggested from tBuOOH-induced mitochondrial swelling experiments [38].

The existence of subpopulations with different sensitivities towards 3,5(Me)2NAPQI may explain why different concentrations of 3,5(Me)₂NAPQI are required to induce near complete Ca2+ release depending on the absence versus presence of ruthenium red. Thus, only a limited fraction of mitochondria releases the sequestered Ca2+ upon addition of 90 nmol 3,5(Me)₂NAPQI/mg protein. In the presence of ruthenium red Ca²⁺ remains extramitochondrially; however, in the absence of ruthenium red Ca²⁺ is taken up by a fraction that is less sensitive to the prooxidant. The increased mitochondrial Ca²⁺ load in turn lowers the sensitivity of these mitochondria towards the prooxidant, thereby leading to a cascade of further, and eventually complete, Ca2+ release.

Taken together the results obtained in the absence versus the presence of ruthenium red allow the following conclusions about the critical factors regulating 3,5(Me)₂NAPQI-induced mitochondrial Ca²⁺ release to be drawn. 3,5(Me)₂NAPQIinduced mitochondrial Ca2+ release is a cooperative phenomenon, which is regulated by at least two different parameters. These are (i) the prooxidant concentration, which acts synergistically with (ii) the mitochondrial Ca2+ load. The open probability of the mitochondrial permeability transition pore (MTP) measured in swelling experiments using depolarized mitochondria was similarly reported to be regulated by the synergistic action of (i) matrix alkalinization, and (ii) the mitochondrial Ca²⁺ content, with matrix acidification being antagonized by the Ca²⁺ content [18]. It is challenging to speculate that the synergism observed here between the Ca2+ load and the prooxidant concentration relate to the synergism observed between Ca2+ load and mitochondrial pH. The prooxidant concentration would be reflected in the matrix pH through the mitochondrial NAD(P)H redox state according to the following equation:

$$NAD(P)H + H^+ + 3,5(Me)_2NAPQI$$

$$\Rightarrow NAD(P)^+ + 3,5(Me)_2NAPQIH_2,$$

where protons are consumed during the reaction. This suggestion is supported by the fact that the chemical oxidation of NADH by $3.5(Me)_2NAPQI$ (25 μ M) is accompanied by an increase in pH of more than 0.4 pH units [M. Weis, unpublished observation].

The agents shown here to inhibit the prooxidantinduced mitochondrial Ca2+ release also inhibited the basal, ruthenium red-induced Ca2+ release by around 45–70%, suggesting that the basal ruthenium red-induced Ca²⁺ release involves, in part, the same pathway. The preventive action of CSA, ADP and Mg²⁺ on mitochondrial Ca²⁺ release is likely to relate to the interaction of the compounds with the mitochondrial pore complex. The mechanism of action underlying the protective action of MIBG is less clear. Thus, MIBG is an inhibitor of protein mono(ADP)ribosylation [39]; however, its instantaneous action in preventing further Ca²⁺ release and pore opening when added after the prooxidant (in the presence of ruthenium red) as well as its increased protective effect at higher mitochondrial Ca2+ load [M. Weis, unpublished observations] suggests the possibility of an additional mode of action, i.e. competition with Ca²⁺ for the Ca²⁺ binding site regulating pore opening.

Little is known about the possible physiological role of the mitochondrial pore. Since pore opening was completely prevented at a concentration of Mg²⁺ that was much lower than that found in the cytosol of mammalian cells [40], pore opening would be expected to be restricted to pathological rather than physiological conditions of the cell. However, we have recently reported that CSA [4] and MIBG [41] lead to a rapid accumulation of Ca²⁺ by mitochondria in isolated hepatocytes. Whether the latter observation reflects a low level of pore opening under physiological conditions or an additional Ca²⁺ release pathway that is sensitive to CSA and MIBG, as suggested by the work of Schlegel *et al.* [20], remains unclear.

In summary, the present data show that the ruthenium red-induced and the 3,5(Me)₂NAPQI-induced mitochondrial Ca²⁺ release occur through a common component. Considering the well known role of several of these inhibitors in regulating the mitochondrial pore, it is proposed that the component mediating prooxidant-induced and ruthenium red-induced Ca²⁺ release is involved in mitochondrial pore formation.

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