



Allograft Inflammatory Factor-1 Induction during Ischemia Reperfusion Injury: A Rat Model of LAD Occlusion

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Abstract

The emerging role of the innate immune response in organ transplantation heightens the significance of molecular markers released from the donor organ due to ischemia/reperfusion injury (IRI) during organ procurement. These molecules are prime activators of innate immunity. Allograft inflammatory factor-1 (AIF-1) has been implicated in the regulation of inflammation and organ rejection. To verify expression level of AIF-1 in cardiac tissues, we developed a rat model of IRI, in which the left anterior descending artery (LAD) was occluded to generate ischemia in the left ventricle (LV) of the heart. Blood and cardiac tissues were tested for the presence of AIF-1 at different time intervals. AIF-1 expression was significantly increased in a time-dependent manner after IRI. AIF-1 was up regulated as early as 10 minutes after reperfusion, and further it was increased several-fold after 60 minutes of reperfusion in PBMCs and PMNCs as compared to the control group. Expression levels of AIF-1 in LV tissues were significantly increased after 60 minutes of ischemia or 30 minutes of ischemia followed by 60 minutes of reperfusion. In addition the level of AIF-1 was 2-fold greater in LV tissues after 60 minutes of reperfusion compared to LV after 30 or 60 minutes of ischemia. Thus, myocardial IRI activates AIF-1 up-regulation, which may result in allograft dysfunction in transplantation settings.

Keywords

Allograft inflammatory factor-1, Cardiac, Ischemia, LAD, Myocardium, Rat, Reperfusion

Introduction

Ischemia reperfusion during organ procurement and recovery is an unavoidable event that contributes significantly to tissue injury and may affect early organ dysfunction after transplantation. However, in clinical settings, the minimum ischemic time that results in a significant impact on cardiac transplantation is not clear. A preservation time >24 hour in kidney as well as lung transplantation has been linked to a decline of organ function during early post-transplantation [1-3]. It has been well documented that ischemia, (the disruption of the blood supply to the organ during procurement)

followed by restoration of blood flow during reperfusion often creates an ultra-structural and metabolic/cellular injury (IRI). This leads to profound changes in the production of reactive oxygen species (ROS) causing myocardial apoptosis and cardiomyocyte death in ischemic regions [4-6]. The consequences of pathological obstruction or the experimental ligation of the coronary artery in the induction of ischemia and myocardial injury is well known. Inflammation is the hallmark of these consequences thus; targeting molecules initiating such inflammatory responses is an emerging therapeutic intervention with significance in the prevention of ischemia induced injury.

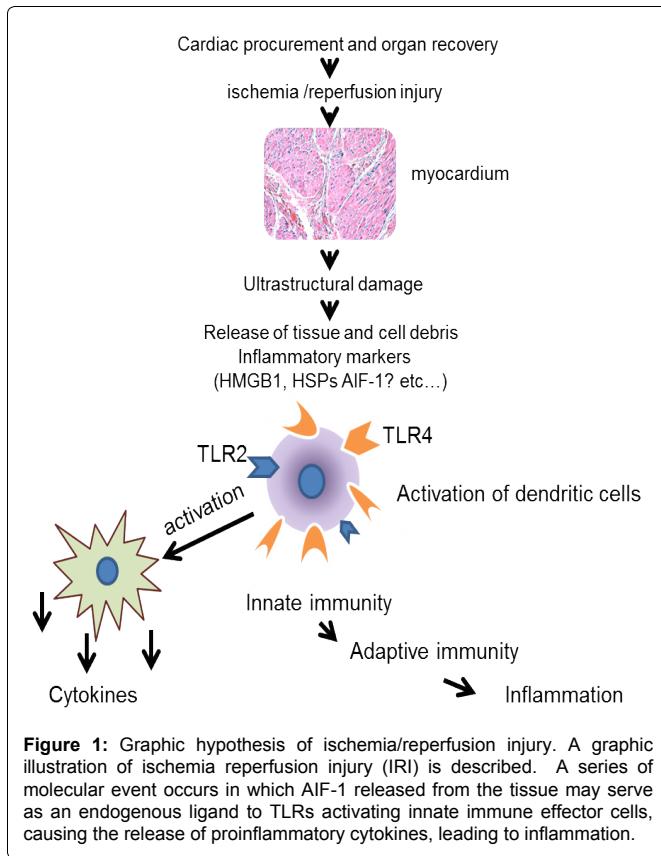
IRI induced inflammation typically occurs in a sterile environment but, shares many characteristics of the host immune response aimed at pathogenic microorganisms [7]. The immune response associated with cardiac IRI is initiated by activation of the myocardial resident dendritic cells and components of the innate immune system. This includes pattern recognition molecules such as Toll-like receptors (TLRs) [8-10], which results in the induction of proinflammatory cytokines and chemokines [7]. Cytokines are secreted by all resident cells in the myocardium and by the leukocytes that migrate into the heart [11]. Eventually, the innate response and the parameters involved in the activation of this early response bridges into adaptive immunity leading to allograft dysfunction and rejection [12]. Given the importance of innate immune responses in the recognition of pathogen or damaged tissue associated molecular patterns known as PAMPs or DAMPs respectively, we postulated that the molecules released from a donor organ during procurement may play a crucial role in the initial inflammatory response to the allograft.

A broad range of non-pathogenic motifs could trigger the induction of an inflammatory episode in the myocardium. It is possible that one of the contributing factors in the initial inflammatory response during IRI is the release of allograft inflammatory factor-1 (AIF-1). AIF-1 is a potential endogenous ligand that engages in activation of TLR-bearing dendritic cells initiating the innate immune response in transplantation settings. An adapted hypothetical model of our study is summarized in Figure 1 [13]. AIF-1 has been implicated in the regulation of inflammation and thus far many

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investigators have shown an association between elevated levels of AIF-1 expression with cellular rejection and coronary vasculopathy after organ transplantation [14-16].

AIF-1 is a 143 amino acid, cytoplasmic, calcium binding protein, inducible by IL-1 β , IFN- γ , and IL-18, as well as T cell conditioned media [17-19]. In humans, the AIF-1 gene maps to the major histocompatibility complex class III region on chromosome 6p21.3. This is a region which is densely clustered with genes associated with the immune response, including TNF- α/β and NF-KB [20], suggesting a fundamental role of AIF-1 in the processing of the inflammatory responses. AIF-1 expression was first reported in 1995, in injured rat muscles in association with rat cardiac allograft rejection [21]. Subsequently AIF-1 was found in humans and is involved in many pathological processes including angiogenesis, cell proliferation, and differentiation [22,23]. It is expressed primarily in dendritic cells and macrophages [24]. In a blood vessel, AIF-1 was increased after balloon injury and angioplasty [25]. Overexpression of AIF-1 is associated with cardiac allograft vasculopathy [14,15,26]. In renal allograft undergoing a rejection episode the infiltrating macrophages with increased AIF-1 signals may distinguish clinical rejection as opposed to subclinical rejection [27,28].

There have been limited reports associated with the early innate immune responses and the parameters involved in the activation of early inflammatory responses after transplantation. Recent evidence suggests that myocardial cell-death and associated inflammation are characteristics of cardiac IRI. Thus, strategies aimed at identification of molecular markers released during IRI could help in the design of blocking experiments to prevent or minimize cardiac injury induced by I/R. Direct evidence identifying the release of AIF-1 from cardiomyocytes during IR has not been established. In this study we investigated the release of AIF-1 during an experimental rat model of IRI, in which the left anterior descending artery (LAD) was occluded at different time intervals. We have compared the level of AIF-1 present in peripheral blood and cardiac tissue specimens before and after IRI. The goal was to provide evidence that AIF-1 is induced during cardiac IR in myocardial tissue and is a crucial biomarker in the injury induced immune response that indicates the magnitude of the inflammatory response initiated by IRI.

Materials and Methods

Surgical preparation of rats

Male Sprague-Dawley rats aged 2-3 months; 300-350 grams were tested in this study. Rats were housed in a temperature; humidity and 12:12 hour light-dark cycle controlled room in the animal facility at the University of Mississippi Medical center (UMMC) before surgery. Two to three rats were kept in one cage, providing free access to tap water and rat chow. All surgeries were performed during the light cycle. The study was conducted according to the guidelines required by Institutional Animal Care and Use Committee (IACUC) for the use of live rodents at UMMC.

Surgical approach

The procedure was performed under an operating microscope with 10X magnification. Surgical protocol for myocardial ischemia reperfusion was adapted from a procedure previously described [29]. Rats were anesthetized in an anesthesia chamber and isoflurane delivery system. Originally with inhalation rates of 3-5 % isoflurane and 2% oxygen via an isoflurane vaporizer (Isotech, Highland Medical Equipment, Tempeca, CA) followed by 1-3% as needed. The rat was placed in the supine position on a heated pad that was fixed on the operating table. The legs of the rat were secured to the side of body with tape, making sure the front legs were not over-stretched, which it may affect respiration. The body temperature was maintained at $37.5^{\circ}\text{C} \pm 0.5^{\circ}\text{C}$ by the heating pad controller. A cotton umbilical surgical ribbon attached to the operating platform was placed horizontally under the front upper incisors to hold the upper jaw in place. The fur from surgical areas, which include: front of the neck, chest and the inner leg were shaved and decontaminated using Betadine, followed by a 70% ethanol swab.

For a successful oral-tracheal intubation, the trachea was exposed by making a 1.5 cm cervical incision, then spreading the lobes of the thyroid gland and the sternohyoideus muscle facilitating viewing of the trachea. Holding the tongue with gauze slightly upwards, an 18-gauge, blunt end cannula with polyethylene (PE) tube attachment was gently inserted through the opening of throat and was viewed through the exposed trachea. The cannula was then secured in the place by a 2-0 suture while being mechanically ventilated using a rodent respirator (SAR-830/AP small animal ventilator). The ventilation rate was adjusted to 55-60 breaths/min, the tidal volume was set at 2-3 ml/100 gram body weight. A left intercostal thoracotomy was performed by a 3-4 cm long incision at a site 0.5 cm away from the left sternal border. Before the ribs were cut, the 3rd- 5th ribs were clamped off with small curved hemostat to prevent the vessels underneath the ribs from bleeding. Then the ribs were spread with a small retractor to expose the heart. The pericardium was then opened with sharp forceps. The LAD, about 2-3 mm below the anterior-inferior edge of the left atrium was ligated with a tapered point needle and a 5-0 silk ligature. For a temporary ligature that can be removed during reperfusion, the ends of the tie was threaded through a 1 cm PE100 tube, which was placed parallel to the coronary artery to form a snare for reversible LAD occlusion. Thus, a local ischemic condition was achieved which was observed by a change in color of the anterior wall of the LV to paleness of myocardium (Figure 2).

Blood was drawn from the femoral vein, after making an incision in the inguinal area, along the angle of the left hind leg. Using blunt-tipped scissors, the connective tissue was spread out until the femoral vein was exposed. A 26-gauge butterfly blood drawing needle attached to a 2 ml sterile syringe was used to collect blood. The syringe was flushed with heparin then filled with 0.1 ml heparin solution (20U/ml), to draw 1-2 ml of blood. At the completion of blood drawing, the needle was gently pulled. A 1-cm in diameter gauze-ball was placed over the vein, secured with a 5-0 silk suture into the muscle layer, and the incision was stitched with 4-0 suture.

Experimental protocol

In a preliminary study, 25 rats were assigned to five groups (5 in

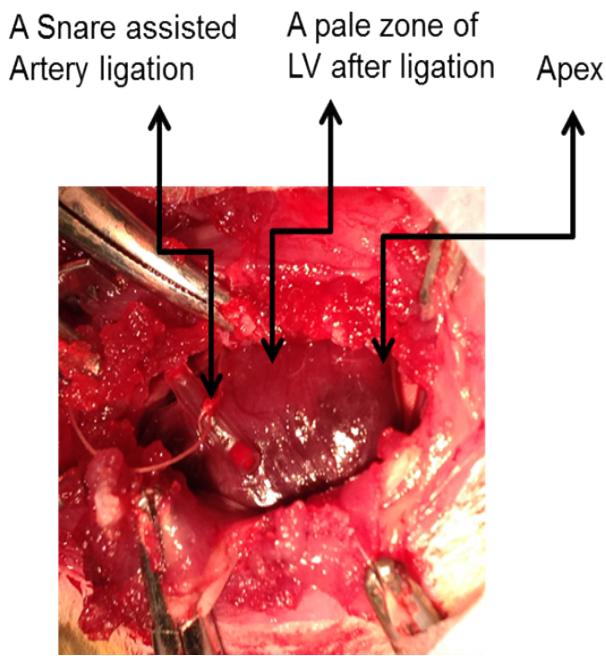


Figure 2: A model of rat LAD ligation and ischemia. A model of rat LAD occlusion and ischemia development is presented. A, demonstrates ischemic condition after 60 minutes, in which a change of color of the anterior wall of the LV to paleness is observed. B, is a video presentation of ischemia. For a temporary ischemia, a 1 cm PE tube was placed parallel to the coronary artery, the ends of the suture was looped around the tube to form a snare for reversible LAD occlusion.

Table 1: Study protocol for determination of AIF-1 release during ischemia vs Reperfusion.

Groups	Incision	Ischemia	Reperfusion
1. Control	yes	No	No
2. 30 minutes Ischemia	Yes	Yes	No
3. 60 minutes Ischemia	Yes	Yes	No
4. 30 minutes Ischemia	Yes	Yes	10 minutes reperfusion
5. 30 minutes Ischemia	Yes	Yes	60 minutes reperfusion

Table 2: Rat primer sequences (5'-3') for Reverse Transcriptase -PCR.

Genes	Sequences
AIF-1 forward primer	5'-agc-cca-gca-gga-aga-gag-gt-3'
AIF-1 reverse primer	5'-tgg-ggg-acc-agt-tgg-ctt-ct-3'
GAPDH forward primer	5'-cag-gga-aga-tgg-tga-gca-tt-3'
GAPDH reverse primer	5'-ctg-ctc-ctc-tgt-cat-ttc-ag-3'

each group): control; 30 minutes LAD occlusion; 60 minutes LAD occlusion; 30 minutes LAD occlusion/10 minutes reperfusion and 30 minutes LAD occlusion/60 minutes reperfusion as summarized in Table 1. In all rats a left thoracotomy was performed, but in the control group no further procedure was applied. In groups 2, 4, and 5 the LAD was circled with suture and the ligation was tied around a small tube as described earlier for reversible LAD occlusion. Then 0.2ml of 1000U/ml heparin in physiological saline was injected through the penile vein to reduce blood coagulation during surgery. In group 2 and 3 at the end of 30 minutes or 60 minutes LAD ligation (ischemia), respectively, blood was drawn for testing of AIF-1 expression levels in the blood. Blood was drawn directly from left ventricle then the heart was excised while the rat was deeply anesthetized (5% isoflurane). In group 4 and 5, the LAD was ligated for 30 minutes followed by 10 minutes or 60 minutes reperfusion respectively. Blood was drawn at the completion of reperfusion for AIF-1 mRNA analysis. The heart was then excised as described earlier, flushed with physiologic-saline, wrapped in saran wrap and stored at -20°C for 30 minutes. Frozen heart was then sliced in two halves along the interventricular septum. Each half of the left and right ventricles (LV and RV) were sliced in 2x2square mm, some stored in formaldehyde for paraffin block preparation, for immunohistochemistry (IHC) staining, or at -80°C freezer to be used for RNA preparation for AIF-1 mRNA transcript.

Measurement of AIF-1 expression in rat peripheral blood cells and cardiac tissue

To determine AIF-1 mRNA transcript levels, RNA was isolated from rat peripheral blood and cardiac tissue specimen. In human, the AIF-1 expression level varies in mononuclear cells (MNCs) versus polymorphonuclear cells (PMNCs). Thus, both types of the cells were tested in this rat model of IRI.

Blood samples were subjected first to Ficoll-paque density gradient for isolation of PBMCs (the interphase layer between plasma and Ficoll-paque). The dense white layer above the red cells (buffy coat) was then suspended in a hypotonic solution prepared from 0.3 M NH₄CL and 0.17 M Tris and layered over inactivated 100% Fetal Calf Serum, followed by centrifugation at 1,500rpm for 10 minutes at room temperature. The pink pellet, which contained PMNCs and residual amounts of red cells was washed 2X with 1X sterile phosphate buffered saline (PBS, pH=7.4), and adjusted to 2-5x10⁶cells/ml. The pellets were immediately homogenized in TRIzol and were stored at -80°C for subsequent RNA extraction. The PBMCs and PMNCs were at ≥95% purity observed by differential cell count stain, and ≥95% viable with 0.4% Trypan blue stain.

RNA from PBMCs and PMNCs were extracted using "Pure Link" RNA kit (Life Technologies, Carlsbad, CA, USA), according to the manufacturer's guidelines. RNA concentration and purity was determined using NanoDrop 2000C (Thermo-Fisher, GA, USA). For RNA preparation from cardiac tissues, frozen sections were homogenized in 800μl TRIzol and extracted using the same "Pure Link" kit. Afterward, the RNA was transcribed using the ImProm-II Reverse Transcription kit (Promega, Madison, WI), followed by semi-quantitative RT-PCR (95°C for 4 min, and a 25 cycles of amplification at 95°C/30sec, 60°C/30sec, 72°C/30sec, and 100nM of each forward and reverse primers) using primer sequences specific for rat AIF-1, and GAPDH shown in Table 2. The amplification products were analyzed in Alpha Innotech System using software v.6.0.14 (Proteinsimple, Santa Clara, CA USA). The mRNA transcript levels were normalized to the GAPDH gene expression levels. The normalized values (Unit) were presented as mean ± SEM.

Immunohistochemistry (IHC)

The formalin fixed paraffin embedded (FFPE) rat cardiac tissue sections were prepared for IHC stain as previously described [15]. The slides were stained with L5-B3662 primary antibody against AIF-1 (ID#1022-5, IgG2b, reactive in human and rat) (Life Span Bioscience Inc., Seattle, WA). The optimal working dilution for LS-B3662 was 1:50. The detail of the staining was previously reported [26]. The stained slides were evaluated by a pathologist to confirm the staining specificity.

Data analysis

Both interstitial infiltrating MNCs and cardiac myocytes (CMCs) were evaluated by IHC staining. Sections were analyzed for the presence and absence of AIF-1 in relation with clinical condition (ischemia vs. reperfusion). The mRNA transcript values were expressed as the mean ± SEM. The group differences for mRNA transcript levels between tissue conditions were determined by one-way analysis of variance. The level of significance was set at p<0.05 for comparison between the groups.

Results

Basic Measurements

A total of 25 rats were studied. The average body weight was 328 ± 21.5. The heart rates (HRs) and blood pressure were not recorded. The HR was stable after the initial anesthesia, and the follow up period (1-2% isoflurane and 1% oxygen). There were no drastic changes among the five groups. In the control group, in which the rats had undergone only a left thoracotomy, no significant HR changes were noticed throughout the observation. In other groups a significant increase of HR was observed during the first 1-5 minutes into ischemia condition,

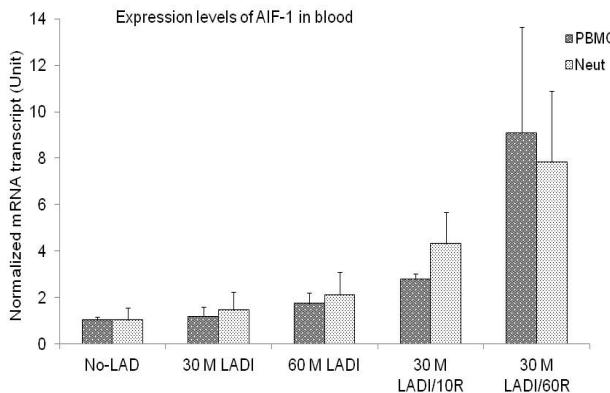


Figure 3: mRNA transcripts of AIF-1. Variations in mRNA transcripts levels were expressed as the mean \pm SEM. AIF-1 mRNA transcripts at different time intervals are shown in PBMCs and Neutrophils/ or PMNCs. The levels of expression were increased after ischemia compared with controls but was not statistically significant. AIF-1 expression was significantly increased after 10 or 60 minutes of reperfusion observed in PBMCs and Neutrophils. For 30 minutes of ischemia (LAD)/10 minutes of reperfusion vs. control the p values were (PBMCs, $p<0.05$, Neutrophils, $p<0.03$). For 30 minutes of ischemia (LAD)/60 minutes of reperfusion vs. control the p values were (PBMCs, $p<0.01$, Neutrophils, $p<0.01$).

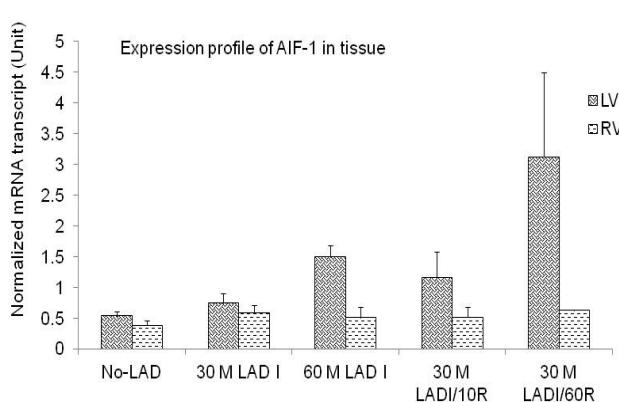


Figure 4: mRNA transcripts of AIF-1. Measurement of AIF-1 expression in cardiac tissues expressed as the mean \pm SEM. After 30 minutes of ischemia AIF-1 mRNA was increased 1.4-fold in LV tissues and 1.5-fold in RV tissues compared to control. After 60 minutes of ischemia AIF-1 mRNA was increased 3.4-fold in LV tissues and 1.3-fold in RV tissues compared to control ($p<0.05$). After 10 minutes of reperfusion AIF-1 mRNA was reduced but it was 2.3-fold greater than the levels in control and 2.2-fold greater than in RV tissues ($p<0.05$). After 60 minutes of reperfusion AIF-1 mRNA was elevated 6.24-fold in LV tissues vs. control ($p<0.01$), and 5.13-fold greater than the levels in RV tissues ($p<0.01$).

but then returned to a stable rate after oxygen was increased to a rate of 1.5–2%. During the reperfusion, there were no significant changes in HR among the group that had undergone reperfusion.

Rat model of LAD ligation and ischemia

The fate of an experimental coronary occlusion is highly dependent on the ideal ligation position. Thus, the procedure was performed under an operating microscope. In the groups with I/R some rats demonstrated a minimal ventricular fibrillation (VF), or ventricular tachycardia (VT), during the early period of both ischemia and reperfusion. The VF or VT did not have impact on the course of the study. A video of a LAD occlusion is shown in Figure 2, where myocardial color changes are visible on the left ventricular wall during ischemic condition.

Measurement of AIF-1 expression in peripheral blood cells

AIF-1 mRNA was tested in PBMCs and PMNCs. The normalized values (Unit) were presented as mean \pm SEM. The levels of expression were increased after 30 minutes and 60 minutes of ischemia as compared with control group in both PBMCs and PMNCs (Controls: 1.03 ± 0.14 and 1.02 ± 0.54 respectively; 30 min LADI: 1.18 ± 0.4

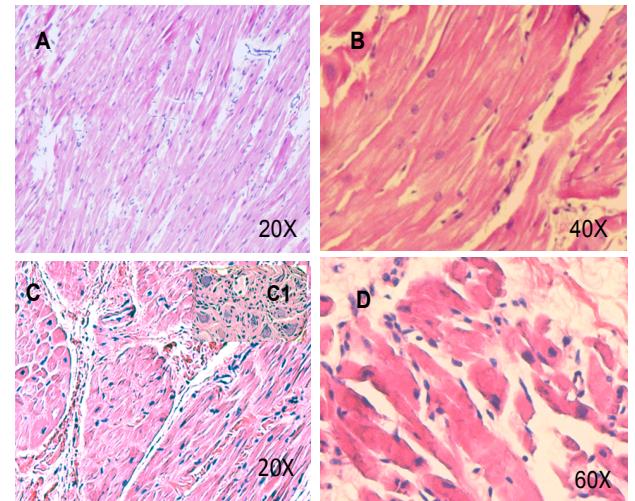


Figure 5: Cross-sections of cardiac LV after ischemia reperfusion. A representative HE staining of LV tissues from control and post IR is shown. A and B, a 20x and 40x illustration of LV from control group, in which myocardial fibers are arranged regularly with clean striation and no pathological changes are observed; C and D, presents after IR in which degenerative changes in myocardial fibers and inflammatory cells in the perimeter of myocytes are observed.

and 1.46 ± 0.75 ; 60min LADI: 1.74 ± 0.45 and 2.1 ± 0.97). However, the data was not statistically significant (Figure 3). Compared to the control group, AIF-1 expression level was significantly increased in the group with 30 min ischemia followed by 10min or 60min reperfusion in both PBMCs and PMNCs (Controls: 1.03 ± 0.14 and 1.02 ± 0.54 respectively; 30 M LADI/10 M R: 2.73 ± 0.23 and 4.3 ± 1.35 ($p<0.05$ and $p<0.03$ respectively); 30 M LADI/60 M R: 9.1 ± 5.56 and 7.83 ± 3.1 ($p<0.01$ and $p<0.01$ respectively).

Measurement of AIF-1 expression in cardiac tissues

We examined mRNA expression levels in rat cardiac LV and RV tissues in all five groups. As shown in Figure 4, the mRNA expression levels were increased compared to control specimens. In group 2, the AIF-1 mRNA was increased 1.4-fold in LV tissues and 1.5-fold in RV tissues as compared with group 1. In group 3, the mean AIF-1 mRNA was 3.4-fold higher in LV as compared to 1.32-fold in RV tissues ($p<0.05$). In group 4, after 10 min reperfusion the AIF-1 levels in LV tissues were 2.3-fold greater than the levels in tissues from group 1, and 2.2-fold greater than RV tissues ($p<0.05$). In group 5, the AIF-1 mRNA was increased 6.24-fold in LV tissues as compared to the levels in group 1 ($P<0.01$). In addition, the mean \pm SEM levels in LV tissues were 5.13-fold greater than the levels in RV tissues ($P<0.01$).

Histopathologic examination and distribution of AIF-1

To localize the expression of AIF-1 after I/R, rat LV tissues were tested by histology and immunostaining. Figure 5, shows a representative hematoxylin and eosin (HE) staining of LV tissues from control and post IR groups. In control group LV myocardial fibers were arranged regularly with clear striation (Figure 5A,5B). No pathological changes were observed. In the post IR group, degenerative changes in myocardial fibers and inflammatory cells were seen in the perimeter of myocytes, which are an indication of myocyte injury (Figure 5C,5D). Immunohistochemistry staining for AIF-1 is shown in Figure 6, section A and section B. Anti-AIF-1 stain as present in both the nucleus of myocytes and in interstitial cells. AIF-1 positive cells were increased after reperfusion (Figure 6B).

Discussion

The scope of this study was to verify the release of AIF-1 after experimental LAD occlusion followed by reperfusion. It has been documented that a series of pathophysiologic and immunologic events occurs when tissue or organ has undergone IRI [30,31]. IRI increases the systemic availability of inflammatory mediators, leading to the activation of myocardial endothelial cells, as well as infiltrating

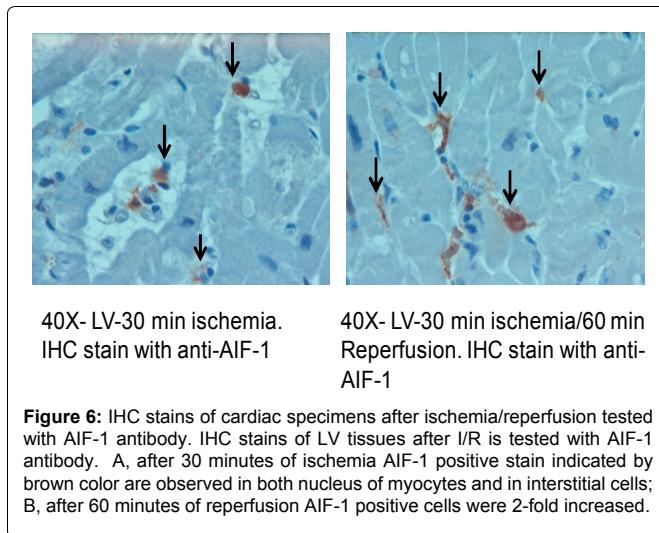


Figure 6: IHC stains of cardiac specimens after ischemia/reperfusion tested with AIF-1 antibody. IHC stains of LV tissues after I/R is tested with AIF-1 antibody. A, after 30 minutes of ischemia AIF-1 positive stain indicated by brown color are observed in both nucleus of myocytes and in interstitial cells; B, after 60 minutes of reperfusion AIF-1 positive cells were 2-fold increased.

MNCs and PMNCs initiating vascular and organ dysfunction. In addition molecular markers associated with innate immunity have been implicated in the regulation of inflammatory responses during IRI [32,33]. In this study we have generated ischemic condition followed by reperfusion in a rat *in vivo* model of LAD occlusion. The practical value of this study was the findings that the AIF-1 was variably expressed in cardiac tissues after IRI. Our data revealed that AIF-1 was up regulated as early as 10 minutes after reperfusion, and further it was increased several-fold after 60 minutes in PBMCs and PMNCs as compared to the control group. Similarly, expression levels of AIF-1 in LV tissues were significantly increased after 60 minutes ischemia or 30 minutes ischemia followed by 60 minutes reperfusion (Figure 4). This indicates that AIF-1 was up regulated during both ischemia and reperfusion. In addition AIF-1 level was 2-fold greater in LV tissues after 60 minutes reperfusion compared to LV after 30 or 60 minutes ischemia, suggesting that AIF-1 upon induction is actively expressed during IRI. This powers the impact of AIF-1 during IRI and thereafter on organ dysfunction.

Recent studies have reported increased levels of AIF-1 expression in IRI [34,35]. In the latter study using human cardiac atrium tissues *in vitro*, it was shown that AIF-1 levels increased in association with IL-18, TLR2, and TLR4 at different time intervals. IL-18 and TLRs are essential receptor molecules on macrophages. They are known for their association with early innate immune responses and several studies have reported the presence of these inflammatory markers in association with allograft rejection [35,36]. On the other hand, AIF-1 is known for its participation in inflammatory responses and regulation of inflammatory pathways of macrophages [37,38]. This was supported by the studies in which AIF-1 cDNA was transfected into a macrophage cell line, Raw264.7. The transfectant cells with over expressed AIF-1 had characteristics of the dendritic cells, with significant production of IL-6, IL-10 and IL-12 upon stimulation with LPS [37]. Moreover, blocking the expression of AIF-1 by siRNA was associated with apoptosis of AIF-1 bearing macrophages [38,39]. Macrophages play a significant role in linking injury, inflammation, and organ dysfunction [40]. The level of AIF-1 expression in rat macrophages has been shown to be augmented by IFN- γ and IL-1 β [17,18]. Furthermore, it was shown in a rat model of IRI AIF-1 expression was parallel with the expression levels of IFN- γ after reperfusion, thus suggesting the immunologic involvement of AIF-1 in proinflammatory processes of macrophages. In our unpublished studies of a rat model of IRI, as well as in *in vitro* studies of human cardiac tissue IR [34], we were unable to demonstrate increased expression levels of IFN- γ . However, we observed a significant increase in the levels of TLR-2 and TLR-4 mRNA transcripts in cardiac tissues after IR. The TLRs particularly TLR-2, -3, -4, and -9 have been shown to be involved in the pathophysiology of IRI [41-43]. Although we have not tested expression levels of TLRs in rat cardiac tissues after IRI, one study reported the impact of TLR-3 in myocardial injury through the recognition of DAMPs during myocardial IRI [44].

We do not know whether AIF-1 is just a marker of immune effector cells associated with inflammation, or directly engaged in regulation of inflammatory processes in cardiomyocytes and resident dendritic cells/macrophages during IR. It has been shown that increased levels of AIF-1 in liver-allograft after cold and warm IR could be reduced with low dose of FK-506 pretreatment [45]. In addition, studies have shown that reducing inflammatory responses during IR significantly protects organ function [46-48].

The functional role of AIF-1 remains to be determined. Hypothetically there is growing evidence suggesting that allograft injury during organ procurement induces innate immune response in which donor-derived damaged molecules may serve as endogenous ligands for TLRs in the activation of an initial inflammatory event. This is an immunologic concept that was originally proposed by Janeway, 1989 in which he described how the innate immune system discriminates noninfectious self from infectious non-self [49]. Later the concept was retackled by Polly Matzinger [50]. These self-derived molecules are known as “danger signals” and are associated with non-pathogenic, host originated damaged tissues causing sterile inflammation. In a similar context we are considering that a segment of AIF-1 may represent a potential candidate ligand and may initiate a danger signal, leading to inflammatory responses. Such a model of cardiac allograft IRI is currently under investigation.

This study was the first to investigate expression levels of AIF-1 in a time-dependent manner from myocardial tissues and peripheral distribution after IRI. Considering the involvement of AIF-1 in the initial inflammatory processes, the specific inhibition or minimizing the expression levels during ischemia may have important clinical implications in transplantation. However, whether such inhibition can protect the cardiac tissue from injury remains to be established. Emerging studies have reported the significance of post-ischemia reconditioning in combination with inflammatory agonist that may attenuate myocardial IRI by inhibiting reperfusion induced pro-inflammatory cytokines [48,51]. Thus, identification of signals and/or activators of myocardial IRI and signaling pathways, which leads to early inflammatory response, suggests that AIF-1 and TLRs may prove efficient therapeutic targets for pharmacologic agents. While the inhibition of such markers will remain crucial for treatment of IRI and organ protection, many other aspects of allograft rejection and allograft tolerance induction need further investigation.

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Ethical Statement

The study was conducted according to the guidelines required by Institutional Animal Care and Use Committee (IACUC) for the use of live rodents at the University of Mississippi Medical Center.

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