

Involvement of the Sarcoplasmic Reticulum Calcium Pump in Myocardial Contractile Dysfunction: Comparison between Chronic Pressure-overload and Stunning

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Summary. Acute as well as chronic forms of heart failure involve mechanical dysfunction during systole and/or diastole. The rapid Ca^{2+} release from and Ca^{2+} reuptake into the tubuli of the sarcoplasmic reticulum are processes that critically determine normal systolic and diastolic myocardial function, which explains why in the last fifteen years so much attention has been paid to understand the performance of the sarcoplasmic reticulum Ca^{2+} pump during myocardial contractile dysfunction. In this communication we have reviewed the literature data on sarcoplasmic reticulum Ca^{2+} pump function in the chronically pressure-overloaded hypertrophied and stunned (post-ischemic reversibly injured) myocardium in the light of some new data from our laboratory. Results on the pressure-overloaded hypertrophied myocardium provide evidence that impaired relaxation is most likely due to a low capacity of the sarcoplasmic reticulum to pump Ca^{2+} , a consequence of a lower density of Ca^{2+} -pumping sites within the sarcotubular membranes. Contractile dysfunction in stunned myocardium is accompanied by an upregulation of the sarcoplasmic reticulum Ca^{2+} ATPase gene resulting in a slight increase of the Ca^{2+} pumping activity. The latter increase is likely an adaptive response of the reversibly injured myocardium which may contribute to the slow recovery of contractile function.

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Normal systolic and diastolic function of the heart requires the release and reuptake of Ca^{2+} . The contractile state of the heart can be altered by two mechanisms which are both Ca^{2+} dependent: changes in the availability of Ca^{2+} to the contractile proteins and changes in the responsiveness of the contractile proteins to activation by intracellular Ca^{2+} [1]. The availability of intracellular Ca^{2+} is regulated by the sarcolemma (SL) and sarcoplasmic reticulum (SR) and the Ca^{2+} responsiveness is controlled by the myofilaments and regulatory troponin-tropomyosin complex. Excitation-contraction (E-C) coupling is initiated when

Ca^{2+} channels are opened by depolarization of SL permitting Ca^{2+} to enter the cytoplasm. This small Ca^{2+} influx induces the release of a much larger quantity of activated Ca^{2+} from the intracellular stores in the SR [2,3]. The released Ca^{2+} interacts with troponin C, which is part of the regulatory complex of the myofilaments, thereby initiating cardiac contraction. Relaxation starts when Ca^{2+} is sequestered by the SR Ca^{2+} pump, so that Ca^{2+} dissociates from the contractile apparatus.

Abnormal handling of intracellular Ca^{2+} at any of the steps in E-C coupling can cause cardiac contractile dysfunction and leads to failure. The signalling function of Ca^{2+} demands a very low ionic concentration of Ca^{2+} inside the myocardial cells (about 10,000 fold lower than outside) and significant changes can therefore be achieved easily. During each depolarization only a very small amount of Ca^{2+} entering the cell needs to be extruded to prevent Ca^{2+} overloading of the myocytes [2]. The bulk of Ca^{2+} released from the SR must be reuptaken to its site of origin (SR) in order to be released during the next cardiac cycle. Several transport systems, including the SR Ca^{2+} pump, have now been documented in heart membranes. These systems are based on 4 basic transport modes: 1) ATP driven (SL and SR Ca^{2+} pumps); 2) Na/ Ca^{2+} antiporters (SL and mitochondria); 3) Ca^{2+} channels (SL and SR) and 4) Ca^{2+} uniporter (mitochondria) [4]. These systems have either a low or a high Ca^{2+} affinity, thus serving different purposes in

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the various phases of the cardiac cycle. Overall, the SR Ca^{2+} pump can be considered to be the transport system that presides over the rapid and fine regulation of intracellular Ca^{2+} linked to the contraction/relaxation cycle.

Acute and chronic forms of heart failure involve mechanical dysfunction during systole and/or diastole [1,9]. Following a brief period of myocardial ischemia, insufficient to cause cell death, myocardial function may remain depressed for several hours to days ("myocardial stunning"), which is a form of reversible myocardial dysfunction. Among the cellular organelles whose altered function might account for the contractile dysfunction are the SR, the contractile filaments and the SL. To date the results of several studies point toward a decrease in the responsiveness of the myofilaments for Ca^{2+} and/or lack of Ca^{2+} available from SR as primary cause of the abnormal E-C coupling in "stunned" myocardium [5–7]. On the other hand, myocardial hypertrophy, which is one of the most important responses of the heart to chronic pressure-overloading, is an adaptive process that enables the heart to compensate for the decrease in cardiac performance. Cells of the hypertrophied, failing myocardium are not normal [1,8,9], however, and it is now apparent that hypertrophy of the overloaded heart is a complex process that ultimately becomes detrimental after initially exerting a beneficial effect. Disturbed relaxation in early experimental cardiac hypertrophy has been noted for some time [9], but it is not until the last decade that the importance of relaxation abnormalities was clearly demonstrated in patients with congestive heart failure [11]. A slower rate of the SR Ca^{2+} pump can explain the impaired relaxation in the hypertrophied heart, but an increase in diastolic stiffness by spontaneous recycling of Ca^{2+} in the SR during diastole must also be considered [8].

SR Ca^{2+} Pump Characteristics of the Normally Functioning Myocardium

SR membranes contain a Ca^{2+} stimulated Mg^{2+} -dependent ATPase, which represents up to 90% of the total protein of the membrane, and which transports Ca^{2+} from the cytosol into the vesicles of the SR with a high velocity and affinity for Ca^{2+} . If one takes the total amount of SR in the heart and combines it with its transport characteristics, it is clear that the amount of Ca^{2+} needed for the cardiac contraction/relaxation cycle can be adequately met by the organelle [2,4]. The pump protein with a molecular weight of about 115 kDa, is distributed asymmetrically across the membrane and consists of dimers of a single polypeptide. For each mole of ATP hydrolyzed, two Ca^{2+} ions are transported into the SR vesicles. Until recent years the major problem regarding the SR function in the entire cardiac cycle was the not yet known

mechanism by which Ca^{2+} is released from the SR in the heart. The pieces of the machinery that couple depolarization to SR Ca^{2+} release, in particular the isolation of the SR Ca^{2+} release channel of 565 kDa molecular weight, have now been identified during recent years [3]. At present, it is believed that the capacity of SR to actively take up Ca^{2+} is present throughout the cardiac cycle. Initially, the small amount of Ca^{2+} entering the cell during depolarization triggers a transient release of Ca^{2+} to exceeding the SR Ca^{2+} pumping rate (Figure 1). When the intracellular free Ca^{2+} rises to saturating Ca^{2+} levels which are sufficient for the activation of myofilaments, the SR Ca^{2+} release channel becomes inactivated [2]. Phospholamban, which is an integral membrane protein of cardiac SR, modulates the Ca^{2+} pump function of cardiac SR [12–14]. When phosphorylated by cyclic AMP-dependent protein kinase or by Ca^{2+} -calmodulin dependent kinase, phospholamban stimulates the Ca^{2+} pump ATPase, predominantly by increasing the affinity for Ca^{2+} . Phospholamban has been purified and its primary structure has been determined by cDNA cloning and sequencing [14]. It consists of five identical monomers of 5.2 kDa and the stoichiometric ratio between phospholamban pentamer and Ca^{2+} pump ATPase is about 1 [14]. Phosphorylation of phospholamban relieves its inhibitory effect on the Ca^{2+} pumping ATPase by increasing the maximal rate as well as the Ca^{2+} affinity of the ATPase. The phosphorylation sites are serine 16 and threonine 17, catalyzed by cyclic AMP- and Ca^{2+} -calmodulin-dependent protein kinase, respectively [14, 15]. Thus, there are two phosphorylation sites activating phospholamban and their stimulatory effects on the ATPase are additive. Similar additive effects of cyclic AMP and Ca^{2+} -calmodulin dependent SL protein phosphorylation on the SL Ca^{2+} pumping ATPase have already been observed earlier [16].

Alterations in SR Ca^{2+} Pump in the Pressure-Overloaded, Hypertrophied Heart

In the mid-seventies many investigations were undertaken to correlate possible modifications of SL and SR function to the altered contractile properties of hypertrophied and failing hearts [17–20]. Myocardial hypertrophy was experimentally induced by various types of chronically increased workload such as aortic constriction, valvular insufficiency, and hypertension. A common observation in these studies was the reduced rate of ATP-dependent Ca^{2+} uptake in isolated SR membrane vesicles from hypertrophied hearts [21–23]. In the meantime it was firmly established that the Ca^{2+} pump in SR in vitro could be stimulated by cyclic AMP-dependent phosphorylation. Data on cyclic AMP levels in the hypertrophied myocardium were not available, but in one study it was hypothesized that the decreased Ca^{2+} accumulation in cardiac

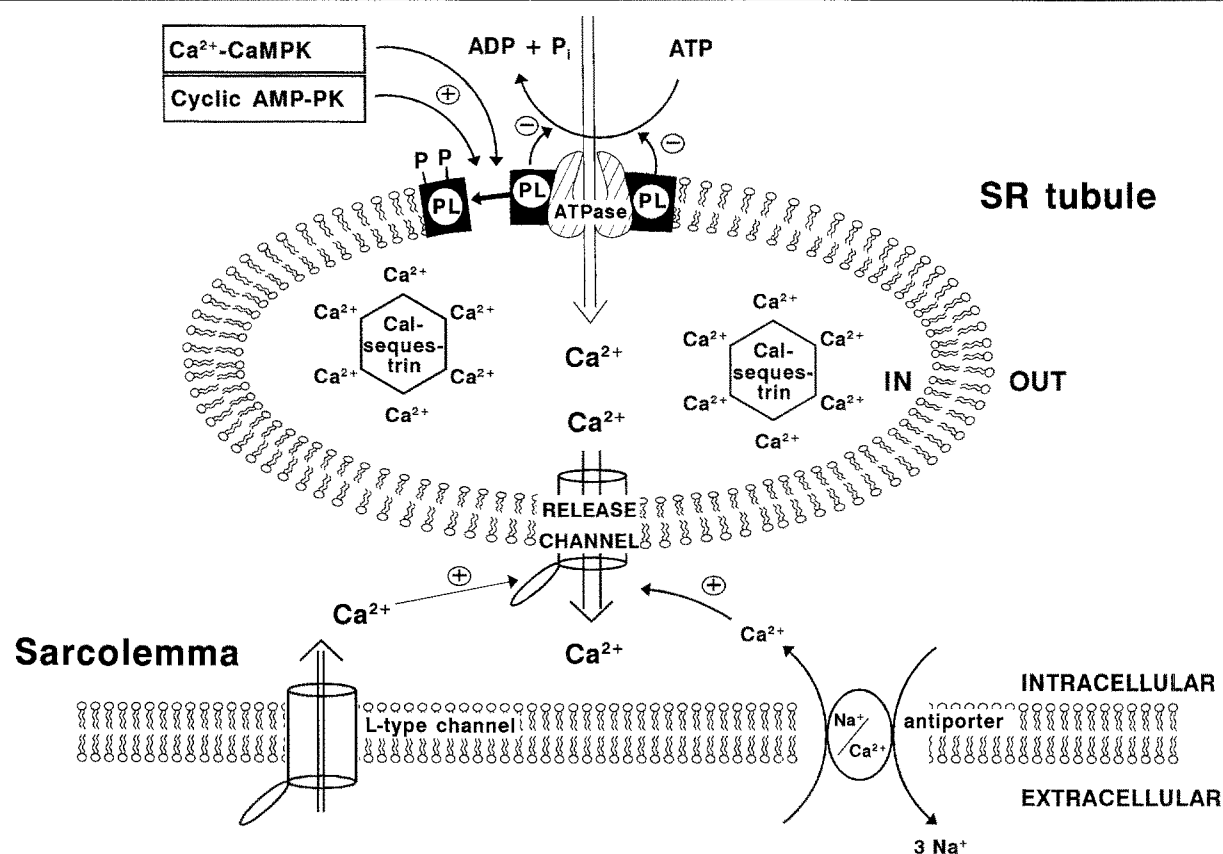


Fig. 1. Concerted operation of the second-messenger regulated Ca^{2+} pump and the Ca^{2+} release channel in SR and the depolarization-induced Ca^{2+} influx pathways through the SL. Notice the relative thickness of the arrows indicating the differences in amounts of Ca^{2+} moving during the phase of depolarization. Ca^{2+} is actively transported into the SR by the Ca^{2+} pump of which the activity can be modulated by protein kinases. Phosphorylation of phospholamban (PL-P) by Ca^{2+} -calmodulin- (Ca^{2+} -CaMPK) and cyclic AMP- (cyclic AMP-PK)-dependent protein kinases switches off the partial inhibition exerted on the Ca^{2+} pump by its unphosphorylated form (PL). The Ca^{2+} release channel is activated by a small rise in cytosolic free Ca^{2+} originating from the depolarization-induced Ca^{2+} influx through the L-type Ca^{2+} channel and the $\text{Na}^{+}/\text{Ca}^{2+}$ antiporter in the SL.

SR from spontaneous hypertensive rats was due to reduced cyclic AMP-dependent protein kinase activity [24]. This postulation prompted us to study the functional changes in microsomes from hypertrophied heart and to investigate whether these changes originate from an altered degree of phospholamban phosphorylation. Left ventricular hypertrophy was induced by clamping the aorta of rabbits for one month [25], in which period left ventricular mass increased 1.6 fold ($p < 0.01$ versus sham-operated animals). In vitro characterization of enriched SR membrane vesicles isolated from control and hypertrophied left ventricle are shown in Table 1. The yield of SR vesicles from control and hypertrophied myocardium was similar for the control and the hypertrophied hearts. The activity of the SR marker-enzyme rotenon-insensitive NADPH cytochrome c reductase was measured as reference for the rate of SR Ca^{2+} pumping. The specific activity of the Ca^{2+} pump proved to be markedly reduced during hypertrophy, whereas the activity of the marker enzyme rotenon-insensitive NADPH cy-

tochrome c reductase remained relatively constant (Table 1). The decrease in Ca^{2+} uptake rate was not due to increased Ca^{2+} release channel activity during the in vitro incubation, because inclusion in the assays of the release channel inhibitor ruthenium red, did not alter the results [26]. Incubation of the SR membranes from control hearts with 10 μM cyclic AMP, 50 μM [γ -³²P] ATP and 2-6 $\text{mU}\cdot\text{ml}^{-1}$ purified cyclic AMP-dependent protein kinase led to enhancement of the ³²P incorporation into SR proteins of which phospholamban was the predominant one [12]. Indeed, prior phosphorylation of SR proteins under the latter conditions increased the Ca^{2+} uptake rate by 1.4 fold in both the control and the hypertrophied hearts (Table 1). This finding indicated that an alteration in the phosphorylation degree of phospholamban was not responsible for the decrease in the rate of SR Ca^{2+} uptake by the hypertrophied heart. Likewise, it was shown that in the failing human heart, the second messenger control of SR Ca^{2+} ATPase was unaffected [27]. In a number of subsequent papers in which hu-

Table 1. Yield, enzymatic and Ca^{2+} transport activities of enriched SR membranes from control and hypertrophied rabbit hearts

	n ^a	Control	Hypertrophied
Yield			
(mg protein.g.tissue ⁻¹)	5	1.46 ± 0.18	1.72 ± 0.14
NADPH cyt c reductase			
(nmol.min. ⁻¹ .mg protein ⁻¹)	5	8.17 ± 0.11	7.77 ± 0.05
Ca^{2+} uptake			
(nmol.min. ⁻¹ .mg protein ⁻¹)			
control	5	28.0 ± 1.8	19.5 ± 1.0 ^b
protein kinase-treated	5	37.5 ± 2.6	27.8 ± 2.7 ^b
³² P incorporation			
(pmol.mg protein ⁻¹)			
control	3	91	—
protein kinase-treated	3	423	—

Left ventricular hypertrophy was produced in New Zealand white rabbits of either sex by constriction of the descending aorta as described before [20,25]. Briefly, the descending aorta at the level of the diaphragm was exposed through the abdominal cavity under anesthesia and sterile conditions. A clip was placed just proximal of the coeliac artery. When animals were killed after one month bodyweights of the sham-operated (3.55 ± 0.11 kg) animals were not different from the animals with aorta constriction (3.49 ± 0.08 kg). But the left ventricle + septum weight (relative to the body weight) was significantly higher (1.75 ± 0.06 versus 1.34 ± 0.03 fold) in the animals with aorta constriction. Enriched SR membranes were isolated from the left ventricle by differential centrifugation [25]. Analysis of rotenon-insensitive NADPH cyt c reductase, ATP-dependent Ca^{2+} uptake and ³²P incorporation into total SR protein, was performed using spectrophotometry (wavelength 340 nm), radioactively labelled ⁴⁵Ca and γ -³²P-labelled ATP, respectively. For details of protein kinase treatment is referred to the text. Details of the incubation conditions for each enzymatic and transport assay can be found elsewhere [25]. ^an = number of rabbits in each group and ^bp < 0.05 versus control animals.

man, rabbit and rat hearts were studied [28–30], the unresolved issue of the cause of the reduction in the rate of SR Ca^{2+} uptake rate during hypertrophy was readdressed. For instance, in severe myocardial hypertrophy, the concentration of the SR Ca^{2+} ATPase messenger RNA (mRNA) decreased to about 70% and that of the protein to 80% of the sham-operated animals, whereas the total left ventricular mRNA and Ca^{2+} ATPase protein was either unchanged or slightly increased [30]. Similarly, a marked decrease (up to about 50% of controls) in SR Ca^{2+} ATPase mRNA was found in human end-stage heart failure [29]. Since there are different isoforms of the SR Ca^{2+} ATPase and their expressions are specifically and developmentally regulated, some studies looked for possible changes in the expression pattern of isoforms at the mRNA level using S1 nuclear mapping analysis and specific cDNA probes for the rat heart SR Ca^{2+} ATPase mRNA. In one of these studies, the 5'-end sequence (1.1 kb fragment) and a 3'-portion (2.4 kb fragment) of the cardiac SR Ca^{2+} ATPase were shared by the normal and the pressure-overloaded hypertro-

phied rat heart [30]. These data indicate that the same Ca^{2+} ATPase mRNA isoenzyme is expressed in both normal and hypertrophied rat myocardium. By using similar molecular biological techniques, it was demonstrated that the same SR Ca^{2+} ATPase is expressed in control and pressure-overload hypertrophied rabbit hearts [28]. Cardiac tissue, whether hypertrophied or normal, expresses the slow-twitch SR Ca^{2+} ATPase isoform (SERCA 2) [9,28,31]. Messenger RNA levels of SERCA 2 isoform of SR Ca^{2+} ATPase were decreased in failing hearts explanted from patients with a dilated cardiomyopathy who underwent a cardiac transplantation [9]. These results indicate that in severe myocardial hypertrophy induced by chronic pressure-overload, the decrease in the SR Ca^{2+} pump activity is not due to an altered second messenger control or an isoenzyme shift. It is more likely that the expression of the SR Ca^{2+} ATPase gene does not follow the overall increase in gene expression that results in cardiac growth. The slower rate of expression of the SERCA 2 gene leads to a decrease in density of Ca^{2+} pump protein molecules embedded into the SR membrane and thus to an impaired Ca^{2+} handling of the hypertrophied myocardium.

Alterations in the SR Ca^{2+} Pump Function in the Ischemic and "Stunned" Myocardium

The effects of ischemia on SR Ca^{2+} pump and release channel have been previously studied with whole cardiac muscle homogenates and isolated SR vesicle fractions [32–38]. Assay of enzymatic activities in both tissue fractions suggested that SR Ca^{2+} uptake and Ca^{2+} ATPase activities are impaired to a similar degree in the ischemic heart. Some of the loss was suggested to have occurred during membrane isolation or to be due to selection of a subpopulation of damaged SR vesicles [36]. However, in several studies parallel changes in SR Ca^{2+} uptake rate in unfractionated homogenates and isolated SR vesicle fractions have been found, ruling out the possibility that assays were performed with selected SR vesicle populations [37,39]. Ca^{2+} release channel inhibitors (ryanodine and ruthenium red) were used to distinguish between ischemic damage to the SR Ca^{2+} release channel and SR Ca^{2+} pump. Periods of ischemia lasting 15 min and 60 min depressed oxalate-supported SR Ca^{2+} pumping rates of rat heart homogenate by 50% and 80%, respectively [39]. No decrease was observed, however, when the Ca^{2+} uptake rate was measured after Ca^{2+} release channel closure by ruthenium red or ryanodine [39]. Therefore it was assumed that ischemia primarily affects the function of the Ca^{2+} release channel, although more recent results do not support this hypothesis [40]. The decrease in SR Ca^{2+} uptake rate usually appears after 10–15 min of ischemia

[33,34,36]. Such a brief period of ischemia does not lead to irreversible myocardial damage, but upon reperfusion, recovery of contractile function is delayed ("stunned myocardium") [5,7,41–45]. In the following paragraphs we present some data characterizing the SR Ca^{2+} pump function in stunned myocardium.

Myocardial stunning

The molecular mechanisms contributing to myocardial stunning are poorly understood, but it is likely that the slow recovery of contractile function is based upon the slow recovery of ischemia-induced changes at molecular function levels. In a recent report, Opie described as the two likely causes of stunning during the very early reperfusion phase excess of cytosolic Ca^{2+} and the formation of free radicals [49]. For instance, it was shown that intracoronary Ca^{2+} infusions during stunning could restore the contractility [47]. Krause *et al.* were the first to demonstrate in a canine model that SR membranes isolated from stunned canine myocardium had a decreased ability to actively transport Ca^{2+} , which suggested a failure of Ca^{2+} handling at the level of the SR [7]. However, Kaplan *et al.* have recently shown, that neither an altered SR uptake of Ca^{2+} nor an abnormal function of SR Ca^{2+} release channels contribute to the contractile dysfunction of stunned myocardium in rabbits [48]. Others have hypothesized that especially during the late reperfusion phase, a decreased Ca^{2+} sensitivity of myofilaments is involved in the contractile dysfunction, although mechanism(s) involved in the alteration of Ca^{2+} activation of contractile apparatus were not provided [5]. This hypothesis of decreased Ca^{2+} sensitivity is supported by a recent study from our laboratory in which we have shown that the Ca^{2+} sensitizing agent EMD 60263, which is devoid of phosphodiesterase inhibitor properties, restored segment length shortening and mechanical efficiency of stunned porcine myocardium, while exerting a negligible effect on the not stunned myocardium [49].

In the porcine model of myocardial stunning (two periods of 10 min coronary artery occlusion separated by 30 min reperfusion) we and others have shown that reperfusion leads to the induction of several genes encoding proto-oncogenes [50,51], Ca^{2+} binding proteins [52] and heat shock proteins (HSPs) [45,53–59]. It has been observed that brief periods of myocardial ischemia and reperfusion do not only cause prolonged contractile dysfunction, but also increase the tolerance for irreversible damage during a subsequent sustained episode of ischemia ("ischemic preconditioning"). It is not yet clear whether the upregulation of these genes contribute to the molecular events involved in ischemic preconditioning. Evidence is accumulating that HSPs have a protective effect on the infarct size limitation which occurs when a long lasting coronary artery occlusion is separated from a brief period of ischemia by a period of 24 h [59]. Elevated levels of HSPs and ubiquitin are believed to have pro-

TECTIVE/adaptive effects [45,53]. However, studies in mammalian cell cultures indicate that the increased synthesis of HSP-27 under conditions of severe stress lead to the disruption of cellular Ca^{2+} homeostasis [60]. In the present work (Figure 2 and [52,65]) the changes in Ca^{2+} pump function are evaluated in relation to the contractile dysfunction during myocardial stunning. Therefore, we examined in the normal and stunned myocardium the steady state mRNA levels of cardiac specific SR Ca^{2+} -ATPase isoform in parallel with SR Ca^{2+} pump function of isolated SR membrane vesicles.

Expression of SR Ca^{2+} -ATPase gene and SR Ca^{2+} uptake activity in stunned reperused myocardium

Cardiac specific SR Ca^{2+} -ATPase mRNA expression pattern was measured by hybridizing the total RNA preparations derived from stunned (90 min) and not stunned myocardium. In stunned (E) myocardium mRNA expression of SR Ca^{2+} -ATPase was significantly enhanced (1.76 ± 0.28 assessed fold by densitometric analysis, $n = 4$) as compared to the not stunned myocardium (Figure 2). A single mRNA species (3.7 kb) migrating just below to the 28S ribosomal RNA hybridized to the cardiac specific SR Ca^{2+} -ATPase cDNA probe [63]. No differences in mRNA levels were seen between stunned and not stunned tissue fractions obtained in experiments carried out with sham-operated animals [52]. Apart from the SR Ca^{2+} -ATPase, other Ca^{2+} transport regulating proteins like phospholamban, calsequestrin and calmodulin also contribute to the maintenance of Ca^{2+} homeostasis [2,4,13,18]. In a detailed study on various Ca^{2+} binding proteins using the same porcine model, we have shown that SR Ca^{2+} -ATPase gene expression did not alter during the first sequence of 10 min occlusion and 30 min reperfusion. After the second occlusion-reperfusion sequence the SR Ca^{2+} -ATPase mRNA levels in stunned myocardium were even reduced but recovered during the further reperfusion period and reached maximal levels at 90 min [52]. The expression of glyceraldehyde-3-phosphate dehydrogenase (GAPDH), a house keeping gene, was used as a standard and to check the integrity of RNA as well as loading differences on the gel. As expected, GAPDH expression did not change during our stunning protocol or even decreased as reported earlier for more severe ischemia leading to myocardial infarction [62]. Hence, our data on the upregulation of SR- Ca^{2+} -ATPase gene expression during late reperfusion may be an adaptive and temporary phenotypic alteration in the stunned myocardium [52]. The increased transcriptional activity of SR Ca^{2+} -ATPase suggests the presence of a repair process at the SR level in the stunned myocardium [64].

In order to compare the mRNA expression data with the SR Ca^{2+} pump function, we also examined ATP dependent Ca^{2+} uptake and cyclic AMP depen-

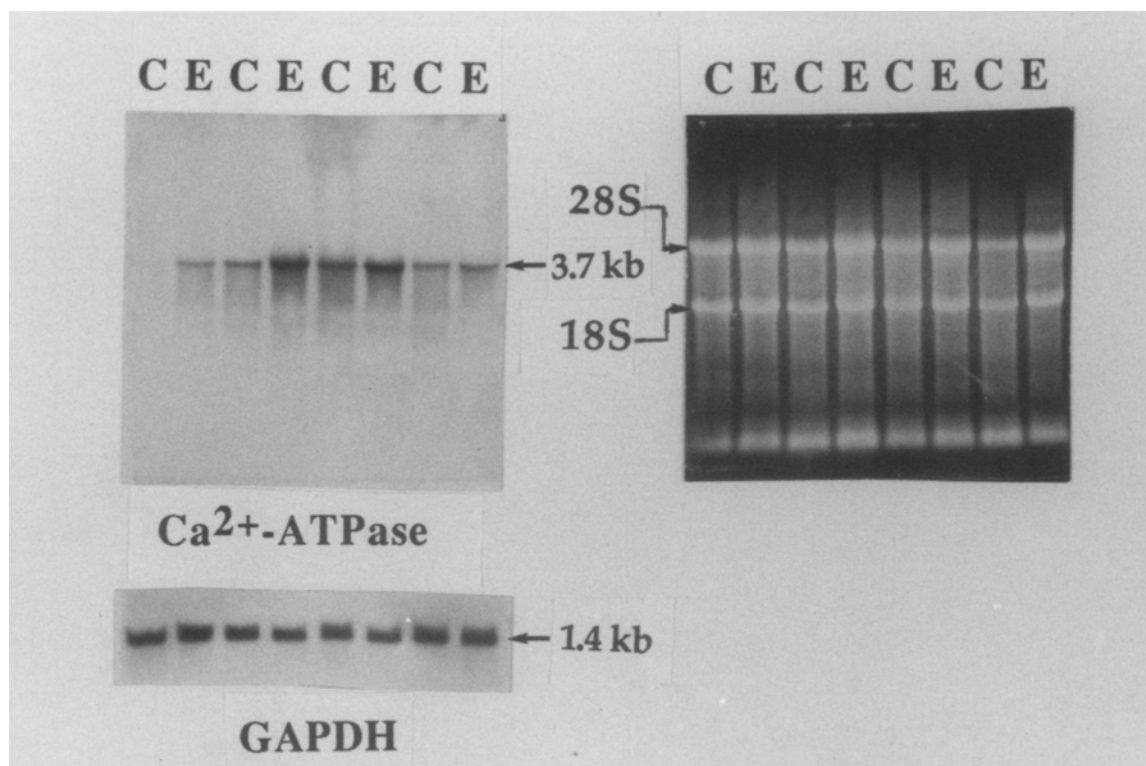


Fig. 2. Expression of SR Ca²⁺-ATPase gene in stunned and not stunned myocardium. Northern blot hybridization showing the expression of cardiac specific SR Ca²⁺-ATPase mRNA signals in stunned (experimental, E) and not stunned (control, C) of four separate experiments. To induce stunning in pigs the left anterior descending coronary artery was occluded for 10 min and reperfused for 30 min, then again occluded for 10 min and reperfused for 90 min. At the end of the second reperfusion (90 min) the pigs were sacrificed and myocardial tissues from the left anterior descending (E) and left circumflex coronary artery (C) regions were excised and processed for Northern hybridization using rat specific cardiac SR Ca²⁺-ATPase cDNA probe. A single mRNA species of 3.7 kb was hybridized to the radioactive cDNA probe [63]. On right panel ethidium bromide stained agarose gel is shown to depict the quality and quantity of total cellular RNA loaded on the gel. The lower panel shows the GAPDH signal (1.4 kb) as a standard and to compare the induction of Ca²⁺-ATPase gene in stunned (E) vs not stunned (C) myocardium.

dent phosphorylation of phospholamban in isolated SR membrane vesicles of stunned reperfused (30 min) and not stunned regions. Ca²⁺ uptake by the SR was increased by 17% in the fractions isolated from stunned myocardium as compared to the control [65]. No differences in Ca²⁺ uptake rate were seen between stunned and not stunned tissue fractions obtained from sham-operated animals (not shown). In the presence of exogenous cyclic AMP dependent protein kinase the amount of ³²P incorporation into phospholamban remained unaltered in both ventricular regions [65]. Our results on SR Ca²⁺-ATPase mRNA levels and SR Ca²⁺ uptake activity, although not measured at the same time of reperfusion (90 and 30 min, respectively), strengthen our hypothesis that the SR Ca²⁺ pump may temporarily be upregulated by a mechanism not related to its regulation by second messengers.

We also observed that phospholamban was not upregulated in stunned myocardium which indicates that the stoichiometric phospholamban/Ca²⁺-ATPase subunit ratio favors a less inhibited state of the

SR Ca²⁺ pump [52]. In the same porcine model the steady-state tissue levels of mRNAs encoding proto-oncogenes [50,51], HSP-27, ubiquitin [45] and HSP-70 are enhanced [53]. Interestingly, α - and β -myosin heavy chain mRNA expressions did not alter in stunned myocardium in our porcine model [52], which suggests that repair processes occur predominantly at the membrane proteins. Our data support the hypothesis that brief episodes of ischemia cause adaptive alterations at the molecular level indicated by an enhanced transcriptional activity of stunned myocardium which may contribute to repair processes during recovery of contractile function.

Although the above described model does not lead to cell death [54], some reversible damage to intracellular Ca²⁺ transport regulatory proteins probably occurred, because we observed a compensatory increase of their mRNA expression [52]. These proteins could be targets of reversible injury which at the same time may play a crucial role in the restoration of contractile function. Upregulation of the SR Ca²⁺-ATPase gene

in response to ischemia and reperfusion points toward its involvement in mechanisms underlying myocardial stunning. Furthermore, we also like to postulate that different stress conditions alter the expression pattern of several sets of genes, and their combined actions will influence the phenotype of cardiomyocytes during stunning.

Conclusions

We reviewed the available literature data on SR Ca^{2+} pump mRNA levels and Ca^{2+} pump protein function in the chronically pressure-overloaded hypertrophied and post-ischemic stunned myocardium in the light of some new data of our laboratory. Results on pressure-overloaded hypertrophied myocardium provide convincing evidence that the impaired relaxation of the hypertrophied heart is likely due to a lower capacity of the SR to pump Ca^{2+} ions as a result of a lower density of Ca^{2+} pump protein molecules within the SR membrane. Apparently, the expression of the SR Ca^{2+} -ATPase gene does not follow the overall increase in gene expression that results in cardiac hypertrophy, rather its expression declines. As far as the contractile dysfunction in ischemic and stunned myocardium is concerned the available data do not point toward a critical involvement of changes in the SR Ca^{2+} pump function during hypo-contractility. Surprisingly, we observed enhanced expression of the Ca^{2+} -ATPase gene accompanied by a slight increase in the SR Ca^{2+} pump activity in stunned myocardium. It is therefore postulated that this increase is an adaptive response of the reversibly injured myocardium which may contribute to the slow process of recovery of contractile function.

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