MCT1 and MCT4 Expression During Myocardial Ischemic-Reperfusion Injury in the Isolated Rat Heart

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Key Words
Lactate • Myocardial ischemia-reperfusion injury • Monocarboxylate transporters

Abstract

Background/Aims: Myocardium ischemia-reperfusion (I/R) injury can be caused by imbalances in cellular metabolism. Lactate, transported by monocarboxylate transporters (MCTs), has been implicated as a mechanism in this process. The present study was designed to investigate the expression and functional role of MCTs in rat hearts during ischemia and reperfusion. Methods: Langendorff-perfused rat hearts were subjected to 20 minutes stabilization, 30 minutes of global ischemia and 60 minutes reperfusion. Hearts were collected serially for detecting expression changes in MCT1, MCT4 during myocardial I/R injury and lactate concentration was measured. Post-ischemic left ventricular function and infract size were determined at end-point, followed by the pretreatment of D-lactate, a competitive inhibitor of MCTs. Results: MCT4 was significantly increased following global ischemia and MCT1 expression was increased during the early stages of reperfusion in isolated rat hearts, while the expression of the ancillary protein CD147 was increased during I/R injury. We determined increases in AMPK phosphorylation status, which was significantly elevated following ischemia and early reperfusion. Blocking monocarboxylate transport by competitive inhibition with D-lactate caused decreased left ventricular performance and increased infarct size. Conclusion: Increased MCT4 expression facilitates lactate extrusion during the ischemic period, while increased MCT1 may facilitate lactate transport into and out of cells simultaneously during early reperfusion, with increases in AMPK phosphorylation status during the myocardial I/R period. Lactate transport by MCTs has a profound protective effect during myocardial ischemia reperfusion injury.
Introduction

Ischemic heart disease, which is caused by an interruption in blood flow to the myocardium, is predicted to become the leading cause of mortality globally by the year 2020 [1]. Emergent reperfusion therapy for the restoration of coronary patency is currently the main treatment. However, immediate reperfusion has been shown to accelerate cell death and increase the extent of infarction, which is undesirable but inevitable. This process is known as myocardial ischemic-reperfusion (I/R) injury [2]. Research to protect the heart against myocardial I/R injury has progressed for decades, both in clinical practice and basic research. Nowadays, many studies are focused on investigating the underlying intracellular mechanisms and signal transduction complexes involved in cardioprotection. Immediate recanalization remains the gold standard for current clinical treatments, although there are known consequences of reperfusion injury. There remains an increasing need for novel therapeutic strategies for cardioprotection against I/R injury.

Myocardial I/R injury is a metabolic event caused by a short supply of energy. Alterations in energy metabolism is a vital issue triggering myocardial I/R injury [3]. Following the onset of ischemia, oxidative phosphorylation of glucose is inhibited and glycolysis is the only mechanism of ATP production [3]. During the early stages of reperfusion, glycolysis continues while glucose oxidation is inhibited [4]. Lactate, which links glycolysis and oxidative metabolism [5], may participate in the imbalance of energy metabolism during I/R injury.

Lactate, the end product of glycolysis resulting from hypoxia, once was treated as a metabolic waste product. However, the potential therapeutic targets of lactate are now being recognized. Lactate plays a central role in cellular metabolism and metabolic communication between tissues [6]. Transport of lactate into and out of cells is facilitated by a family of monocarboxylate transporters (MCTs) [7]. Among the isoforms of MCT, MCT1 and MCT4 are considered to be the key transporters in cardiac muscle [8]. The transportation function of these isoforms is associated with CD147 (EMMPRIN) [9], a key glycoprotein, which assists MCT1 and MCT4 in folding, stability, membrane expression, and functionality. Moreover, lactate dehydrogenase (LDH) is involved in the shuttling of lactate between cells via the cell-cell lactate shuttle [10].

Recent reports have shown that AMP-activated protein kinase (AMPK) activation increases MCT1 and MCT4 protein expression in skeletal muscle [11-13]. Furthermore, the current view is that AMPK protects the heart from I-R injury by sustaining an energy supply [2, 14]. It therefore follows that there may be an analogous correlation between MCTs and AMPK in cardiac muscle.

The purpose of the present study was two-fold: 1. Identify MCT1 and MCT4 expression in isolated rat hearts during acute I/R injury; and 2. Investigate the functional role of MCTs in acute I/R injury.

Materials and Methods

Adult male Sprague-Dawley rats (250–300g) were purchased from the Hunan Weaselyg Scene of Experimental Animals Co, LTD (Hunan China). All animal protocols were performed in accordance with the U.S. National Institute of Health guidelines (NIH Publication No.85-23, revised 1996) and approved by the Animal Care and Use Committee of Tongji Medical College of Huazhong University of Science and Technology.
series. Series 1: hearts were perfused with a 20 minute stabilization period (T0, n=8), or perfused with
20 minutes stabilization, 30 minutes of global ischemia (I30, n=8), or reperfused for 15 minutes following
ischemia (R15, n=8), or reperfused for 60 minutes following ischemia (R60, n=8). These hearts were
collected to determine the expression and level of MCT1, MCT4, CD147, LDH and AMPK phosphorylation.
Coronary effluent at the end of the study in each group was collected for lactate release measurements.
Once series 1 was accomplished, series 2 was performed to study lactate transport during the I/R period
mediated by MCTs and to test whether lactate itself influences heart performance. Series 2: hearts were
perfused with 20 minutes stabilization, 30 minutes global ischemia, and 60 minutes reperfusion (I/R, n=9),
and compared with hearts perfused for 5 minutes with 100 mM of either D-lactate (DLAC, n=9) or L-lactate
(L-LAC, n=9) prior to ischemia. During perfusion hemodynamic parameters, including left ventricular (LV)
function, were recorded continuously. At the end of perfusion, besides the measurements determined in
series 1, hearts were collected for the determination of myocardial apoptosis and necrosis, and myocardial
infarct size.

Real-time RT-PCR

MCT1 and MCT4 mRNA levels in rat cardiomyocytes were determined using quantitative real-time
PCR analyses. Myocardium samples were homogenized with Trizol reagent (Invitrogen, USA) and total
RNA was extracted according to the manufacturer’s protocol. Concentration and purity of total RNA was
determined using a BioPhotometer (Eppendorf, Germany). Total RNA was then reverse-transcribed into
complementary DNA (cDNA) using a cDNA synthesis kit (TOYOBO, Japan) which was performed according
to the manufacturer’s protocol. Real-time PCR was performed using a StepOne Real-Time PCR System
(Applied Biosystems, USA) and Thunderbird SYBR qPCR Mix (TOYOBO, Japan). The RT-PCR conditions were
42°C/30 min, and 80°C/5 min for reverse transcription. PCR involved pre-denaturing at 95°C for 1 minute,
then 40 cycles of 95°C for 15 seconds, 58°C for 20 seconds, and 72°C for 20 seconds. β-actin was used as
the housekeeping gene. Levels of MCT1 and MCT4 mRNA were calculated based on the 2^-ΔΔCt
comparative quantification method. The PCR primers used included: MCT1 forward primer (5′-CTC TGG
GCG CCG GAT AC-3′), reverse primer (5′- CAA CTA CCA CCG CCC AGC CC-3′); MCT4 forward primer
(5′-CCA GCC CGA CCG GCC CCA CGG CAG GTT TC-3′), reverse primer (5′-GCC ACC GTA GTC ACT
GGC CG-3′); β-actin forward primer (5′-ACG TTG ACA TCC GTA AAG AC-3′) and reverse primer
(5′-GAA GGT GGA CAG TGA GGC-3′). All samples for each gene
were run in duplicate.

Western blotting

LV tissue was homogenized and protein concentration of the supernatant was determined using a
BCA method (Pierce, USA). The supernatant containing 50 μg of protein was separated using a 10%SDS-
PAGE and transferred onto polyvinylidene difluoride membranes (Millipore, USA) by electrophoresis. The
membranes were then blocked in TBS containing 0.1% Tween-20 and 5% nonfat milk at room temperature
for 1 hour. The membranes were incubated in primary antibodies (anti-Actin, anti-MCT1, anti-MCT4 and
anti-CD147, 1:250, Santa Cruz, USA; anti-AMPK-α, anti-phospho-AMPK-α, 1:1000, Cell Signalling Technology,
USA) at 4 °C overnight, followed by a HRP-conjugated anti-rabbit IgG secondary antibody or anti-mouse IgG
secondary antibody (1:5000, Proteintech, China). Target proteins were detected using the ChemiDocXRS+
chemiluminescence imaging system (Bio-Rad, USA). Protein bands were quantified by Image Lab image
acquisition and analysis software (Bio-Rad, USA).

Measurement of myocardial LDH content and activity

Cardiac tissues were homogenized in PBS (pH 7.4) and centrifuged for extracting supernatant.
Myocardial LDH levels were measured using a rat LDH enzyme-linked immune sorbent assay (ELISA) kit
(IBL, Germany). Myocardial LDH activities were measured using LDH kits (Nanjing Jiancheng Biotech,
China).

Lactate concentration in coronary effluent

Lactate concentration in coronary effluent was measured using a lactic acid kit (Nanjing Jiancheng
Biotech, China) according to the manufacturer’s instructions.
**Cardiac function measurements**

Cardiac LV function was recorded continuously via a water-filled balloon placed into the left ventricle and attached to a pressure transducer after the heart was mounted on a Langendorff apparatus (ADInstruments, Australia). Cardiac functional assessment was then collected and analyzed using a PowerLab data-acquisition system with LabChart software (version 7 for Windows, ADInstruments, Australia). Measured parameters included LV developed pressure (LVDP), LV end-diastolic pressure (LVEDP), and maximum LVDP increase (+dp/dt) and decrease (−dp/dt) rates. Coronary flow rate (CF) was calculated from the coronary effluent.

**Myocardial infarct size assessment**

At the end of the experiment, the hearts were frozen at -20°C for 2 hours and then sliced transversely into 2-mm-thick sections and incubated in a 1% solution of 2,3,5-triphenyltetrazoliumchloride (TTC; Amresco, USA) for 15 minutes at 37°C, followed by transfer into a 4% paraformaldehyde–phosphate-buffered saline solution for fixation at 4°C. The infarcted tissue remained unstained (white), whereas normal tissue was stained red. Infarct area was traced and the total area was calculated using Image J software (NIH, USA). Infarct size was expressed as a percentage of total ventricular area.

**Determination of myocardial apoptosis and necrosis**

Terminal dUTP nick-labelling (TUNEL) was assessed using a kit (Roche, Switzerland) according to the manufacturer’s instructions in order to determine cardiomyocyte apoptosis. TUNEL-positive nuclei with chromatin condensation and fragmented nuclei were considered as probable apoptotic cells. Quantification of apoptosis was determined by counting the TUNEL-positive myocyte nuclei from three random fields per section using light microscopy at x200 magnification by an investigator who was blinded to the study. LV tissues were fixed in 4% paraformaldehyde and embedded in paraffin. Sections were stained using haematoxylin and eosin (HE), and then histologically assessed by a pathologist for acute myocardial necrosis, interstitial oedema, and cellular swelling.

**Statistics**

All values are shown as mean ± standard error of the mean (SEM). GraphPad Prism (version 5 for Windows, GraphPad software, USA) was used for all statistical analyses. Differences in the recovery of post-ischemic hemodynamic parameters were tested using a two-way analysis of variance (ANOVA) followed by a Bonferroni’s post-hoc test for multiple comparisons. Other data were subjected to one-way ANOVA followed by a Tukey’s post-hoc test. Statistical significance was set at P<0.05.

**Results**

**MCT1 and MCT4 mRNA expression in cardiomyocytes**

Series 1: Compared with the level prior to ischemia, MCT1 mRNA was significantly elevated following 15 minutes reperfusion (P<0.01; Fig. 1A), while MCT4 mRNA was rapidly increased after 30 minutes ischemia (P<0.001; Fig. 1B). There was no obvious difference when the rest group was compared to T0. Series 2: compared with the I/R group, MCT1 and MCT4 mRNA was increased in the D-LAC group (P<0.01, respectively, Fig. 1C, D), while the L-LAC and I/R groups were similar.

**Western blotting**

Series 1: Western blotting showed that MCT4 protein expression was significantly increased following 30 minutes ischemia (P<0.01; Fig. 2B). After 15 minutes reperfusion, protein expression decreased to the level occurring before ischemia until 60 minutes reperfusion. MCT1 expression was markedly increased after 15 minutes reperfusion (P<0.01; Fig. 2A) and the level was normalized after 60 minutes reperfusion. CD147 protein expression was increased during the I/R period compared to the level before ischemia (P<0.05; Fig. 2C). AMPK phosphorylation status [phospho-AMPK (Thr172)/total AMPK] was significantly elevated following 30 minutes ischemia and 15 minutes reperfusion (P<0.01, P<0.05, respectively, Fig. 2D). After 60 minutes reperfusion, the level was normalized. Series
There were no statistically significant differences between groups with regard to MCT1, MCT4 and CD147 protein expression (Fig. 2E, F, G). However, AMPK phosphorylation status was significantly elevated in the D-LAC group compared with both L-LAC and I/R groups (P<0.01, respectively, Fig. 2G), while no statistical differences were observed between the L-LAC and I/R groups.

**LDH content and activity in coronary effluent**

Series 1: LDH content was elevated after 30 minutes ischemia and 15 minutes reperfusion (P<0.01, P<0.05, respectively, Fig. 3A). After 60 minutes reperfusion, the level dropped to that observed before ischemia. The activities of LDH were in parallel to the content during the I/R period (P<0.05, P<0.05, respectively, Fig. 3B). Series 2: There were no statistically significant differences between groups (Fig. 3C, D).

**Lactate concentration**

Series 1: Lactate concentration was elevated after 30 minutes ischemia (P<0.01; Fig. 4). The concentration rapidly fell to the level before ischemia after 15 minutes reperfusion. Series 2: There were no statistically significant differences between groups (Fig. 4C, D).

**Cardiac LV function**

Following sustained ischemia, hearts pretreated with D-lactate had a worse performance in the recovery of post-ischemic hemodynamic parameters. Compared with hearts pretreated with L-lactate or not pretreated, LVDP, ±dp/dt were significantly depressed in D-lactate (P<0.01, P<0.01, respectively, Fig. 5A, B, C). LVEDP ascended during reperfusion in D-lactate compared with L-lactate or I/R, respectively (P<0.01, P<0.01, respectively, Fig. 5D). CF of hearts pretreated with D-lactate was also reduced compared with L-lactate or I/R (P<0.01,
Fig. 2. (A-D) Western blot analyses of MCT1 and MCT4, CD147 expression and AMPK phosphorylation status during myocardial I/R period. Actin was used as an internal reference. Values are expressed as means±SEM. *P < 0.05 vs T0 group. (E-H) Western blot analyses of MCT1 and MCT4, CD147 expression and AMPK phosphorylation status in hearts perfused with L- and D-lactate for 5 minutes or without lactate before 30 minutes of global ischemia and 60 minutes reperfusion. Values are expressed as means±SEM. *P < 0.05 vs I/R group, # P < 0.05 vs L-LAC group.
P<0.05, respectively, Fig. 5E). There were no differences between groups L-lactate and I/R during reperfusion in these parameters except ±dp/dt at the early stage of reperfusion (P<0.05, Fig. 5B, C).

**Infarct size**

Infarct size in each group was evaluated after 60 minutes reperfusion. D-lactate significantly increased the infarct size percentage compared with L-lactate and I/R (P<0.05, respectively, Fig. 6). There were no differences between L-lactate and I/R groups.

**Myocardial apoptosis and necrosis**

Using the TUNEL assay, a large number of TUNEL-positive cells were observed in hearts pretreated with D-lactate, while few TUNEL-positive cells were visible in the L-lactate and
I/R groups. The percentage of apoptotic cells in D-lactate were significantly increased compared with the L-lactate or I/R groups (P<0.01, P<0.01, respectively, Fig. 7). There was no difference between the L-lactate and I/R groups. D-lactate increased the extent of cellular swelling and extracellular matrix oedema (Fig. 8).
Under physiological conditions, myocardial ATP production is dependent chiefly on oxidative decarboxylation of fatty acids, providing 60–80% of the energy demand. The remaining ATP requirement is supplied by glucose and lactate in nearly equal proportions [15], and other substrates such as amino acids and ketone bodies. Co-utilization is determined
by both the amount of substrate energy generated in the circulation and complex intracellular regulatory mechanisms [16]. However, this regulated balance is altered significantly during the myocardial I/R period. ATP generated from glycolysis prioritizes the function of the ion pump in the cell membrane, while ATP from glucose oxidation promotes the recovery of normal contractile function in the post-ischemic heart [17, 18]. Therefore, reductions in ATP generation during the I/R period results in the accumulation of intracellular ions and myocardial dysfunction. Ischemic-reperfusion injury then occurs. As mentioned above, lactate is not just the production of glycolysis, but the energy substrate for glucose oxidation via the Krebs cycle. Since the heart is an efficient lactate consumer and producer [19], we speculate that lactate transport is associated with energy imbalance during the myocardial I/R period.

Alterations in the expression of the lactate shuttle machinery may affect cellular transport and oxidation of lactate [20]. In earlier studies of lactate transport in the heart in relation to myocardial ischemia [21], Halestrap et al. assumed that there are two MCT isoforms in the heart, one being MCT1 which is found to be expressed in almost all tissues [7], while the other is MCT4, or at least this has been argued [8, 22]. Halestrap believed MCT4 is widely expressed and especially restricted in glycolytic tissues while MCT4 is absent in the heart. However, hypoxia-induced MCT4 expression has been reported by McClelland and Brooks [20]. In addition, another study demonstrated that MCT4 is rapidly induced by hypoxia through the HIF-1α pathway [23]. MCT4 is adapted to export lactate, although it has low affinity for L-lactate because of its very low affinity for pyruvate (Km, 150 mM) while other isoforms are extremely high, which would be of particular importance since it prevents the loss of pyruvate from the cell. This is essential because high rates of glycolysis requires that NADH produced in glycolysis is re-oxidised, which is achieved by pyruvate being converted to lactate [24]. In a global Langendorff rat heart model, glycolysis is the only means for energy generation. During global ischemia, it slows gradually at 30 minutes of ischemia, and may continue even longer until lower pH mediates inhibition of glycolysis before complete glycogen depletion [25, 26]. In humans, myocardial ischemia is a common occurrence which extends from hours to days preceding death. Therefore, the hypothesis that MCT4 is totally absent from the heart is unsound. In our study, MCT4 was markedly expressed following 30 minutes ischemia with lactate concentrations in the coronary effluent very high. Moreover, expression of LDH and CD147 is also increased following ischemia. This result suggests that it is MCT4 which mainly facilitates lactate extrusion during the ischemia period.

After 15 minutes reperfusion, we found that MCT1 is significantly increased in the isolated heart. This result is consistent with Martinov’s study in mice [27]. The predominant role of MCT1 is to facilitate unidirectional proton-linked transport of L-lactate across the plasma membrane. Intracellular and extracellular pH achieved 5.9 and 5.5, respectively after 20 minutes ischemia [26]. MCT1 may facilitate lactate extrusion during the ischemia period and the onset of reperfusion to maintain non-acidity in the intracellular environment. However, intracellular pH in cardiomyocytes returns to normal within a few minutes of reperfusion [26], and our results showed that after 15 minutes reperfusion, lactate concentration in the effluent nearly drops to the level observed before ischemia. In addition, LDH and CD147 are also active at this time, which indicates that lactate metabolism is still active. Therefore, it seems that the function of the abundant expression of MCT1 at this moment is in question. During the early stages of reperfusion, glycolysis may exceed regional imbalance between glycolysis and glucose oxidation [4, 28, 29]. As lactate is a preferred fuel for the injured heart [30], and it is proven to provide simultaneous lactate uptake and extrusion as functionally separate metabolic pathways within the heart [31], we speculate that MCT1 affords lactate transport into and out of the cells simultaneously at this moment. This requires further study.

Considering the lack of common inhibitors of both MCT1 and MCT4 [22], D-lactate is a valuable tool with which to investigate L-lactate transport [32]. When MCTs were competitively blocked using D-lactate, left ventricular function was depressed and infarct size increased. This indicates that lactate transport by MCTs has a profound protective effect on myocardial ischemia reperfusion injury. Although myocardial MCT1 and MCT4 expression
were not influenced by either L- or D-lactate in this study, we found that their mRNA expression and AMPK phosphorylation status increased following D-lactate pretreatment. Considering lactate may be a major energy resource for the failing heart, and may preserve or even improve myocardial performance [30, 33], this possibly indicates that the hearts are failing and have a greater reliance on carbohydrates. It is suggested that regulation of MCT expression seems to be translational [13]. We did not find increased MCT expression, which may be due to the time frame of observation.

AMPK has been proposed as a putative target to limit the extent of I/R injury [14]. It protects the heart from I/R injury by sustaining energy supply during ischemic stress and reducing the likelihood of cardiomyocyte cell death. Furthermore, in other studies, AMPK activation induced increases in MCT1 and MCT4 protein content in rat skeletal muscle [11-13]. In the current study, we also found that MCT1 and MCT4 protein expression was increased with increases in AMPK phosphorylation status during the I/R period. As a master metabolic regulator in mammalian cells, AMPK activity may be important and required for up-regulating MCT1 and MCT4 expression, to facilitate lactate uptake and release in the heart for meeting cellular homeostasis and energy demands during the myocardial I/R period. The specific mechanism needs further study.

There are several limitations in our study. First, our model did not take fatty acid into account in energy imbalance during myocardial I/R injury. We know that oxidation of fatty acid has a competitive relationship with glucose in myocardial energy metabolism [34]. Fatty acid may cause a more complicated, moot result. Second, D-lactate and L-lactate may influence cellular pH in cardiomyocytes, but no common inhibitors are available.

Our study first showed that MCT1 and MCT4 protein expression are increased during the myocardial I/R period with increases in AMPK phosphorylation. Pharmacological targeting of MCTs may provide new treatment strategies for myocardial ischemia–reperfusion injury. In addition, AMPK may be a potential modulation pathway for MCTs.

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Conflict of Interest

None.

References


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