Temperature Threshold and Preservation of Signaling for Mitochondrial Membrane Proteins during Ischemia in Rabbit Heart¹

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Temperature modulates both myocardial energy requirements and production. We have previously demonstrated that myocardial protection induced by hypothermic adaptation preserves expression of genes regulating heat shock protein and the nuclear-encoded mitochondrial proteins, the adenine nucleotide translocator isoform 1 (ANT₁), and the β subunit of F₁-ATPase (β F₁-ATPase). This preservation is associated with a reduction in ATP depletion similar to that noted in cardioplegic arrested hearts preserved at a critical temperature (30°C) or below. We tested the hypothesis that expression of these genes may also be subject to this temperature threshold phenomenon. Isolated perfused rabbit hearts were subjected to ischemic cardioplegic arrest at 4, 30, or 34°C for 120 min. Cardiac function indices and steady-state mRNA levels for ANT₁, β F₁-ATPase, and HSP70-1 were measured prior to ischemia (B) and after 45 min of reperfusion. Cardiac function was significantly depressed in the 34°C group. Ischemia at 34°C reduced steady-state mRNA levels for ANT₁ and β F₁-ATPase from B, but these levels were similarly preserved at 4 and 30°C. HSP70-1 levels were mildly elevated (fourfold) above B to similar levels at all three temperatures. These results indicate that mRNA expression for ANT₁ and β F₁-ATPase is specifically preserved in a pattern consistent with the temperature threshold phenomenon. HSP70-1 expression is not influenced by ischemic temperature. Preservation of gene expression for these mitochondrial proteins implies that signaling for mitochondrial biogenesis or resynthesis is maintained after ischemic insult. © 1998 Academic Press

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Temperature protects ischemic myocardium (2, 3, 6, 7, 15, 34, 36, 38) and modulates both myocardial energy requirements and production (27). The relationship between energy supply and demand in the cardioplegic arrested heart remains fairly constant in the temperature range between 4 and 30°C (27), which is frequently used during cardiac surgical procedures. The myocardial supply/demand ratio demonstrates a threshold phenomenon by deteriorating rapidly at temperatures above 30°C (27). Accelerated ATP depletion and inability to reestablish highenergy phosphate stores during reperfusion accompany this deterioration. Thus, 30°C represents a critical temperature point (39), which

triggers acceleration of metabolic energy utilization rate above the energy production rate. Provision of glucose as substrate in the cardioplegia solution can partially ameliorate this metabolic imbalance by promoting anaerobic ATP production (27).

Recently, we have demonstrated that hypothermic exposure at 30°C prior to warm ischemia preserves myocardial function and ATP stores after reperfusion (28). This response represents an adaptation, whereby the heart becomes tolerant to a second exposure of the same or alternative type of stress. Hypothermic exposure induces a metabolic down-regulation, which persists during subsequent warm ischemia and may represent the principal factor in this down-regulation. This response differs from simple cold exposure during ischemia. Consistent with other forms of coldinduced adaptation (18, 20, 21), alterations in expression of specific genes occur. These include

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preservation of steady-state mRNA levels for genes regulating heat shock proteins and the nuclear-encoded mitochondrial membrane proteins, the adenine nucleotide translocator (ANT), and the β subunit of F₁-ATPase (β F₁-ATPase). This preservation is associated with a reduction in ATP depletion (28) that is similar to that noted in cardioplegic arrested hearts preserved at 30°C or below. Accordingly, we posed the hypothesis that expression of these genes may be subject to this temperature threshold phenomenon. Ischemic temperatures above 30°C may affect expression of these genes. To test this hypothesis steady-state mRNA levels were measured after reperfusion in rabbit hearts subjected to different ischemic temperatures during cardioplegic arrest. These data show that gene expression for the mitochondrial membrane proteins is consistent with the temperature threshold hypothesis. However, HSP70-1 gene expression does not vary according to ischemic temperature.

MATERIALS AND METHODS

Preparation of Isolated Heart

White New Zealand rabbits (male or female, 2.3-2.7 kg body wt) were heparinized (700 U/kg, intravenously) and anesthetized with sodium pentobarbital (45 mg/kg, intravenously). Under median sternotomy, the heart was rapidly excised and immersed momentarily in ice-cold physiological salt solution (PSS), pH 7.4, containing 118.0 mmol/L NaCl, 4.0 mmol/L KCl, 22.3 mmol/L NaHCO₃, 11.1 mmol/L glucose, 0.66 mmol/L KH₂PO₄, 1.23 mmol/L MgCl₂, and 2.38 mmol/L CaCl₂ (27, 28). The aorta was cannulated in the Langendorff mode and the heart was perfused with PSS equilibrated with 95% 02-5% CO2 at 37°C. Perfusion pressure was maintained at 90 mm Hg and PSS passed twice through 3.0-µm-pore size filters. An incision was made in the left atrium, and a fluidfilled latex balloon was passed through the mitral orifice and placed in the left ventricle. The balloon was connected to a pressure transducer for continuous measurement of left ventricular pressure (LVP) and the first derivative of LVP (dP/dt). The caudal vena cavum, the left and

right cranial vena cava, and the azygous vein were ligated. The pulmonary artery was cannulated for collection of coronary effluent. The coronary flow was measured with a flow meter (T201, Transonic Systems Inc., Ithaca, NY, U.S.A.).

The analog signals of left ventricular mechanical function were continuously recorded on a pressurized ink chart recorder (Gould, Inc., Cleveland, OH, U.S.A.), and an on-line computer (Macintosh, Biopac Analog Signal Acquisition System). To characterize cardiac function, developed pressure (DP) is defined as peak systolic pressure minus end-diastolic pressure. To provide an estimate of myocardial work, the product of DP and heart rate (HR) was calculated (PRP, mm Hg/min). Myocardial oxygen consumption (MV_{O2}) was calculated as the expression $MV_{O_2} = CF \times [(Pa_{O_2} - Pv_{O_2}) \times$ (c/760)], where CF is coronary flow (ml/min), $(Pa_{O_2} - Pv_{O_2})$ is the difference in the partial pressure of oxygen (Po, mm Hg) between perfusate and coronary effluent flow, and c is the Bunsen solubility coefficient of O2 in perfusate at 37°C (22.7 μ l O₂ × atm⁻¹ × ml⁻¹perfusate) (27). Oxygen extraction was calculated as the expression $O_2EXT = MV_{O_2}$ /oxygen content in the perfusate. Procedures followed were in accordance with institutional and NIH guidelines on treatment of animals.

RNA isolation. After removal of excess fat and connective tissues, the left ventricular wall was briefly blotted on nine-layer gauze, then frozen in liquid nitrogen, and stored at -80° C. An aliquot (100 mg) of the frozen tissue was pulverized and homogenized and total RNA was extracted with a RNA Isolation kit (Ambion Inc., Austin, TX, U.S.A.). RNA samples were tested by ultraviolet absorption at A260 nM to determine the concentration. The quality and concentration of the RNA samples were further confirmed by electrophoresis on denatured 1% agarose gels (28).

Northern blot analysis. For Northern blot analysis, 15 μ g of RNA was denatured and electrophoresed in a 1% formaldehyde agarose gel, then transferred to a nitrocellulose transfer membrane (Micron Separations Inc., Westboro, MA, U.S.A.), and cross-linked to the membrane with short-wave ultraviolet cross linker. The prehybridizing and hybridizing solutions contained 50% formamide, 1× Denhardt's solution, $6 \times$ SSPE, and 1% sodium dodecyl sulfate (SDS). Complementary DNA (cDNA) probes were labeled with [³²P]dCTP by random primer extension (PRIME-IT II, Stratagene, La Jolla, CA, U.S.A.) and added to the hybridizing solution to a specific activity. Hybridization was carried out at 42°C for 18 h. The blots were then washed several times with a final wash in $2 \times$ standard sodium citrate and 0.1% SDS at 65°C. The relative amount of mRNAs was evaluated using a PhosphorImager (Model 400S, Molecular Dynamics, Sunnyvale, CA, U.S.A.). The same size area at each band was taken to measure the intensity and the same size area at the closest upstream position of each band was taken as the background of the image, respectively. The blots were also exposed on Kodak X-Omat film (Eastman Kodak Co., Rochester, NY, U.S.A.) at -70°C. RNA loading was normalized by comparison to that of 28s ribosomal RNA (28). Adenine nucleotide translocase isoform 1 (ANT₁) mRNA levels were detected using a 1.4-kb ANT₁ cDNA fragment cloned from a human skeletal muscle library (ATCC, Rockville, Maryland) (23, 35). βF₁-ATPase mRNA levels were detected using a 1.8-kb βF₁-ATPase cDNA fragment cloned from a human HeLa cell line library (ATCC) (4, 37). An inducible heat shock protein (HSP70-1) mRNA level was detected by a 1.7-kb cDNA fragment cloned from human hippocampus (ATCC) (17, 21). To compare different mRNA levels in the same myocardial sample, aliquots of 15-µg total RNA from the myocardium were analyzed by means of sequentially reprobing the membranes with 28s, ANT_1 , βF_1 -ATPase, and HSP70-1 cDNA probes.

Experimental Protocols

After instrumentation and performing calibrations were completed, left ventricular balloon volumes were varied over a range of values to construct left ventricular function curves. In this manner, it is possible to define a specific balloon volume that is associated

with a developed pressure between 100 and 140 mm Hg. This volume was maintained at the same level during baseline and reperfusion conditions. The intraventricular balloon volumes were not adjusted to produce specific end-diastolic pressures (rather, we defined a level of systolic pressure development), but end-diastolic pressures at baseline greater than 10 mm Hg were not considered acceptable (26). Data from hearts characterized by developed pressures less than 100 mm Hg or greater than 140 mm Hg were not used. Baseline data were obtained after an equilibration period of approximately 30 min. The same procedures were followed in each experiment. During the baseline period, data were obtained with the hearts maintained at 37°C by water circulated through the organ bath. The pulmonary outflow temperature was monitored continuously with a thermal probe to adjust the temperature of the infusion. During ischemia, the PSS infusion was stopped and 60 mL of oxygenated St. Thomas' cardioplegic solution (CP) at 4°C was injected into the aorta at a rate of 1 mL/s to begin the 2-h ischemia. The organ bath temperature was changed to 4, 30, or 34°C during ischemia. Fifteen milliliters of St. Thomas cardioplegic solution (4°C) was injected every 30 min thereafter. The St. Thomas cardioplegic solution contained 109.0 mmol/L NaCl, 25.0 mmol/L KCl, 21.9 mmol/L NaHCO₃, 16.0 mmol/L MgCl₂, and 0.8 mmol/L CaCl₂. Repeated administration of cardioplegic solution is used commonly in the clinical arena and is consistent with experimental studies that have reported previously (9, 19, 25, 28). When the 2-h ischemic period was ended, the hearts were reperfused with oxygenated PSS at 37°C and the water bath temperature was increased to 37°C. During the 45 min of reperfusion, hemodynamic data were recorded to compare with baseline data and to determine the degree of functional recovery in each heart. After reperfusion, hearts were quickly placed in liquid nitrogen for RNA measurements. Figure 1 briefly illustrates the protocol.

In previous studies, the functional recovery

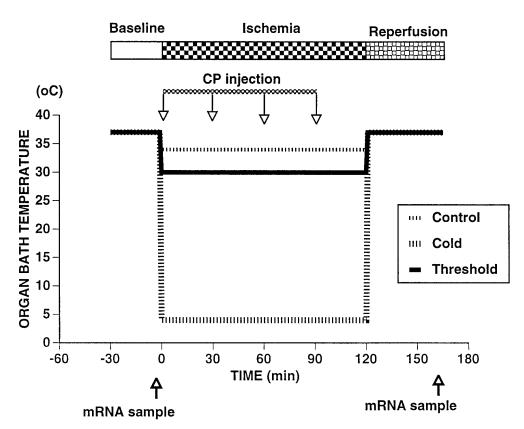


FIG. 1. Experimental protocol. The abbreviations used are: CP, cardioplegic solution; Control, control hearts treated at 34°C ischemia; Cold, hearts treated at 4°C ischemia; Threshold, hearts treated at 30°C threshold temperature. Function signals were continuously recorded during the experiments.

was similar at 4, 18, and 30°C ischemia, but the recovery was dramatically decreased at 34°C ischemia (27). To simplify the protocol, in this study 20 hearts were divided into 4°C (n = 4), 30°C (n = 4), and 34°C (n = 8) ischemic groups as well as a baseline control group (n = 4). In order to evaluate the alteration of mRNA levels of genes, values of ischemic groups were compared to the baseline control value.

Statistical Analysis

Values reported are means \pm standard error (SE) in the text, tables, and figures. The Statview 4.5 (FPV) program (Abacus Concepts, Inc., Berkeley, CA, U.S.A.) was used for statistical analysis. Data were evaluated with repeated measures analysis of variance within groups and single-factor analysis of variance across groups. When significant *F* values were obtained, Scheffe's test was used to distinguish which groups differed from each other significantly. The criterion for significance was taken to be P < 0.05 for all comparisons.

RESULTS

Functional Parameters

Baseline. The heart weights were similar in the groups (6.6 \pm 0.5, 6.9 \pm 0.4, and 7.5 \pm 0.8 g in the 4, 30, and 34°C groups, respectively; P > 0.05). The end diastolic left ventricular pressure was adjusted to similar levels (2.5 \pm 0.5, 3.4 \pm 0.3, and 1.7 \pm 0.3 mm Hg in the 4, 30, and 34°C groups with the ventricular volume at 1.2 \pm 0.1, 1.3 \pm 0.1, and 1.7 \pm 0.1 ml, respectively) at baseline conditions. There

Hemodynamics (Means \pm SE)			
Group	$4^{\circ}\mathrm{C}\ (n=4)$	$30^{\circ}{ m C}$ (<i>n</i> = 4)	$34^{\circ}\mathrm{C}\ (n=8)$
DP (mm Hg)			
Baseline	116.2 ± 5.0	114.2 ± 4.9	119.0 ± 4.5
REP	69.5 ± 4.1	66.1 ± 3.0	$38.0 \pm 6.4*$
dP/dt_{max} (mm Hg/s)			
Baseline	2037 ± 168	2024 ± 235	1875 ± 211
REP	1095 ± 59	1049 ± 103	$529 \pm 101*$
$-dP/dt_{max}$ (mm Hg/s)			
Baseline	1420 ± 65	1423 ± 111	1260 ± 159
REP	977 ± 57	887 ± 37	$449 \pm 85^{*}$
HR (Beats/min)			
Baseline	200.8 ± 20.4	212.0 ± 17.5	187.9 ± 11.8
REP	226.8 ± 16.1	214.8 ± 17.3	173.0 ± 18.3
PRP (10 ³ mm Hg/min)			
Baseline	23.29 ± 2.37	24.14 ± 1.84	20.81 ± 2.71
REP	15.75 ± 1.42	14.19 ± 1.20	$6.04 \pm 0.95^{\circ}$
MV_{O_2} (µl/g/min)			
Baseline	122.3 ± 13.9	111.0 ± 5.4	94.0 ± 5.1
REP	87.3 ± 12.3	80.2 ± 3.0	$34.0 \pm 5.3^{*}$

TABLE 1 Hemodynamics (Means \pm SE)

Note. The hemodynamic indices were determined in isolated reperfused hearts at baseline and after 45 min of reperfusion (REP) as described under Materials and Methods at 4, 30, and 34°C ischemic groups, respectively. The abbreviations used are: DP, left ventricular developed pressure; dP/dt_{max} , positive maximum of the first derivative of left ventricular pressure; $-dP/dt_{max}$, negative maximum of the first derivative of left ventricular pressure; HR, heart rate; PRP, product of HR and DP; MV_{Q2}, myocardial oxygen consumption. **P* < 0.05 vs 4°C group.

were no significant differences between groups in DP, $\pm dP/dt_{max}$, HR, PRP, or MV_{O2} (Table 1).

Functional recovery during reperfusion. In Table 1 and Fig. 2, the data demonstrate that hypothermic hearts provide greater functional

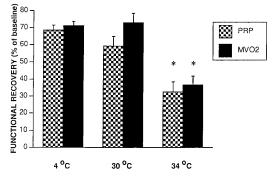


FIG. 2. Effect of hypothermia on functional recovery during reperfusion. The reperfused cardiac function parameters are plotted as a percentage of baseline (37°C). The abbreviations used are: PRP, product of HR and DP; MV_{O_2} , myocardial oxygen consumption. *P < 0.05 vs 4°C.

recovery than those observed in the 34°C hearts. There were no significant differences between hearts after 4 and 30°C ischemia. The hypothermic hearts were characterized by higher developed pressures, higher $\pm dP/dt_{max}$ values, higher products of heart rate and DP, and higher myocardial oxygen consumption, indicating that heart function was improved in the hypothermic groups.

Ischemic contracture. As noted under Materials and Methods, a specific balloon volume was adjusted and maintained throughout the experiment, allowing comparisons of left ventricular pressure under constant end-diastolic volume. After injecting cardioplegic solution, the left ventricular pressure was always near 0 mm Hg. The beginning of ischemic contracture was defined by the initial rise in left ventricular pressure above 2 mm Hg. Ischemic contracture started in the 34°C control group after 64.9 \pm 4.8 min of ischemia but there was no contracture in the hypothermic groups.

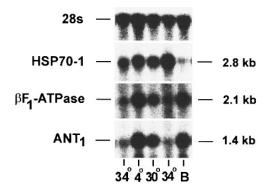


FIG. 3. A representative Northern blot. Each lane was loaded with 15 μ g total RNA from ventricular myocardium and probed specifically for 28s, β F₁-ATPase, ANT₁, and HSP70-1. Samples were taken from hearts at baseline control and after 2 h ischemia at 4, 30, or 34°C followed by 45 min reperfusion (see text). The abbreviations used are: 4°, 30°, and 34°, ischemic temperature at 4, 30, and 34°C, respectively; B, baseline.

ANT₁, βF_1 -ATPase, and HSP70-1 mRNAs

Representative Northern blot analyses for mRNAs of ANT₁, β F₁-ATPase, and HSP70-1 are shown in Fig. 3. Compared to hearts perfused at baseline, both 4 and 30°C treatment resulted in no significant differences in mRNA levels. Steady-state mRNA levels for ANT₁ β F₁-ATPase and HSP70-1, normalized to 28s ribosomal RNA intensity, are shown in Figs. 4 and 5 for tissue obtained during the protocol. On the other hand, postischemic steady-state levels of ANT₁ and β F₁-ATPase mRNA was

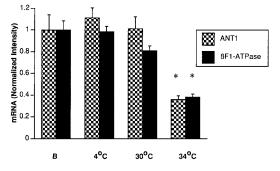


FIG. 4. Steady-state mRNA levels of ANT₁ and β F₁-ATPase. All transcript levels are relative to the 28s band intensity and normalized to the B lane. Abbreviations are defined in the legend to Fig. 3. **P* < 0.05 vs B.

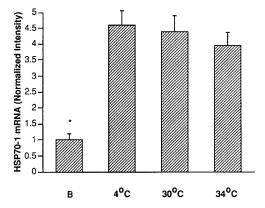


FIG. 5. Steady-state mRNA levels of HSP70-1. All mRNA levels are relative to the 28s band intensity and normalized to the B lane. Abbreviations are defined in the legend to Fig. 3. *P < 0.05 vs 34°C.

markedly diminished in 34°C hearts (Figs. 3 and 4). The HSP70-1 mRNA level was fourfold higher in 34°C ischemic hearts than baseline condition and there were no noticeable differences from that in 4 and 30°C ischemia (Fig. 5).

DISCUSSION

Previous work using the current model demonstrated that preservation of postischemic function was closely related to ATP levels during reperfusion (25, 27, 28). These levels cannot be considered indices of mitochondrial function or damage, as they are dependent on the balance between energy production and utilization during ischemia, as well as the status of the purine salvage pathways (30). Reductions in ATP utilization caused by temperatures below the threshold preserve ATP stores during ischemia and prevent nucleotide loss through purine degradation pathways (27). Preservation of ATP or rapid restoration during reperfusion ensures continuation of processes, which are ATP dependent. These processes clearly include those utilizing ATPases for energy conversion, such as the myosin-actin interaction.

However, the role of maintained ATP stores in preservation of pathways necessary for the cardiomyocyte to initiate repair after ischemia has not been fully established. These pathways might include those responsible for mitochondrial repair as mitochondrial dysfunction and damage occur during hypoxia/ischemia and reoxygenation or reperfusion (30). This dysfunction has been linked to reductions in activities of both the adenine nucleotide translocator and the mitochondrial ATPase (8, 14). The adenine nucleotide translocator and the BF1-ATPase together account for greater than 30% of mitochondrial protein (12). Logically, one might assume that restoration of normal mitochondrial function following ischemic injury would require resynthesis and/or reconfiguration of these major mitochondrial membrane proteins (1, 4, 10, 11, 23, 32, 37). As expression of the adenine nucleotide translocator protein in particular closely coordinates with steady-state mRNA levels for the corresponding gene (31), one would expect that preserved gene expression would be necessary for eventual protein resynthesis. Maintenance of ATP levels have been closely linked to preservation of these mRNA levels, which are considered indices of signaling for mitochondrial biogenesis (28). Similarly, the current studies demonstrate preservation of steady-state mRNA levels after ischemia at the critical temperature or below, where we have previously demonstrated reductions in ATP depletion (27).

Increased expression of cytosolic heat shock proteins such as HSP70-1 has been related to preserved myocardial function after reperfusion (5, 18, 22, 24). The mechanisms responsible for this relationship are not fully elucidated. However, cytosolic stress proteins do interact with mitochondrial protein precursors in the cytoplasm (16, 34), prevent their aggregation, and promote their import across the mitochondrial membrane. Stress agents such as heat shock or hypothermia applied prior to ischemia enhance HSP-70 expression in isolated hearts or cells after ischemia (5, 13, 20, 22, 29). Application of these stress agents yields a substantial elevation in steady-state HSP70-1 mRNA during reperfusion in isolated rabbit hearts (28). Previously, a near 12-fold increase in these steady-state levels occurred after hypothermic adaptation and subsequent ischemia (28). In comparison, the current protocol produced a modest 4-fold increase in HSP70-1 mRNA after reperfusion. This elevation occurred at all levels of ischemic temperature studied during these experiments. This may be considered a response to ischemia, but it does not appear to be ischemic temperature dependent and does not relate to the level of postischemic myocardial function or ATP preservation.

In summary, these data show that maintenance of postischemic steady-state mRNA levels for ANT₁ and the β F₁-ATPase occurs at or below the previously defined critical temperature (30°C) for superior postischemic ATP and cardiac function preservation. Preservation of these mRNA levels implies, though does not prove, that the corresponding protein levels will be amplified. However, these data provide the basis of a novel hypothesis, whereby hypothermia promotes myocardial protection through maintenance of transcriptional signaling pathways.

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