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## Calcium signaling in cardiac mitochondria

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### Abstract

Mitochondrial Ca signaling contributes to the regulation of cellular energy metabolism, and mitochondria participate in cardiac excitation-contraction coupling (ECC) through their ability to store Ca, shape the cytosolic Ca signals and generate ATP required for contraction. The mitochondrial inner membrane is equipped with an elaborate system of channels and transporters for Ca uptake and extrusion that allows for the decoding of cytosolic Ca signals, and the storage of Ca in the mitochondrial matrix compartment. Controversy, however remains whether the fast cytosolic Ca transients underlying ECC in the beating heart are transmitted rapidly into the matrix compartment or slowly integrated by the mitochondrial Ca transport machinery. This review summarizes established and novel findings on cardiac mitochondrial Ca transport and buffering, and discusses the evidence either supporting or arguing against the idea that Ca can be taken up rapidly by mitochondria during ECC.

### Keywords

excitation-contraction coupling; heart; intracellular calcium; mitochondrial Ca transport; Ca buffering

## 1. Introduction: Cardiac excitation-contraction coupling and mitochondrial Ca

Cardiac contraction is regulated by beat-to-beat elevations of cytosolic calcium ( $[Ca]_i$ ) by a process termed excitation-contraction coupling (ECC) [1] where membrane depolarization induced by an action potential leads to Ca entry through voltage-activated L-type Ca channels. Entering Ca triggers Ca release from the sarcoplasmic reticulum (SR) Ca store via ryanodine receptor (RyR) Ca release channels by a mechanism known as Ca-induced Ca release (CICR). CICR increases global  $[Ca]_i$  which activates proteins of the contractile apparatus and initiates cell contraction. Subsequent relaxation occurs by removal of Ca from the cytosol via four main pathways including reuptake via the SR Ca-ATPase (SERCA), extrusion via sarcolemmal Na/Ca exchange (NCX) and the sarcolemmal Ca-ATPase. A fourth avenue of Ca sequestration potentially involves mitochondrial Ca uptake since

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### Disclosures

None.

mitochondria are equipped with an efficient machinery for Ca transport and are capable of storing large amounts of Ca. Mitochondria of cardiomyocytes are known to accumulate Ca during elevations in cytosolic  $[Ca]_i$  (for reviews cf. [2–8]), however, the kinetics of mitochondrial Ca cycling and buffering during ECC have remained highly controversial [9]. The question of whether and how beat-to-beat changes of cytosolic  $[Ca]_i$  during ECC translate into changes of mitochondrial matrix Ca concentration ( $[Ca]_m$ ) has important ramifications for cardiac physiology and pathophysiology. First, mitochondrial Ca uptake and buffering have the potential to shape the cytosolic Ca transient and therefore contribute to the regulation of contractile activity, and second, mitochondrial Ca uptake regulates cellular metabolism and energy supplies required for contraction. The latter occurs through the Ca-dependence of key enzymes of the tricarboxylic acid (TCA) cycle and possibly also Ca-dependent regulation of various sites of the electron transport chain (ETC) and the mitochondrial  $F_1/F_0$  ATP synthase [10–12].

In this review we will briefly summarize the elements of the mitochondrial Ca transport machinery and recent novel findings on mitochondrial Ca buffering. Furthermore, we discuss how mitochondria encode rapid beat-to-beat cytosolic Ca oscillations and critically review arguments in favor and against rapid mitochondrial Ca uptake during ECC.

## 2. Mitochondrial Ca transport

Mitochondria are cytosolic double-membrane organelles that have been dubbed 'power plants' of the cell [13, 14] for their ability to generate ATP to satisfy cellular energy demands. Mitochondria, however, participate in a myriad of other cellular processes such as ion homeostasis, redox signaling, apoptotic and necrotic cell death, as well as the control of cell cycle and cell growth [15]. Furthermore, mitochondria undergo remodeling in cardiac disease, including arrhythmia and heart failure, that has profound effects on mitochondrial structure-function that become important determinants of the course of the disease [6, 16–19]. In cardiac myocytes mitochondria occupy ~35% of the cell volume [20, 21] reflecting the high energy demands of these cells. Mitochondria possess an elaborate system of Ca uptake and extrusion mechanisms and pathways (Fig. 1) that allow for a fine-tuned regulation of  $[Ca]_m$  [22, 23].

### 2.1. Mitochondrial Ca uptake

Several mechanisms for mitochondrial Ca uptake (Fig. 1A) have been described and proposed for cardiac myocytes - some well established, others still controversial or a matter of debate: the mitochondrial Ca uniporter (MCU) [24–27], a rapid mode of Ca uptake (RaM) [28–31], the mitochondrial ryanodine receptor type 1 (mRyR1) [32–35] and the recently proposed leucine-zipper-EF-hand-containing transmembrane protein 1 (LETM 1) [36–38], all, together with additional putative Ca uptake pathways (see discussion below), localized to the inner mitochondrial membrane (IMM).

The best established and most characterized pathway for mitochondrial Ca uptake is through the MCU [24–27], driven by the large electrochemical gradient (mitochondrial membrane potential  $\Delta\Psi \sim -180$  mV) for Ca across the IMM. Many characteristics of the MCU (kinetic and electrophysiological [39] properties, Ca-dependence and ion selectivity) have been known for many years. MCU is regulated by a number of modulators (activators and inhibitors) that include divalent cations, lanthanides, adenine nucleotides, and ruthenium compounds (for review see [3, 40]). Among the latter RU-360 is preferentially used experimentally to inhibit MCU [41]. Nonetheless, it was not until recently that the molecular identity of the transporter has been revealed, essentially at the same time by two different laboratories [26, 27]. Both studies are in agreement that a transmembrane protein previously identified as CCDC109A is the molecular identity of the MCU (but see [42]). CCDC109A

(MCU) has two transmembrane domains that are likely to form the Ca channel pore. The Ca binding protein MICU1 (mitochondrial calcium uptake 1) has been postulated to be required for mitochondrial Ca uptake [43], and may act as an auxiliary regulatory protein to MCU. Recently, MICU1 was shown to serve as a 'gatekeeper' for MCU that determines the Ca threshold for uptake and prevents mitochondrial Ca overload [44]. Furthermore, Ca/calmodulin-dependent protein kinase II (CaMKII), a key signaling molecule involved in cardiac pathologies including heart failure, promotes mPTP opening and myocardial cell death by increasing mitochondrial Ca uptake via MCU [45].

In addition to the MCU other avenues of Ca entry into mitochondria have been proposed. RaM (demonstrated in isolated heart mitochondria) operates transiently during the initial phase of pulsatile elevations of extramitochondrial [Ca] ( $[Ca]_{em}$ ) [28–31]. Ca uptake via RaM is several hundred times faster than uptake via MCU, however, the recovery of RaM following a Ca pulse in isolated heart mitochondria required more than 60 s [28] rendering this mechanism potentially inactivate during cardiac Ca oscillations.

Of more controversial nature are ryanodine receptors that have been located in the IMM and proposed to be one of the mitochondrial Ca uptake pathways [32–35]. Reconstitution into lipid bilayers yielded cesium-conducting, Ca-sensitive, large conductance (500–800 pS) channels [34] with characteristics similar to SR RyRs. The mitochondrial RyR exhibits biochemical, pharmacological, and functional properties similar to the skeletal muscle RyR (type 1, RyR1), and is therefore called mRyR1.

Additional putative mitochondrial Ca uptake mechanisms and pathways have been proposed, including two Ca-selective conductances referred to as mCa1 and mCa2 [46], Coenzyme Q10 (an element of the mitochondrial electron-transport chain; [47]), the uncoupling proteins UCP2 and UCP3 [48], and LETM1 [36–38]). A debated issue is whether the LETM1 protein can function as a high affinity mitochondrial Ca influx mechanism. LETM1 was identified first as a mitochondrial potassium/proton antiporter, but has since been argued to function as a Ca/H antiporter [37]. It remains to be determined, however, whether normal electrochemical gradients for protons and Ca would allow LETM1 to act as a Ca influx pathway, or whether LETM1 might actually act as Ca efflux rather than influx mechanism [38].

## 2.2. Mitochondrial Ca efflux

Ca extrusion from the mitochondrial matrix occurs by Na-dependent and Na-independent mechanisms (Fig. 1B). The predominant Ca extrusion pathway in cardiac mitochondria is Na-dependent [49, 50], while Na-independent efflux (particularly proton/Ca exchange, mHCX) plays little to no role [51, 52]. The Na-dependent extrusion occurs via mitochondrial Na/Ca exchange (mNCX, with NCLX (Na/Ca/Li exchanger) suggested as its molecular identity [53, 54]), i.e. mediated by an antiporter that utilizes the Na gradient across the inner membrane. The  $[Na]_i$  dependence of mNCX is sigmoidal with a half-maximal activity at ~4–8 mM [55–58], i.e. in the range of resting  $[Na]_i$  observed under physiological conditions in cardiomyocytes [59–62], making mNCX potentially sensitive to physiological fluctuations in cytosolic  $[Na]_i$  [56]. However, no significant variations in bulk  $[Na]_i$  are typically observed during the normal cardiac cycle, and only unphysiological increases in stimulation frequency or pathological conditions such as heart failure resulted in significant changes of bulk  $[Na]_i$  [60, 63, 64]. mNCX plays an important role in modulating the steady-state balance between extra- and intramitochondrial Ca [65]. Cellular Na overload, as observed in heart failure, disrupts this equilibrium, resulting in changes of pyridine nucleotide redox potential and increase in reactive oxygen species (ROS) generation, with ultimately detrimental effects on mitochondrial bioenergetics and mismatch of cardiac energy supply and demand [63, 66, 67]. Na-dependent efflux via mNCX is

inhibited by divalent cations and a number of drugs, including Ca channel blockers (e.g. diltiazem, verapamil) and benzodiazepine derivatives (e.g. clonazepam, CGP-37157) [3, 7, 40]. Furthermore, it has long remained elusive and controversial [2, 50] whether mNCX is an electroneutral or electrogenic [68, 69] antiporter. Nonetheless, the majority of data agree that mNCX exchanges 3 Na ions for 1 Ca, and thus is electrogenic [7, 70–72]. As an electrogenic exchanger, Ca extrusion would tend to depolarize  $\Delta\Psi$ , thereby reducing the electrical gradient for Ca uptake and favor net mitochondrial Ca extrusion. The IMM also hosts a Na/proton exchange system (mNHX) [73], which serves as the pathway for Na extrusion. Through this mechanism the energy requirement for Ca extrusion via mNCX is coupled to proton movement across the ETC during mitochondrial respiration.

Of considerable controversy remains whether the mitochondrial Permeability Transition Pore (mPTP) can function as a Ca efflux pathway [7, 74]. The mitochondrial permeability transition defines a sudden increase in the permeability of the IMM to ions and solutes with molecular weights up to ~1.5 kDa [75]. Elevated levels of matrix Ca and ROS are known inducers of mPTP opening [40, 76]. The permeability transition process is attributed to the opening of a voltage- and Ca-dependent, cyclosporin A (CsA)-sensitive [77], high-conductance channel. The molecular identity of mPTP is largely unknown (see [76, 78, 79] for recent reviews) and the original model of the mPTP proposed a multi-protein complex that included the adenine nucleotide translocase in the IMM, the voltage-dependent anion channel of the outer membrane (VDAC), the  $F_1/F_0$  ATP-synthase, and the protein cyclophilin D (Cyp D) [80]. In addition, complex I of the ETC has also been suggested to be a part of the mPTP complex [81], and modulation of mPTP by complex I and Cyp D may share a common mechanism [82]. However, this mPTP model has been challenged by recent genetic studies [83, 84]. Electrophysiological studies revealed that the mPTP is a large-conductance (1.3 nS) channel with numerous subconductance states and may flicker rapidly between a fully closed and a subconductance state [85, 86]. mPTP is activated by high matrix [Ca], oxidative stress and depolarization. Repetitive opening and closing of the mPTP has been demonstrated in individual isolated heart mitochondria under conditions of oxidative stress [87, 88]. However, whether mPTP can serve as a mitochondrial Ca release channel [25] under physiological conditions has remained controversial. The existence of small and possibly ion-selective subconductance states of the mPTP [89] could potentially allow brief openings of the mPTP and serve as a mechanism for fast dissipation of  $\Delta\Psi$  and subsequent Ca efflux without causing dramatic changes to the matrix environment, but possibly acting as a Ca 'relief valve' and provide protection against cell injury under mitochondrial Ca overload conditions. However, experimental evidence for the participation of mPTP in mitochondrial Ca signaling under physiological conditions remains scarce [90–92] and controversial [93]. Under pathological conditions of heart failure mice lacking Cyp D exhibit a substantially more pronounced maladaptive phenotype and a reduction in myocardial function that was associated with altered mPTP-mediated Ca efflux which resulted in elevated matrix Ca, leading to a mismatch of energy metabolism and myocardial workload [91]. Under conditions of  $\Delta\Psi$  dissipation Ca can enter mitochondria through the mPTP [55] potentially maintaining a mitochondrial Ca sink for cytosolic Ca overload when MCU is inactivated.

### 3. Mitochondrial Ca signals during ECC: The controversy of beat-to-beat mitochondrial Ca transients

Historically, two different theories (Fig. 1C) have evolved on how mitochondria decode rapid cytosolic Ca transients (for review see for example refs. [2–4, 7, 9]). In model I, originally introduced by Crompton [94], Ca uptake into mitochondria is slow and balanced by an even slower release of accumulated Ca ions. Fast cytosolic beat-to-beat Ca oscillations are integrated by the Ca transport machinery of the IMM. The response to a train of

cytosolic Ca transients is a gradual amplitude- and frequency-dependent increase of  $[Ca]_m$  until a new steady-state is reached when the amount of Ca gained during a single cycle equals Ca removal from the matrix compartment. Consequently, beat-to-beat  $[Ca]_m$  changes are small, and energetic requirements of mitochondrial Ca transport are minimal. In contrast, model II describes how fast cytosolic Ca oscillations are efficiently translated into rapid beat-to-beat changes of  $[Ca]_m$  of considerable amplitude. For this to occur, rapid mitochondrial Ca uptake as well as fast Ca efflux mechanisms are mandatory prerequisites, with Ca uptake being large enough to overcome the buffering capacity of both the cytosol and the mitochondrial matrix compartment, thus requiring high SR Ca release fluxes. With this scenario, mitochondrial Ca uptake would effectively buffer cytosolic Ca transients and therefore play a critical role for shaping the cytosolic Ca transient during ECC, and thus potentially regulate contraction on a beat-to-beat basis. With approximately one third of cell volume being occupied by mitochondria in cardiac cells, the additional SR Ca fluxes (and energy requirement linked to them) would have to be substantial [2].

The controversy surrounding beat-to-beat changes in  $[Ca]_m$  in the heart is certainly (at least in part) related to experimental limitations of available methods for reliable measurements of  $[Ca]_m$  [4]. This caveat applies to many such studies irrespective whether the outcome favored model I or II. Below we discuss experimental data (and their limitations) supporting either model I or model II.

### 3.1. Model I: $[Ca]_m$ reflects the slow integration of cytosolic Ca transients

Evidence in support of slow mitochondrial integration of cytosolic Ca transients are based on studies utilizing electron probe microanalysis (EPMA) and fluorescence microscopy techniques. EPMA uses rapidly frozen tissue samples and has the advantages of a resolution close to electron microscopy, however it measures total mitochondrial Ca ( $[Ca]_{m,tot}$ ) instead of free  $[Ca]_m$ . EPMA studies on hamster [95, 96] and rat [97] papillary muscles were unable to resolve fast changes in  $[Ca]_{m,tot}$ , not even after  $\beta$ -adrenergic stimulation to increase Ca cycling [96, 98].

In a majority of studies, fluorescent dyes (such as the membrane permeable ester forms of indo-1, rhod-2 or fluo-3) known to compartmentalize into mitochondria were used to monitor  $[Ca]_m$  directly. To eliminate the cytosolic component of the fluorescent signals, cells were treated with manganese [99–102] or cobalt [103], exposed to higher temperature [104, 105], membrane permeabilized [106, 107] or dialysed [66, 101]. Using indo-1 loaded rat [99, 105], hamster [100], ferret and cat [101] ventricular myocytes with subsequent Mn-quenching of cytosolic dye, it was shown that an increase in the stimulation frequency from 0.2 to 4 Hz in the presence of  $\beta$ -adrenergic stimulation [99, 104] or cellular Ca loading via sarcolemmal NCX [101] led to a slow rise of  $[Ca]_m$  from ~100–200 to ~500–800 nM, however no beat-to-beat changes in  $[Ca]_m$  were observed. In the absence of  $\beta$ -adrenergic stimulation, only modest increases in  $[Ca]_m$  could be achieved in rat myocytes by electrical stimulation at 2 Hz [100], which indicated that only large amplitude cytosolic Ca transients were sensed by mitochondria. Miyata et al. [99] demonstrated an exponential relationship between  $[Ca]_m$  and  $[Ca]_i$ , with a threshold for mitochondrial Ca uptake being at a  $[Ca]_i$  of ~500 nM. Similarly, Zhou et al. [101] reported that under conditions of high cellular Ca load imposed by membrane depolarization in ferret and cat myocytes, phasic increases of  $[Ca]_m$  could be detected, although they were slow and only observed at diastolic  $[Ca]_i > 400$  nM. The authors concluded that mitochondria of intact cells did not take up detectible amounts of Ca during individual contractions. These findings are in agreement with a recent study where  $[Ca]_m$  changes during cytosolic Ca transients were quantified [108]. Changes in  $[Ca]_m$  during individual cytosolic Ca transients amounted to only approximately 2–10 nM per beat, but integrated gradually to a new steady-state. Total mitochondrial Ca uptake for larger  $[Ca]_m$  transients amounted to only about 1% of the SR Ca uptake during a normal

cytosolic Ca transient. However, there is evidence for species-dependent differences in the kinetics of  $[Ca]_m$  as demonstrated in a comparison between rat and guinea pig cardiomyocytes where  $[Ca]_m$  transients were observed in indo-1 loaded guinea-pig myocytes, while no changes in  $[Ca]_m$  were observed in rat cardiomyocytes [105].

A general criticism of the experimental approach used in some of the studies outlined above is the possible interference of Mn with Ca transport across the IMM. Mn can be sequestered by the MCU, and Mn can potentially quench the fluorescence of the dye compartmentalized into mitochondria. While Miyata et al. [99] provided evidence that Mn had no effect on mitochondrial Ca uptake or cell shortening, other studies demonstrated that Mn significantly inhibited Ca efflux [109]. Nonetheless, removal of cytosolic indo-1 by heat treatment gave similar results to the Mn quench technique [104, 105]. Moreover, when fura-2 loaded rat heart mitochondria were exposed to Ca oscillations of 100 cycles/min,  $[Ca]_m$  increased proportionally to the average rise in extramitochondrial  $[Ca]$ , but independently of oscillation frequency [110].

A novel experimental approach was developed in our laboratory by Sedova et al. [106] to study  $[Ca]_m$  kinetics in response to cytosolic Ca spikes (Fig. 2). The approach allowed the simulation of fast cytosolic Ca transients in membrane-permeabilized cells.  $[Ca]_m$  was measured with fluo-3 entrapped inside mitochondria. Permeabilized cells have the unique advantage that the cytosolic environment can be controlled precisely (including complete removal of cytosolic Ca indicator dye) while the arrangements and interaction between intracellular membranes and organelles (SR, mitochondria) remain structurally and functionally intact [111, 112]. Control experiments indicated that mitochondrial Ca uptake was initiated only when  $[Ca]_{em}$  was elevated above  $0.5 \mu M$ , confirming the existence of a threshold for MCU activation [101, 107]. Ca entry via MCU exhibited a sigmoidal  $[Ca]_{em}$ -dependence (half-maximal uptake rate at  $[Ca]_{em}=4.4 \mu M$ ). To simulate cytosolic Ca transients (Fig. 2A), cells were placed in the laminar flow of a Ca-free solution and then exposed to rapid ejections of a solution containing  $100 \mu M$  Ca from a micropipette positioned upstream of the cell with regard to the direction of the bulk flow. Calculations revealed that with the applied technique cells were exposed to  $\sim 5 \mu M$   $[Ca]_{em}$  during each ejection pulse, which is in the range of physiological cytosolic Ca transient amplitudes and close to half-maximal activation of MCU. With this technique rapid beat-to-beat changes in  $[Ca]_i$  were simulated by rapidly switching  $[Ca]_{em}$  between low and high levels. The technique allowed to precisely vary pulse duration and pulse frequency. Exposure to a train of Ca transients evoked a gradual, but pulse duration- (Fig. 2B) and frequency-dependent (Fig. 2C) elevation of  $[Ca]_m$ , however no  $[Ca]_m$  oscillations were observed. As shown in Fig. 2D transient elevations of  $[Ca]_m$  in response to a cytosolic Ca pulse could be observed, provided a sufficiently long time interval between pulses was allowed. The recovery time of the  $[Ca]_m$  transients was in the range of minutes, i.e. orders of magnitude longer than the time interval between typical cytosolic Ca oscillations. Taken together, the data suggest that in cat ventricular myocytes fast cytosolic Ca transients are integrated by mitochondrial Ca transport systems resulting in a frequency- and pulse duration-dependent net accumulation of Ca in the matrix, thus supporting model I of mitochondrial decoding of fast cytosolic Ca signals. These data are complemented by the quantitative estimates of mitochondrial Ca uptake and its kinetics in response to SR Ca release by Andrienko et al., as discussed above [108].

### 3.2. Model II: Beat-to-beat oscillations of $[Ca]_m$

Evidence in support of a beat-to-beat translation of cytosolic Ca transients into  $[Ca]_m$  oscillations (model II; Fig. 1C) was first obtained in guinea pig myocytes using EPMA [113, 114]. Although changes  $[Ca]_{m,tot}$  were resolved only in myocytes in which the SR Ca content and diastolic  $[Ca]_i$  was enhanced, beat-to-beat variations in  $[Ca]_{m,tot}$ , which

followed cytosolic Ca transients with 15–25 ms delay, peaked within 30–45 ms, and reversed by 90 ms, could be demonstrated. The decline of the  $[Ca]_{m,tot}$  transient was attributed to mNCX activity. In a later study [115] the same authors found that subsarcolemmal mitochondria responded to changes in  $[Ca]_i$  with  $[Ca]_{m,tot}$  transients, while central mitochondria did not.

Further support for fast mitochondrial  $[Ca]_m$  transients came from studies using laser scanning confocal microscopy in combination with fluorescent Ca indicators (e.g. fluo-3, indo-1, rhod-2) entrapped in mitochondria and identifying mitochondrial locations by co-staining with 'mitochondrial markers' such as the voltage-sensitive dye TMRM or rhodamine 123 [116–119]. Recordings obtained during electrical or  $\beta$ -adrenergic stimulation revealed  $[Ca]_m$  transients with an identical time course in mitochondria and cytosol. The lack of kinetic differences between the two signals raises the possibility that the mitochondrial signal was 'contaminated' to a significant degree with cytosolic signal due to the presence of indicator dye in the cytosol. Refining the dye loading protocol using different incubation temperatures it could be demonstrated that the majority of the signal arose from mitochondria since  $[Ca]_m$  measured with rhod-2 was sensitive to ruthenium red, while  $[Ca]_i$  measured with fluo-3 was only slightly affected [118, 119]. Moreover, Mackenzie et al. [120] were able to record mitochondrial  $[Ca]_m$  transients in rat atrial myocytes with rhod-2 that were abolished by the mitochondrial inhibitors antimycin and oligomycin, while under identical experimental conditions cytosolic Ca transients could still be evoked. These authors also confirmed the importance of mitochondrial location, showing that during electrical stimulation peripherally located mitochondria sequestered most of the released Ca.

To circumvent the problem of signal contamination mentioned above, novel fluorescent probes were developed that could be genetically encoded and targeted specifically to the mitochondrial matrix compartment (for review see [4, 121]). Using the Ca-sensitive photoprotein aequorin and green fluorescent protein-based Ca indicator pericam as such probes [122], it could be demonstrated that spontaneous cytosolic Ca oscillations in cultured neonatal rat ventricular myocytes were synchronous with mitochondrial  $[Ca]_m$  oscillations. Elevation of extracellular  $[Ca]$  or  $\beta$ -adrenergic stimulation resulted in a substantial increase in spike amplitude in both compartments, and increased inter-spike  $[Ca]_m$ , suggesting that Ca extrusion from mitochondria was slower than mitochondrial Ca uptake, resulting in mitochondrial Ca accumulation. With a similar technique (adenoviral infection with aequorin targeted to cytosol and mitochondria)  $[Ca]_m$  and  $[Ca]_i$  were shown to increase simultaneously in adult rat myocytes in response to electrical stimulation, however the decay of  $[Ca]_m$  was much slower [123], and the observation of  $[Ca]_m$  transients required increased extracellular  $[Ca]_o$  or  $\beta$ -adrenergic stimulation. The mitochondrial fluorescence signal was insensitive to the MCU inhibitor Ru360 (presumably due to poor membrane permeability of Ru360 in these cells), increased by the mNCX blocker clonazepam, and decreased by the mitochondrial uncoupler FCCP in combination with oligomycin.

Szalai et al. [124] showed that activation of RyRs by Ca, ryanodine or caffeine evoked cytosolic Ca oscillations that were synchronized with  $[Ca]_m$  oscillations in cardiac H9c2 myotubes. However, the frequency of  $[Ca]_m$  oscillations observed in this study was nearly an order of magnitude slower than physiological heart rates, and an increase in basal (inter-spike)  $[Ca]_m$  was observed, suggesting that the rate of mitochondrial Ca removal was not fast enough to extrude all the Ca before the next cytosolic Ca spike. Addition of 30  $\mu$ M Ca to the bath was required to match the rates of rise of the caffeine-induced  $[Ca]_m$  transients, indicating that the local Ca in the microdomain near the mitochondria during SR Ca release is much higher than the average cytoplasmic concentration. In another study by the Hajnoczky group, more direct evidence of a local fast communication between the SR Ca release units and mitochondria was forwarded. Also in permeabilized H9c2 myotubes, local

SR Ca release events in form of Ca sparks elicited  $[Ca]_m$  transients in adjacent mitochondria, referred to as Ca marks (i.e. mitochondrial Ca sparks) [125]. This and other studies (e.g. [126, 127]) have provided strong evidence for the importance of a SR-mitochondria Ca signaling microdomain where a close physical association between the SR Ca release apparatus and mitochondria exists (for recent reviews see e.g. [128–130]). The functional and structural connection between SR and mitochondria is attributed to inter-organelle tether proteins [131–133], such as mitofusin 2 [134]. This microdomain enables elevations of local  $[Ca]$  possibly into the tens of micromolar range [135, 136] bringing it into the range of the relatively low Ca affinity of the uniporter [106, 129].

Evidence for fast mitochondrial Ca uptake also came from a recent study where cytosolic and mitochondrial Ca was monitored simultaneously in guinea-pig cardiomyocytes using a differential Ca dye loading technique [66]. The study showed that cytosolic Ca transients elicited by voltage-clamp depolarization were accompanied by fast  $[Ca]_m$  transients, however the detection of  $[Ca]_m$  transients required conditions of enhanced Ca cycling ( $\beta$ -adrenergic stimulation, elevated extracellular  $[Ca]$  or increased stimulation frequency). During  $\beta$ -adrenergic stimulation, the decay of  $[Ca]_m$  was  $\sim 2.5$ -fold slower than of  $[Ca]_i$ , leading to a stepwise accumulation  $[Ca]_m$  during rapid pacing. Remarkably, the upstroke of the  $[Ca]_m$  transient preceded the rise of  $[Ca]_i$ . This raises the question of possible contamination of the mitochondrial Ca signal by cytosolic rhod-2 as traces of the indicator dye could have remained in the cytosol even after dialysis. Nonetheless, directionally opposite effects on the  $[Ca]_m$  and  $[Ca]_i$  signals were elicited by inhibitors of mitochondrial Ca uptake or efflux, underpinning the mitochondrial origin of the signal and indicating that a significant fraction of the Ca released by the SR was buffered by the mitochondria with every beat. Maack et al. [66] also presented a new computational model that included mitochondrial microdomains in cardiomyocytes, where pulses of high  $[Ca]_{em}$  (10 or 20  $\mu M$ ) and 50 ms duration were simulated. This model predicted changes in  $[Ca]_m$  reminiscent of both, model I and model II described in this review. Similarly to model II, beat-to-beat  $[Ca]_m$  oscillations were simulated with extramitochondrial Ca pulses applied at 1 Hz, but with increasing frequency or amplitude of stimulation, diastolic  $[Ca]_m$  rose slowly until a new steady-state level was reached (reminiscent of model I). Recently beat-to-beat mitochondrial  $[Ca]_m$  transients were also reported using the Ca-sensitive inverse pericam Mitycam [137] targeted to mitochondria with a mitochondria-targeting sequence (subunit VIII of human cytochrome c oxidase) and adenovirally expressed in cardiomyocytes [138]. With the Mitycam probe sustained and transient phases of mitochondrial Ca signals were observed, which were dependent on  $[Ca]_i$  levels and required a functional MCU. Furthermore, in rat neonatal cardiomyocytes cytoplasmic Ca transients were reduced or enhanced by MCU overexpression or siRNA silencing, respectively, using novel targeted Ca probes. The data provide evidence that mitochondrial Ca uptake contributes to buffering of cytoplasmic Ca peaks in cardiomyocytes [139].

### 3.3. Fast versus slow mitochondrial Ca uptake: mutually exclusive or requirement for both?

Whether mitochondria can sequester Ca, even in large quantities, is not at debate. The controversies center around the kinetics and magnitude of mitochondrial Ca uptake. Arguments have been made in favor of or against beat-to-beat mitochondrial Ca uptake and the question of energetic cost efficiency of the translation of cytosolic Ca oscillations into oscillations of  $[Ca]_m$  have been raised [3, 9]. The mitochondrial Ca sink function that helps protect against cytosolic Ca overload requires high Ca buffering power, rather than fast uptake. From an ECC point of view, substantial beat-to-beat mitochondrial Ca sequestration would curtail the cytosolic Ca transient, and consequently, larger quantities of Ca are required to be shuttled, at increased energetic costs, between extracellular space, cytosol, SR



and mitochondria to achieve  $[Ca]_i$  levels necessary for contraction. However, rapid and frequent changes in metabolic demands of the working heart may indeed necessitate a rapid mitochondrial Ca response and fast mitochondrial Ca uptake to stimulate Ca-dependent dehydrogenases of the TCA cycle [67]. Future studies are likely to demonstrate that there is a role for both, rapid uptake in response to cytosolic Ca signals and a slow integration of changes in  $[Ca]_i$ . Indeed, consistent with this notion a recent study demonstrated two modes of operation of MCU, one with a proposed role for energetic signaling, the other serving as a cellular Ca sink [140] (see also section 4).

#### 4. Mitochondrial Ca and mPTP

Excessive mitochondrial Ca uptake and Ca accumulation bears the risk of mPTP activation [3], potentially leading to irreversible  $\Delta\Psi$  collapse, cessation of mitochondrial respiration and ultimately cell death. Activation of mPTP is facilitated by elevated levels of matrix  $[Ca]_m$  and ROS. We have recently shown that a mitochondrial Ca-dependent nitric oxide synthase [141] can become uncoupled and turn into a significant source of ROS, which together with enhanced mitochondrial Ca accumulation, significantly increased the risk of deleterious mPTP opening. According to the redox-optimized ROS balance model proposed by Aon et al. [142] mitochondria are required to operate in an intermediate redox state to maximize energy output and keep ROS generation minimal. In an attempt to define the relationship between total mitochondrial Ca uptake, changes in mitochondrial matrix free Ca, and mPTP activation, Wei et al. [140] measured changes in  $[Ca]_{em}$  and  $[Ca]_m$  during Ca uptake in isolated cardiac mitochondria and identified two components of Ca influx, termed  $MCU_{mode1}$  and  $MCU_{mode2}$ , with differential sensitivity to the MCU inhibitor Ru360, Ca transport kinetics and capacities, and Ca buffering associated with the respective pathway. Intra-mitochondrial Ca buffering has been assumed to be in the order of 100:1 (bound:free Ca) [108], however, according to Wei et al. [140] mitochondrial Ca buffering is highly dynamic, and the differential responses of  $[Ca]_m$  to Ca entry results from a two-component buffer system comprised of static Ca buffers and dynamic Ca buffering by phosphate that enters together with Ca. The authors interpreted their results that the role of  $MCU_{mode1}$  might be to modulate oxidative phosphorylation in response to intracellular Ca signaling, whereas  $MCU_{mode2}$  and a dynamic high-capacity Ca buffering system through calcium-phosphate complex formation constitute a Ca sink function. Furthermore, evidence was provided that the trigger for mPTP activation is unlikely to be  $[Ca]_m$  itself, but rather a downstream byproduct of total mitochondrial Ca loading. In agreement with this study separate components of mitochondrial Ca accumulations were also identified in brain and liver mitochondria that serve matrix dehydrogenase regulation, buffering of extramitochondrial free Ca, and mPTP activation [143].

#### 5. Inorganic polyphosphate - a mediator of Ca-dependent mPTP activation?

A critical link between mitochondrial Ca influx, mitochondrial Ca buffering and activation of the mPTP has recently been proposed by Seidlmayer et al. [144, 145]. In this study we tested the hypothesis that the adverse effect of mitochondrial Ca accumulation followed by mPTP opening is mediated by its interaction with inorganic polyphosphate (polyP), a polymer of orthophosphates (average length of 25 orthophosphates) that is found in cardiac mitochondria in significant amounts ( $280 \pm 60$  pmol/mg of protein). Depletion of polyP in mitochondria of rabbit ventricular myocytes led to significant inhibition of mPTP opening. This effect was observed when mitochondrial Ca uptake was stimulated by increasing cytosolic  $[Ca]_i$  in permeabilized myocytes mimicking mitochondrial Ca overload as observed e.g. during ischemia-reperfusion injury [19]. The results of this study indicate that inorganic polyP is a previously unrecognized major activator of mPTP and lend support to the hypothesis that the adverse effects of polyP might be caused by its ability to form stable

complexes with Ca and directly contribute to IMM permeabilization, raising even the possibility that a Ca/PolyP complex forms an integral part of the mPTP.

## 6. Conclusions

While substantial efforts were undertaken in recent years to characterize the properties and kinetics of mitochondrial Ca cycling, the experimental approaches and techniques used to date have essentially failed to reach unequivocal conclusions whether and how cardiac mitochondria respond to cytosolic Ca oscillations in a rapid beat-to-beat fashion. Despite experimental and technical limitations and shortcomings, it is needless to say that Ca is a key second messenger for the regulation of mitochondrial tasks and represents a crucial link for the role of mitochondria for excitation-metabolism as well as excitation-contraction coupling in the heart. Recent findings indicate that mitochondrial Ca transport capacities and kinetics are more complex and variable than previously assumed. The coexistence of rapid Ca uptake capabilities and slower, but high-capacity Ca buffering properties simply underpins the versatility of Ca signaling that is genuine for these organelles. Furthermore (and only discussed with a few illustrative examples in this review) the development of advanced computational models of energy metabolism, mitochondrial ion dynamics, redox signaling and mPTP regulation has significantly advanced the field and generated unprecedented novel insights into these complexities of mitochondrial dynamics [146–153]. Combined computational-experimental approaches [66, 154] will increasingly help overcome experimental limitations, validate computational approaches and generate new hypotheses on energy metabolism and mitochondrial signaling.

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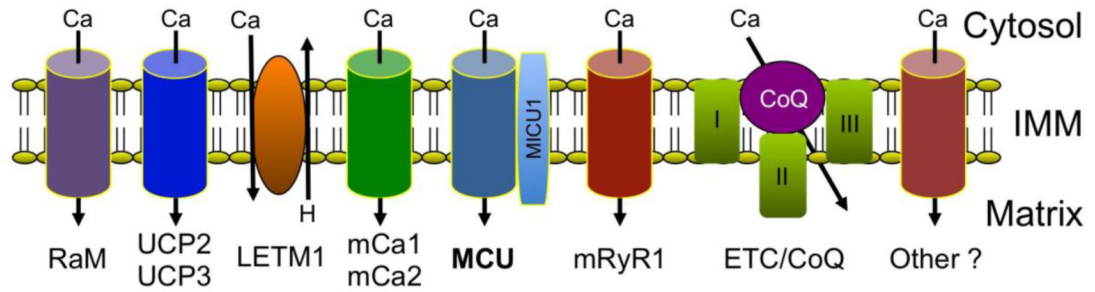
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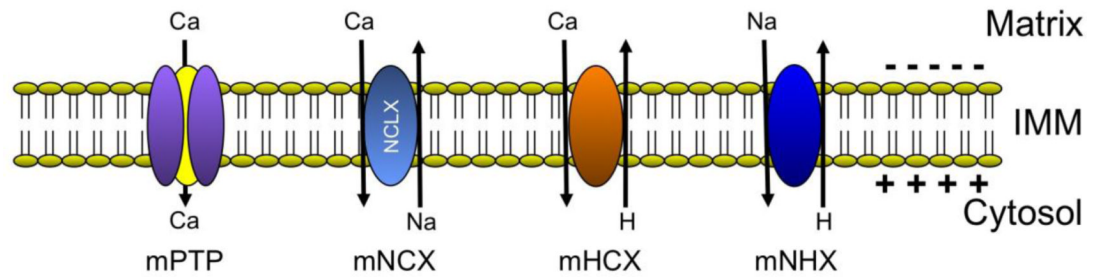
### Highlights

- Mitochondria occupy a third of the cell volume of cardiac myocytes
- Mitochondria have an elaborate system of calcium uptake and extrusion mechanisms
- Mitochondrial Ca signaling is crucial for excitation-metabolism-contraction coupling
- Mitochondria decode cytosolic Ca signals to respond to energy demands and buffer Ca
- Beat-to-beat transmission or slow integration of cytosolic Ca signals are debated

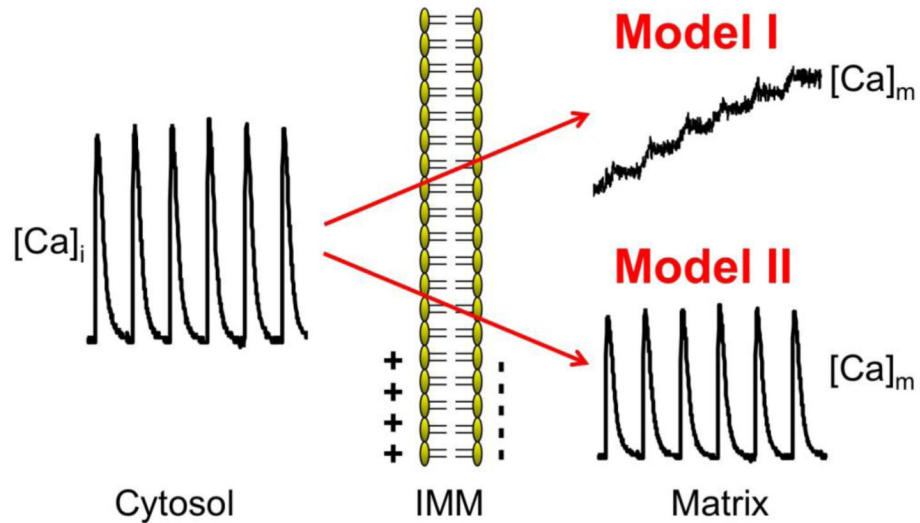
## A Ca influx



## B Ca efflux



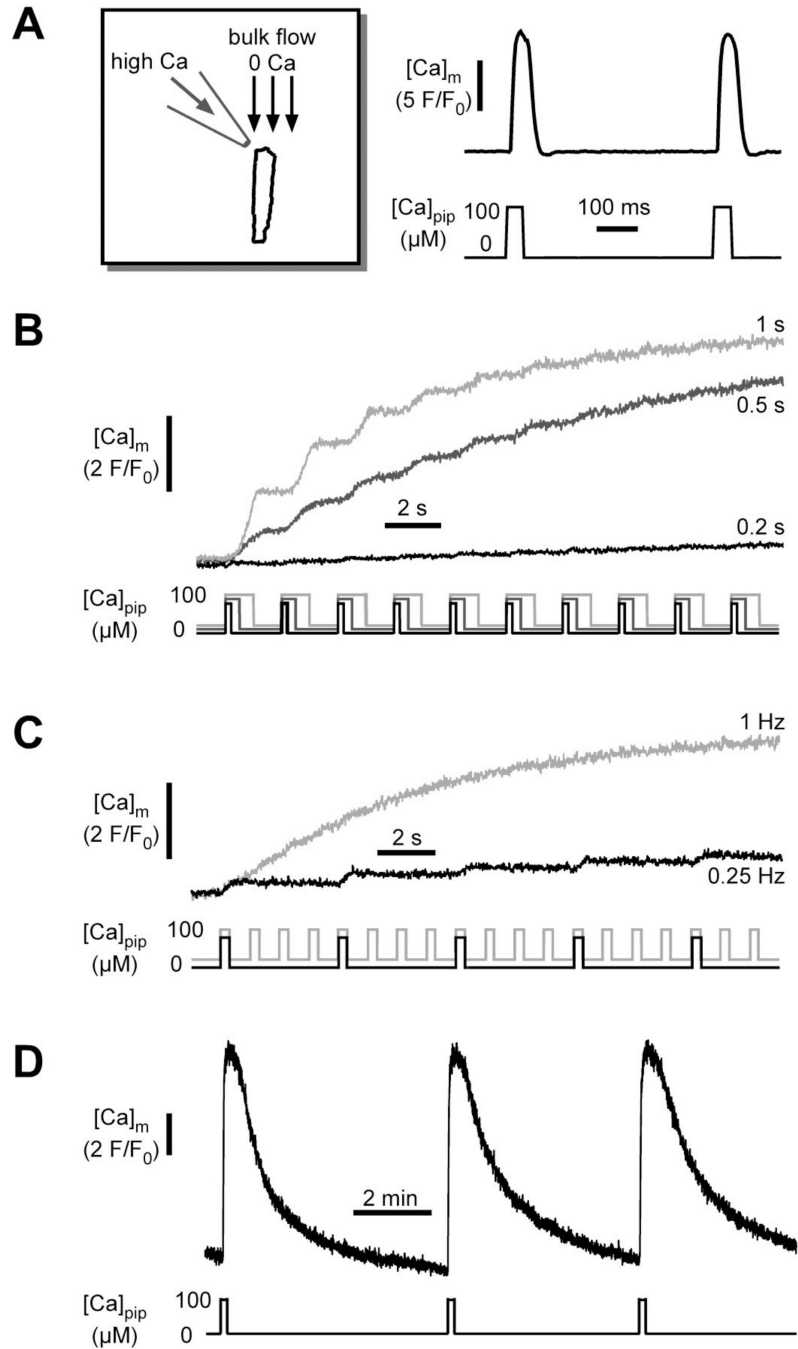
## C



**Fig. 1. Mitochondrial Ca transport and mitochondrial decoding of cytosolic Ca signals**

**A)** Mitochondrial Ca uptake mechanisms and pathways located at the inner mitochondrial membrane (IMM). From left: RaM, rapid mode of Ca uptake; UCP2 and UCP3, uncoupling proteins 2 and 3; LETM1, leucine-zipper-EF-hand-containing transmembrane protein 1; mCa1 and mCa2, Ca-selective IMM conductances 1 and 2; MCU, mitochondrial Ca uniporter, with the associated protein MICU1; mRyR1, mitochondrial ryanodine receptor type 1; CoQ, Coenzyme Q10; ETC, electron transport chain. **B)** Mitochondrial Ca extrusion mechanisms and pathways located at the IMM. mPTP, mitochondrial permeability transition pore; mNCX, mitochondrial Na/Ca exchange with NCLX as suggested molecular identity;

mHCX, mitochondrial proton/Ca exchange; mNHX, mitochondrial Na/proton exchange. C) Models of transmission of fast cytosolic Ca transients to matrix  $[Ca]_m$ : Model I, slow integration of cytosolic Ca spiking. Model II: rapid, beat-to-beat transmission of cytosolic Ca oscillations.



**Fig. 2. Measurements of  $[Ca]_i$  to  $[Ca]_m$  transmission in single permeabilized ventricular myocytes**

**A)** Technique used to simulate the effect of repetitive rapid cytosolic Ca transients on  $[Ca]_m$ . Mitochondria were loaded with the fluorescent Ca indicator fluo-3. Cells were placed in the laminar flow of Ca-free solution (1 mM EGTA) and exposed to computer controlled pressure ejections of a 100  $\mu$ M Ca containing solution from a micropipette. **B)** Effect of Ca pulse duration (0.2, 0.5 and 1 s) on  $[Ca]_m$  at constant stimulation frequency (0.5 Hz). **C)** Effect of Ca pulse frequency (0.25 and 1 Hz) on  $[Ca]_m$  at constant pulse duration (0.5 s). (Panels A, B and C from Sedova, Dedkova & Blatter, *Am. J. Physiol. Cell Physiol.*, 291(5): C840-50, 2006). **D)**  $[Ca]_m$  transients elicited by single Ca puffs applied at extended

intervals. Extracellular  $[Na]_o$  (20 mM) was set to allow for maximal Ca extrusion rates via mNCX.