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Inflammation in cardiac injury, repair and regeneration

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Abstract

Purpose of the review—Cardiomyocyte necrosis activates an inflammatory response that serves to clear the injured myocardium from dead cells, and stimulates repair, but may also extend injury.

Recent findings—Recently published studies have identified Interleukin (IL)-1 α and RNA released by necrotic cardiomyocytes as key danger signals that trigger the inflammatory response following infarction. IL-1 promotes activation of a pro-inflammatory phenotype in leukocytes and fibroblasts, and delays myofibroblast transdifferentiation. Inhibitory lymphocytes play a crucial role in negative regulation of the post-infarction inflammatory response by modulating macrophage and fibroblast phenotype. Cardiac macrophages exhibit significant heterogeneity and phenotypic plasticity and may orchestrate the reparative response following infarction. In neonatal mice, resident embryonic macrophage subpopulations may promote a regenerative response. In contrast, in adult animals replacement of resident macrophage populations with monocyte-derived macrophages may induce inflammation while inhibiting cardiac regeneration. These exciting observations highlight the crucial role of macrophages in cardiac injury and repair, but should be interpreted with caution considering the limitations of murine models of neonatal myocardial injury.

Summary—Design of novel strategies to reduce cardiac injury, improve repair and promote regeneration is dependent on understanding of the cell biology of the inflammatory response.

Keywords

inflammation; myocardial infarction; cytokine; macrophage; lymphocyte

Introduction

Cardiomyocyte necrosis triggers an intense inflammatory cascade that serves to clear the injured myocardium from dead cells and sets the stage for activation of the reparative process. In myocardial infarction, necrotic cardiomyocytes release alarmins, activating innate immune signals, and inducing recruitment of pro-inflammatory leukocytes. Although inflammation is required for phagocytotic removal of dead cells and for activation of

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reparative mesenchymal cells, timely suppression and spatial containment of the inflammatory reaction is necessary to prevent extension of injury. Overactive, dysregulated, temporally prolonged, or spatially expanded inflammatory responses may cause death of viable cardiomyocytes, enhance matrix degradation (thus promoting dilative remodeling), and extend fibrosis[1]. The inflammatory cascade in myocardial infarction and heart failure is an attractive therapeutic target; however, implementation of anti-inflammatory strategies is challenging due to the pleiotropic and multifunctional effects of inflammatory mediators that may activate both injurious and reparative processes. Thus, dissection of the inflammatory signals implicated in cardiac injury and repair and identification of the effector cells involved in regulation of inflammation is of outstanding significance. This manuscript will present recent advances that significantly contributed to our understanding of myocardial inflammation.

Activation of pro-inflammatory cascades following cardiac injury. The “danger signals”

Necrotic cardiomyocytes are capable of releasing a wide range of damage-associated molecular patterns (DAMPs) that activate innate immune pathways triggering the inflammatory response (Figure 1)[2]. Although high mobility group box-1 (HMGB1) has been suggested as important cardiomyocyte-derived alarmin[3], [4]; the identity of the key primary stimulus that may trigger the inflammatory reaction following infarction remains unknown. Using a combination of in vivo and in vitro approaches, Lugrin and co-workers[5] demonstrated that necrotic cardiomyocytes release Interleukin (IL)-1 α , a key early damage signal that activates the abundant fibroblasts in the myocardium, promoting their pro-inflammatory activation through MyD88-dependent, toll like receptor (TLR)-independent signaling (Figure 1). IL-1 signaling is critically implicated in activation of the post-infarction inflammatory response: in addition to IL-1 α released by necrotic cells, de novo synthesis and activation of IL-1 β stimulates type 1 IL-1 receptor (IL-1R1) signaling in pro-inflammatory leukocytes and in fibroblasts[6]. In both leukocytes and fibroblasts, activation of IL-1 signaling induces cytokine expression and promotes matrix-degrading properties[7]. Moreover, IL-1 suppresses fibroblast proliferation[8] and inhibits transdifferentiation of fibroblasts into myofibroblasts[7], [9], delaying activation of a reparative response until the infarct is cleared from dead cells and matrix debris. IL-1 α is not the only endogenous danger signal released by cardiomyocytes in the infarcted region. Chen et al identified release of RNA (including several microRNAs) by necrotic cardiomyocytes as an important injury-activated pro-inflammatory stimulus in the infarcted heart[10]. Extracellular RNA contributed to myocardial inflammation by activating a TLR-3/Toll/IL-1 receptor domain-containing adaptor inducing IFN- β (Trif) cascade. In the pressure overloaded heart, mitochondrial DNA that escapes from autophagy has been reported to trigger TLR-9-dependent inflammation extending myocardial injury[11]. Whether a similar pro-inflammatory pathway is activated in the infarcted heart remains unknown. The cellular targets of these danger signals include cardiomyocytes, vascular cells, fibroblasts and infiltrating leukocytes[12]. Whether specific alarmins activate distinct cell types driving injurious or reparative functions remains unknown. A growing body of evidence suggests that activation of innate immune signaling in hematopoietic cells may exert deleterious

actions, extending injury. Bone marrow transplantation experiments in mice suggested that activation of TLR2, or stimulation of complement-activated responses in leukocytes extend ischemic injury[13, 14].

Removal of apoptotic cells as a mechanism for suppression of post-infarction inflammation

Repair of injured tissues is dependent on timely suppression and resolution of pro-inflammatory signaling. In the healing infarct, negative regulation of the inflammatory response is critical for protection of the infarcted heart from adverse remodeling. Clearance of dead cells by phagocytes activates an inhibitory program, serving as a key mechanism for termination of the pro-inflammatory cascade. Wan and co-workers provided the first demonstration of a critical role for cardiomyocyte efferocytosis (the process by which dying cells are removed by phagocytes) in suppression of acute inflammation following myocardial infarction[15]. Induction of macrophage myeloid-epithelial-reproductive tyrosine kinase (MERTK) in myocardial phagocytes was necessary for effective engulfment and removal of apoptotic cardiomyocytes from the infarct. Disruption of cardiomyocyte efferocytosis resulted in prolonged inflammation and accentuated adverse remodeling. Whether clearance of other apoptotic cell types involves distinct pathways remains unknown. In the healing infarct, the majority of apoptotic cells are non-cardiomyocytes; infiltrating neutrophils represent a large pool of cells programmed to undergo apoptosis. Their clearance from the infarcted heart may also activate an inhibitory program leading to resolution of inflammation.

Lymphocyte subpopulations: key effector cells implicated in negative regulation of inflammation

Although descriptive studies in large animal models have demonstrated that the infarcted hearts recruits lymphocytes expressing anti-inflammatory cytokines (such as IL-10)[16], the critical role of lymphocyte subpopulations in regulation of the inflammatory and reparative response has only recently been appreciated. Zouggari and co-workers demonstrated that B cells produce the CC chemokine CCL7 and mobilize pro-inflammatory monocytes in mouse infarcts increasing cardiomyocyte injury[17]. During the reparative phase, lymphocyte subsets with inhibitory properties, such as regulatory T cells (Tregs) are recruited in the infarct through pathways that may involve distinct chemokine/chemokine receptor pairs[18]. Recent studies using both genetic and antibody-mediated depletion strategies have suggested a critical role for Tregs in suppression of the inflammatory reaction following myocardial infarction[19]. Tregs may act by promoting differentiation of macrophages to an anti-inflammatory phenotype[19] and by modulating protease synthesis by cardiac fibroblasts[20]. The protective actions of Tregs may involve both secreted mediators and contact-dependent cell-cell interactions[21]. Selective activation of invariant Natural killer T cells (iNKT cells) through administration of α -galactosylceramide also suppressed inflammation and attenuated injury in experimental models of myocardial infarction[22], [23]. These studies suggest that, despite their small numbers[20], [24], infiltrating T cell

subpopulations are capable of exerting profound effects on many other cell types involved in cardiac inflammation, repair and remodeling. (Figure 2)

Are surviving cardiomyocytes a source of protective cytokines in the healing infarct?

In the early hours after infarction, necrotic cardiomyocytes release danger signals that initiate inflammation and may extend injury. However, spatial containment of the inflammatory response into the area of infarction is critical for optimal repair, and may be dependent on activation of inhibitory pathways in the border zone[25], [26]. Do viable cardiomyocytes in the infarct border zone secrete cytoprotective and anti-inflammatory mediators that may serve as a protective “barrier” preventing expansion of the inflammatory response? A recent study by Rainer and co-workers suggests that cardiomyocytes may secrete a wide range of cytoprotective mediators; expression of these signals may be regulated by the local cytokine environment[27]. Mice with cardiomyocyte-specific loss of Transforming Growth Factor (TGF)- β signaling were protected against rupture through an increase in levels of protective mediators, such as IL-33, Thrombospondin-4 and growth differentiation factor-15. The findings highlight the pleiotropic actions of TGF- β , suggesting that in addition to its effects on leukocytes, fibroblasts and vascular cells[28], it may also inhibit activation of a protective program in cardiomyocytes of the infarct border zone.

The cardiac macrophages

Despite their abundance in the infarcted heart, macrophages have remained understudied, in part due to their functional heterogeneity[29]. Over the last 2 years, several high-impact investigations have shed new light into the role of macrophages in cardiac homeostasis, injury and repair. The adult mouse heart contains a heterogeneous population of macrophages that can be subclassified into several subsets with distinct ontological origins [30], [31]. The cardiac macrophage population is derived from yolk-sac macrophages and from fetal monocyte progenitors and is maintained through local proliferation. As mice age, these resident macrophages are progressively replaced by monocyte-derived cells, even in the absence of inflammatory stimulation[32]. Following infarction, the heart is infiltrated by abundant macrophages; the majority of these newly recruited cells are derived from circulating Ly-6C(high) monocytes attracted in the healing infarct through interactions involving the CC chemokine monocyte chemoattractant protein-1/CCL2[33], [34], [35]. Infiltrating monocytes may be derived not only from the bone marrow, but also from the spleen[36], [37]. Expression of the transcription factor interferon regulatory factor 5 (IRF5) is increased in macrophages during the inflammatory phase of cardiac repair and drives a pro-inflammatory program[38]. As the infarct heals, reparative Ly-6C(low) F4/80 (high) macrophages, derived from the Ly-6C(high) monocyte subpopulation, proliferate and express inhibitory mediators, promoting repair[34]. If the majority of infarct macrophages are derived from monocytes, what is the fate of the resident cardiac macrophage subpopulations? Much like for other populations of interstitial cells, the susceptibility of macrophages to ischemic injury remains unknown. Evidence suggests that some resident macrophages may undergo apoptosis following permanent coronary occlusion[35]. Whether

there is complete removal and replacement of the original resident macrophage population remains unknown.

Do macrophages hold the key to cardiac regeneration?

Recent experimental evidence suggests that macrophages may be critically involved in cardiac repair and regeneration. In an apical resection model, neonatal mice exhibited myocardial regeneration; this capacity was lost at 7 days of age[39]. Depletion of macrophages abrogated regeneration of the neonatal heart, suggesting that neonatal cardiac macrophages exhibit a unique regenerative phenotype and secrete soluble mediators that contribute to formation of new myocardium[40]. Similar observations were noted in a model of genetic cardiomyocyte ablation driven by a diphtheria toxin receptor-based system. In neonatal mice, cardiomyocyte ablation resulted in expansion of embryonic-derived resident macrophages that promoted cardiomyocyte proliferation. In contrast, in adult hearts, recruitment of pro-inflammatory monocytes inhibited repair. These findings suggested that embryonic resident macrophage populations may be essential cellular effectors in cardiac regeneration[41].

Although the emerging body of evidence suggesting a potential role of macrophage subsets in cardiac regeneration is of outstanding interest, caution is advised regarding interpretation of the findings. First, the observations on regeneration of the neonatal heart are predominantly derived from experimental models with a pathology that is quite different from infarction of ischemic origin. In the apical resection model, removal of cardiac tissue eliminates the main source of “danger signals”, responsible for activation of inflammation in infarctive cardiac injury. Myocardial regeneration may be much more complicated in an environment that contains abundant dying cells and fragmented matrix proteins. On the other hand, in the cardiomyocyte ablation model, preservation of vascular and interstitial cells, and possible differences in the mechanism of cardiomyocyte death, may result in distinct reparative responses that do not recapitulate repair of ischemic infarction. Second, to what extent the observations made in neonatal mice can be generalized in higher mammals, and in particular in humans, is unknown. The profile and properties of macrophages in human hearts has not been systematically investigated. Moreover, whether human neonatal macrophages exhibit reparative and regenerative properties has not been tested.

Clinical implications

The negative experience with implementation of early anti-inflammatory strategies in myocardial infarction[42] and the complex effects of inflammatory pathways in injury and repair have discouraged attempts to target inflammation in heart disease. In patients with overactive, or dysregulated, post-infarction inflammatory responses, inhibition of inflammation may be a promising strategy to attenuate adverse remodeling. Anti-IL-1 approaches are particularly attractive and have already been tested in pilot studies with promising results. Treatment of patients with ST elevation myocardial infarction with recombinant IL-1 receptor antagonist attenuated adverse remodeling [43] and reduced the incidence of heart failure [44]. Approaches targeting the MCP-1/CCR2 system may selectively inhibit pro-inflammatory monocytes preventing adverse remodeling[45].

Identification of patients with excessive post-infarction inflammation using suitable biomarkers, or imaging approaches, may be needed for optimal selection of patients likely to benefit from anti-inflammatory strategies[46]. Expansion of our understanding on the immune cell subsets that promote cardiac regeneration may lead to identification of new cell therapy approaches to generate new myocardium.

Conclusions

Cardiomyocyte death activates resident myocardial populations that serve to sense injury and to repair the damaged heart. These non-cardiomyocyte populations exhibit remarkable plasticity and undergo dynamic changes in the environment of the infarct. Moreover, danger signals released by dying cells recruit leukocyte subsets that play a critical role in clearance of the wound and in activation of a reparative program. Studies published over the last 12 months suggested that unique subpopulations of macrophages may be capable of inducing myocardial regeneration, implying that the inflammatory reaction to myocardial injury can be modulated towards a regenerative path. A lot remains to be learned regarding these leukocyte subsets, the basis for their reparative properties, and the pathways driving their differentiation. However, these observations stress the importance of the non-cardiomyocyte compartment in determining the fate of cardiomyocytes, and highlight the potential therapeutic benefits that could be gained by unraveling the secrets of the inflammatory response.

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Abbreviations

DAMPs	Damage-associated molecular patterns
HMGB1	high mobility group box-1
IL	Interleukin
MyD88	myeloid differentiation primary response gene 88
TLR	toll-like receptor
Trif	IL-1 receptor domain-containing adaptor inducing IFN- β
Mertk	macrophage myeloid-epithelial-reproductive tyrosine kinase
Tregs	regulatory T cells
iNKT cells	invariant Natural killer T cells
(TGF)-β	Transforming Growth Factor
(IRF5)	interferon regulatory factor 5

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Key points

- Necrotic cardiomyocytes release danger signals that activate innate immune pathways and may extend injury, but also serve a crucial role in clearance of dead cells and in cardiac repair
- Lymphocyte subpopulations play an important role in negative regulation of the post-infarction inflammatory response.
- Macrophage subsets orchestrate reparative responses following cardiac injury.
- Neonatal resident cardiac macrophage may promote cardiac regeneration.

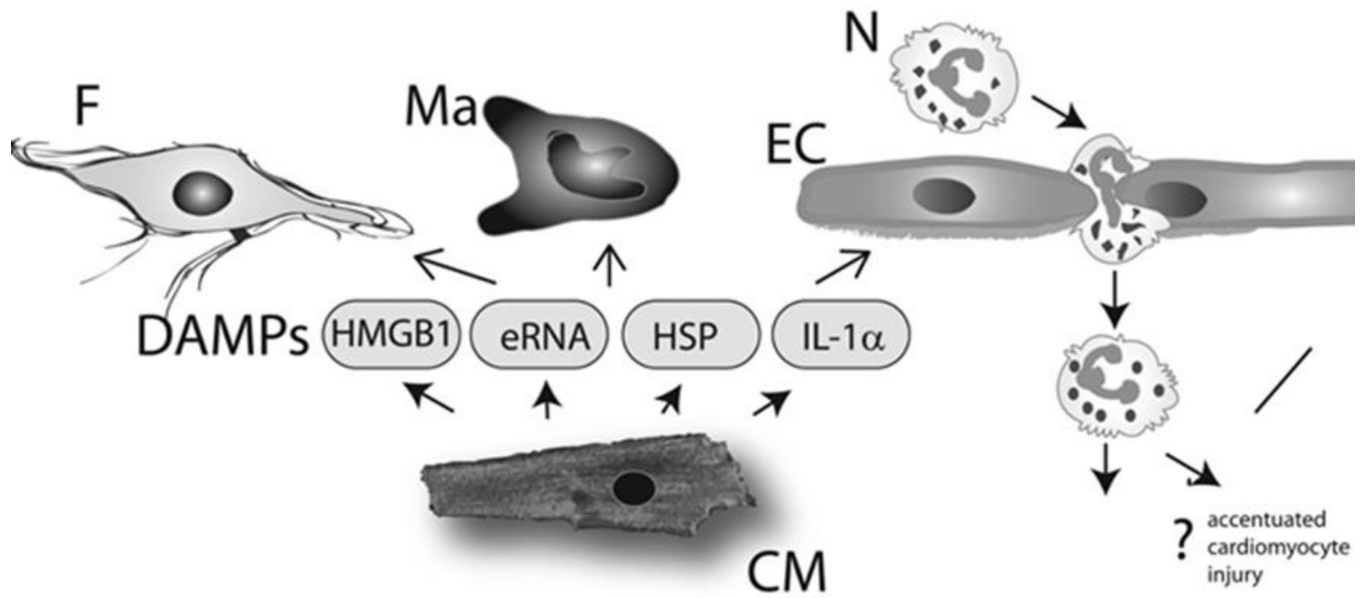


Figure 1.

Release of damage-associated molecular patterns (DAMPs) in the infarcted myocardium initiates the inflammatory response and may extend injury. High mobility group box-1 (HMGB-1), extracellular RNA (eRNA), heat shock proteins (HSP) and Interleukin (IL)-1 α are released by necrotic cardiomyocytes (CM) and activate innate immune signaling pathways in leukocytes (N, neutrophil; Ma, macrophage), endothelial cells (EC) and fibroblasts (F).

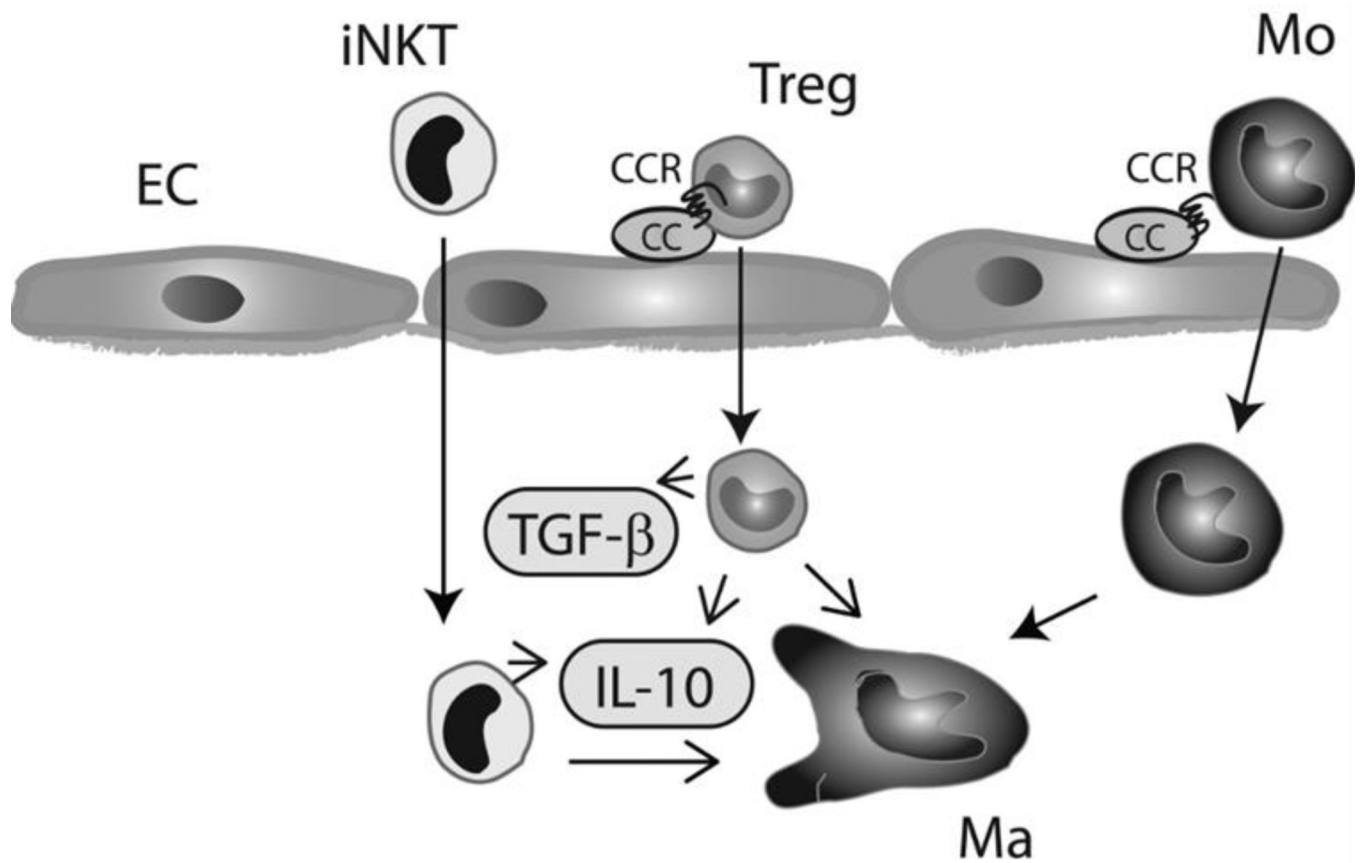


Figure 2.

T cell subsets, such as regulatory T cells (Tregs) invariant Natural killer T cells (iNKT) are recruited in the infarct through CC chemokine-dependent pathways and may be important cellular effectors in negative regulation of the post-infarction inflammatory response. Lymphocyte subpopulations may modulate macrophage (Ma), endothelial cell (EC), and fibroblast phenotype by secreting soluble mediators (such as IL-10 and TGF- β), or through contact-dependent interactions (Mo, monocyte).