



Published in final edited form as:

Transl Res. 2016 January ; 167(1): 152–166. doi:10.1016/j.trsl.2015.07.002.

Inflammation as a therapeutic target in myocardial infarction: learning from past failures to meet future challenges

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Abstract

In the infarcted myocardium, necrotic cardiomyocytes release danger signals, activating an intense inflammatory response. Inflammatory pathways play a crucial role in regulation of a wide range of cellular processes involved in injury, repair and remodeling of the infarcted heart. Pro-inflammatory cytokines, such as tumor necrosis factor- α and interleukin (IL)-1, are markedly upregulated in the infarcted myocardium and promote adhesive interactions between endothelial cells and leukocytes, by stimulating chemokine and adhesion molecule expression. Distinct chemokine/chemokine receptor pairs are implicated in recruitment of various leukocyte subpopulations in the infarcted myocardium. Over the last 30 years, extensive experimental work has explored the role of inflammatory signals and the contributions of leukocyte subpopulations, in myocardial infarction. Robust evidence derived from experimental models of myocardial infarction has identified inflammatory targets that may attenuate cardiomyocyte injury, or protect from adverse remodeling. Unfortunately, attempts to translate the promising experimental findings to clinical therapy have failed. This review manuscript discusses the biology of the inflammatory response following myocardial infarction, attempts to identify the causes for the translational failures of the past, and proposes promising new therapeutic directions. Because of their potential involvement in injurious, reparative and regenerative responses, inflammatory cells may hold the key for design of new therapies in myocardial infarction.

INTRODUCTION

In myocardial infarction, coronary artery occlusion causes ischemia of the territory subserved by the vessel, eventually leading to death of up to a billion cardiomyocytes (1). The adult mammalian heart has negligible regenerative capacity; thus, acute myocardial infarction is associated with loss of a large amount of myocardium that is replaced by a collagen-based scar. Necrosis of ischemic cardiomyocytes triggers an intense inflammatory

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The authors have no conflicts to disclose. All authors have reviewed and approved the manuscript, and have read the journal's policy on disclosure of potential conflicts of interest and the journal's authorship agreement.

reaction that serves to clear the wound from dead cells and matrix debris, and contributes to formation of a collagen-based scar (2). Abundant leukocytes infiltrate the infarcted myocardium and are predominantly localized in the infarct border zone where they may interact with viable cardiomyocytes. In the 1980s and 1990s a large body of experimental evidence suggested that inflammatory leukocytes may extend ischemic injury, exerting potent cytotoxic effects on border zone cardiomyocytes (3). These observations generated significant enthusiasm regarding the potential use of targeted anti-inflammatory strategies to reduce infarct size and to attenuate injury following myocardial infarction. Unfortunately, clinical trials inhibiting leukocyte integrins and the complement cascade in order to attenuate post-infarction inflammation were disappointing (4),(5). The failures of the clinical studies markedly dampened enthusiasm regarding the translational potential of the field. This is unfortunate, because inflammatory signaling is implicated in repair and remodeling of the infarcted heart. Thus, targeting inflammatory mediators may exert beneficial actions by attenuating dilative remodeling of the infarcted heart. Moreover, recent observations suggesting cytoprotective and regenerative actions of inflammatory signals have (once again) fueled interest in the field.

This review manuscript deals with the potential role of the inflammatory cascade as a therapeutic target in myocardial infarction. After a brief overview of the cellular effectors and molecular signals implicated in the post-infarction inflammatory reaction, we will discuss promising therapeutic approaches targeting the inflammatory response. Dissection of the molecular signals regulating induction and resolution of post-infarction inflammation needs to be complemented by understanding of the pathophysiologic complexity of the clinical context, in order to design effective therapeutic approaches for patients with myocardial infarction.

THE INFLAMMATORY RESPONSE IN MYOCARDIAL INFARCTION

Healing of the infarcted heart can be divided in three distinct but overlapping phases: the inflammatory phase, the proliferative phase and the maturation phase (6). During the inflammatory phase, danger signals (alarmins) released from dying cardiomyocytes activate innate immune pathways inducing chemokine and cytokine synthesis, and stimulating adhesion molecule expression on vascular endothelial cells (Figure 1). Experimental studies have suggested that high mobility group box-1 (HMGB1), heat shock proteins, adenosine, extracellular RNA, matrix fragments, and Interleukin (IL)-1 α released from necrotic cardiomyocytes may stimulate the innate immune response initiating the post-infarction inflammatory cascade (7),(8),(9),(10). The relative role of specific cardiomyocyte-derived alarmins in activation of the post-infarction inflammatory response remains unknown. Alarmins released by necrotic cells activate toll-like receptor (TLR) and receptor for advanced glycation end-products (RAGE)-dependent pathways in vascular endothelial cells, immune cells and fibroblasts inducing synthesis of chemokines and cytokines (11),(12). Activation of the complement cascade is also prominent in the infarcted heart and stimulates downstream pro-inflammatory pathways (13),(14),(15). Pro-inflammatory cytokines, such as Tumor Necrosis Factor (TNF)- α , IL-1 β and IL-6, released by leukocytes, fibroblasts, endothelial cells and cardiomyocytes induce adhesion molecule and chemokine synthesis promoting recruitment of leukocytes into the infarcted area (16),(17),(18). IL-1 signaling

plays a central role in activation of the post-infarction inflammatory reaction, stimulating chemokine expression by leukocytes and fibroblasts (19), while preventing premature conversion of the abundant cardiac fibroblasts into matrix-synthetic myofibroblasts (20). The inflammasome, a molecular platform that triggers caspase-1 activation and mediates subsequent cleavage of active pro-IL-1 β into IL-1 β , plays an important role in generation of bioactive IL-1 in the infarct. *In vivo* evidence has demonstrated that activation of the inflammasome is localized in both cardiomyocytes and interstitial cells in the infarcted heart (21), (22).

Chemokine upregulation is a hallmark of the post-infarction inflammatory response and mediates recruitment of pro-inflammatory leukocyte subsets in the infarcted heart (23). CXC chemokines containing the Glu-Leu-Arg sequence (the ELR motif), such as CXCL8/IL-8, are secreted in the infarct (24), and stimulate recruitment of neutrophils, whereas CC chemokines, such as monocyte chemoattractant protein-(MCP)-1/CCL2 and CCL7, mediate recruitment of pro-inflammatory monocytes (25),(26),(27). Infiltrating leukocytes phagocytose dead cells and matrix debris and set the stage for activation of reparative cells.

Apoptosis of infiltrating neutrophils marks the end of the inflammatory phase. Professional phagocytes clear the infarct from apoptotic cells (28),(29); this process known as efferocytosis, is associated with release of anti-inflammatory cytokines, such as IL-10 and Transforming Growth Factor (TGF)- β , and may play an important role in suppression of the post-infarction inflammatory response. In addition to the role of secreted anti-inflammatory mediators, activation of endogenous intracellular signals that restrain the immune response (such as Interleukin-receptor associated kinase-M) suppresses pro-inflammatory cascades in the healing infarct (30).

Suppression of inflammatory signaling and resolution of the inflammatory infiltrate are dependent on infiltration of the infarct with inhibitory leukocyte subsets, such as anti-inflammatory monocyte subpopulations (31) and regulatory T cells (32),(33). Moreover, modulation of cardiac macrophages towards an anti-inflammatory phenotype may contribute to suppression of inflammatory cytokine and chemokine expression (34),(35). Surviving cardiomyocytes in the infarct border zone may also limit inflammation by secreting mediators with anti-inflammatory properties (29),(36). Because unrestrained activity, temporal prolongation, or spatial expansion of the inflammatory reaction result in accentuated cardiac remodeling and worse dysfunction, defective negative regulation of the post-infarction inflammatory response may be implicated in the pathogenesis of heart failure in patients surviving an acute myocardial infarction (Figure 2) (37).

As pro-inflammatory signaling is suppressed, activated myofibroblasts become the predominant cell type in the healing infarct. Infarct myofibroblasts are phenotypically modulated fibroblasts, primarily localized in the infarct border zone that express contractile proteins (such as α -smooth muscle actin and the embryonal isoform of smooth muscle myosin) (38, 39), but do not synthesize the SM1 and SM2 isoforms of smooth muscle myosin heavy chain (38). Their origin is debated: proliferation and activation of resident fibroblast populations, endothelial to mesenchymal transdifferentiation, recruitment of circulating and resident fibroblast progenitors and modulation of cardiac pericytes have been

proposed as potential sources for the abundant myofibroblasts in the infarct border zone. Recent lineage-tracing studies have suggested that resident epicardium-derived fibroblasts may be the main source of myofibroblasts in healing myocardial scars (40). Infarct myofibroblasts secrete both structural and matricellular proteins in the healing infarct. Deposition of structural matrix proteins (such as collagens) preserves the integrity and geometry of the ventricle. On the other hand, incorporation of matricellular proteins into the matrix network plays an important role in transducing cytokine and growth factor-mediated signals from the cardiac interstitium to the cellular elements, thus contributing to the plasticity of the infarct environment (41),(42),(43),(44). The renin-angiotensin-aldosterone system, TGF- β , Platelet-Derived Growth Factor (PDGF) and the mast cell-derived proteases tryptase and chymase play an important role in activating fibroblasts in the healing infarct (45),(46),(47),(48).

During the maturation phase, proliferative activity of the fibroblasts is suppressed and deposition of new matrix proteins is inhibited. Little is known regarding the pathways and inhibitory signals that terminate the fibrotic response in the infarct. During the inflammatory and proliferative phase of infarct healing, the anti-fibrotic chemokine interferon- γ -inducible protein (IP)-10/CXCL10 is upregulated in the infarcted area and inhibits fibroblast migration, preventing an overactive fibrotic response (49),(50). Whether, IP-10 or other anti-fibrotic mediators are involved in scar maturation remains unknown. As the scar matures, the collagenous matrix is cross-linked and reparative infarct myofibroblasts become quiescent, or may undergo apoptosis. However, in large infarcts, interstitial cell activation persists in the infarct border zone and in remodeling non-infarcted myocardial segments, likely reflecting regional alterations of the cardiac mechanics due to loss of contractile myocardium, and the effects of pressure and volume overload on the surviving myocardium.

TARGETING INFLAMMATION IN MYOCARDIAL INFARCTION. LESSONS FROM PAST TRANSLATIONAL FAILURES

The concept of cytotoxic inflammatory injury

Several observations fueled the notion that inflammation may extend ischemic injury following myocardial infarction. First, the idea that the inflammatory reaction triggered to clear the infarcted heart from dead cells may be inherently overactive for the delicate requirements of the myocardium. Inflammatory cascades did not evolve to protect the injured heart. In most species, cardiovascular disease is an uncommon cause of death. In humans, heart disease is a major cause of morbidity and mortality; however, myocardial infarction occurs past the reproductive age and would not be expected to impose any evolutionary pressures on the inflammatory cascade. Thus, inflammatory processes evolved as a response to traumatic injury and as a protective shield from pathogenic organisms, and may be excessive for the sterile environment of the infarcted heart (51). Second, in the infarcted myocardium, leukocytes infiltrate the viable border zone, where they may interact with surviving cardiomyocytes. Through the production of proteases, cytokines and reactive oxygen species, neutrophils and pro-inflammatory monocytes may exert cytotoxic actions on viable cardiomyocytes extending ischemic injury. In a large animal model of reperfused myocardial infarction, neutrophil depletion was reported to reduce infarct size by more than

40%, providing strong support for a leukocyte-driven mechanism responsible for extension of ischemic injury (52). The deleterious effects of neutrophils on cardiomyocytes were supported by *in vitro* evidence suggesting that activated neutrophils adhere to stimulated cardiomyocytes through integrin/intercellular adhesion molecule (ICAM)-1 interactions, and induce oxidative injury (3). It should be noted that experimental evidence did not consistently support the deleterious effects of the neutrophils in accentuating cardiomyocyte death following myocardial ischemia, as neutrophil depletion studies failed, in some cases, to reduce infarct size (53). Thus, the role of the neutrophils in extending ischemic myocardial injury has remained controversial (54).

Broad-range anti-inflammatory strategies may have catastrophic consequences on repair of the infarcted heart

If inflammation extends ischemic injury, broad inhibition of the inflammatory response with corticosteroids or non-steroidal anti-inflammatory drugs (NSAIDs) could confer significant protection to the infarcted myocardium. It should be noted that the therapeutic use of corticosteroids in acute myocardial infarction was proposed before the development of the inflammatory injury hypothesis, based on experimental studies suggesting protective actions on cell survival and cardiomyocyte metabolism (55). Clinical studies on the use of corticosteroids in patients with myocardial infarction have produced conflicting results and, in some cases, raised significant concerns regarding the safety of this approach (56),(57). *In vivo* experiments have suggested that corticosteroid treatment impairs clearance of the infarct from dead cells and may disrupt fibroblast function (58); these effects would be expected to be detrimental in repair of the infarcted heart.

NSAIDs also exert a broad range of anti-inflammatory actions mediated through inhibition of cyclooxygenase. In animal models of myocardial infarction, use of NSAIDs had both beneficial actions (on acute infarct size) and detrimental effects (on the reparative response). Treatment with ibuprofen reduced infarct size (59); the protective effects were presumed due to attenuated leukocyte infiltration (60). However, observations in a canine model of non-reperfused infarction suggested that reduced inflammation in animals treated with NSAIDs is associated with marked thinning of the scar (61). Clinical investigations showed an association between the use of NSAIDs and adverse outcome following myocardial infarction, due at least in part to increased incidence of cardiac rupture (62). Thus, non-selective inhibition of the inflammatory cascade has potentially catastrophic consequences on the reparative response. Based on this concern, current guidelines recommend against the use of broad range anti-inflammatory therapy (corticosteroids and NSAIDs) in patients with acute myocardial infarction (63).

Selective inhibition of inflammatory signaling

Advances in understanding of the biology of inflammation suggested that targeted inhibition of selected inflammatory pathways may afford protection to the infarcted heart without disturbing the reparative response. Extensive experimental evidence demonstrated that neutralization of specific inflammatory mediators (including leukocyte integrins, endothelial adhesion molecules, cytokines and chemokines) has impressive beneficial effects in large animal models of reperfused myocardial infarction, markedly reducing the size of the

infarct. Approaches targeting CD11/CD18 integrins seemed particularly promising: the bulk of experimental evidence derived from a wide range of animal models, ranging from rats to primates, showed impressive reduction in infarct size upon treatment with neutralizing antibodies (4),(64),(65),(66),(67),(68),(69). The protective actions were presumed due to reduced infiltration of the infarct with neutrophils (65) and to attenuated neutrophil-induced cardiomyocyte injury. Unfortunately, the beneficial effects of anti-integrin targeting in animal models could not be reproduced in clinical studies. In three small clinical trials, anti-integrin approaches failed to reduce the size of the infarct in patients with myocardial infarction (70),(5),(71). Approaches targeting the complement system, an upstream activator of the innate immune response, were equally disappointing. In the Assessment of Pexelizumab in Acute Myocardial infarction (APEX-AMI) clinical trial, 5745 patients with STEMI received the anti-C5 antibody pexelizumab as an intravenous bolus prior to percutaneous intervention followed by continuous infusion over the next 24h. Pexelizumab administration did not affect 30-day mortality and the composite endpoint of death, cardiogenic shock and congestive heart failure (72). Moreover, administration of the P-selectin inhibitor inclacumab in patients with acute coronary syndromes reduced the release of enzymes associated with cardiomyocyte necrosis, but was associated with trends towards worse clinical outcome (73),(74). Considering the great enthusiasm generated by the impressive results of the animal model studies, what are the possible causes of these translational failures?

The anti-inflammatory strategies used in clinical trials may have been suboptimal

Translation of a therapeutic approach from the animal model to the clinical context is not dependent solely on implementation of a sound pathophysiologic concept, but also requires careful planning of the strategy to maximize the likelihood of success. Selection of an appropriate molecular target, identification of an effective therapeutic agent, correct timing and optimal delivery of the agent in the infarcted area are all crucial for successful translation of the approach. Ultimately, understanding the basis for failure of anti-integrin and complement inhibition strategies requires assessment of the effectiveness of these strategies in inhibiting the myocardial inflammatory response. Unfortunately, such data are lacking. In a subgroup of patients enrolled in the APEX-AMI trial, assessment of circulating biomarkers suggested that pexelizumab was not successful in suppressing most indicators of inflammation (75). Although use of circulating biomarkers as a window to the myocardial inflammatory response is inherently problematic, ineffective suppression of inflammation may provide an explanation for the negative trial. Moreover, experimental studies suggested that the protective actions of integrin inhibition on the infarcted myocardium are variable and may be dependent on the specific characteristics of the antibody used (76). Protective actions could only partially be predicted by the extent of inhibition of neutrophil infiltration (76). Future strategies targeting post-infarction inflammation need to include systematic assessment of the effectiveness of the approach in attenuating several distinct aspects of the inflammatory response through a combination of *in vivo* imaging approaches and analysis of circulating biomarkers.

Inflammation may not extend cardiomyocyte death following myocardial infarction

An alternative explanation of the translational failure of anti-inflammatory approaches may be that, despite the abundant evidence in animal models of myocardial infarction, acute inflammatory cardiomyocyte injury may be of limited significance in human myocardial infarction. Considering the intense inflammatory reaction triggered following infarction in both human patients with STEMI and in animal models of infarction, the disconnect between the human clinical studies and the animal model investigations is unlikely to be explained only by species-specific effects, but may reflect the limitations of animal model experiments. First, enthusiasm regarding the effectiveness of anti-inflammatory strategies in myocardial infarction derived by early inhibition experiments in large animal models may have been premature and somewhat misguided. In contrast to the impressive protective effects of antibody neutralization in large animal models, a more recent body of work in genetically targeted mice has challenged the notion that post-infarction inflammation extends ischemic cardiomyocyte injury. Mice lacking P-selectin and ICAM-1 (77), animals with defective IL-1 signaling (19), and MCP-1 knockouts (25) had no significant reduction in infarct size, despite a marked attenuation of the post-infarction inflammatory response.

Second, animal models cannot recapitulate the pathophysiologic complexity and heterogeneity of the clinical context. Outcome in human patients with myocardial infarction is affected by a wide range of factors. Differences in gender, age, genetic profile, the pattern of coronary disease, comorbid conditions (including diabetes, obesity, hypertension and hyperlipidemia), and treatment with various pharmacologic agents have profound effects on the pathophysiologic response to myocardial infarction. In contrast, in a well-designed animal study, the goal is to eliminate variability in order to test a specific hypothesis. Experimental animals are healthy, matched for gender and age and have an identical genetic profile so that the consequences of a very specific genetic or pharmacologic intervention can be studied. For this reason, animal model studies are optimally used to gain pathophysiologic insights and not to predict effectiveness of a therapeutic approach. Experiments in senescent animals illustrate the impact of age on the inflammatory and reparative response following myocardial infarction. Senescent mice exhibited significantly suppressed (and somewhat prolonged) inflammatory activation following myocardial infarction, associated with defective activation of growth factor signaling and impaired collagen deposition (78). Considering that the conclusions on the effectiveness of anti-integrin approaches in myocardial infarction were derived from experimental studies performed in young mammals (known to exhibit very robust inflammatory reactions), the translational failure may reflect, at least in part, the suppressed inflammatory activation in aged human populations presenting with myocardial infarction.

USE OF TARGETED ANTI-INFLAMMATORY STRATEGIES TO IMPROVE REPAIR AND TO REDUCE ADVERSE POST-INFARCTION REMODELING

The failures of anti-inflammatory strategies in myocardial infarction may reflect the limited role of inflammatory cardiomyocyte injury during the early stages of infarction. However, inflammation is critically involved in repair and remodeling of the infarcted heart. Inflammatory pathways have been implicated in recruitment of progenitor cells that may

play a crucial role in infarct angiogenesis and cardiac repair (79). Chemokines (such as stromal cell derived factor (SDF)-1/CXCL12 and MCP-3) mediate homing of progenitor cell subpopulations in the infarcted myocardium (80),(81). Growth factors, such as stem cell factor, hepatocyte growth factor and vascular endothelial growth factor are also upregulated in the infarcted myocardium (82),(83) and may be involved in recruitment and activation of stem cell subsets. On the other hand, prolonged or expanded pro-inflammatory signaling may accentuate adverse remodeling by activating proteases, transducing pro-apoptotic cascades in cardiomyocytes, and promoting matrix degradation. Extensive experimental work suggests that overactive, temporally prolonged (30), or spatially expanded (42) inflammation may cause dilative remodeling following myocardial infarction. Highly-selective approaches to inhibit inflammation-driven protease activation and to promote recruitment of reparative cells may exert beneficial actions on the infarcted heart by stimulating repair, by reducing adverse remodeling and by preventing the development of post-infarction heart failure. Several inflammatory mediators have shown promise as therapeutic targets.

THE CHEMOKINES

The chemokines are a large family of small (8–14 kDa) chemotactic cytokines with a crucial role in regulating immune function and inflammatory responses (84). On a structural basis, chemokines are classified into 4 subfamilies according to the presence of one or more aminoacids between the N-terminal cysteine residues region (CC, CXC, CX₃C and XC subfamilies). Most chemokines belong to CC and CXC subfamilies. The structural differences between the chemokines have important functional implications. CC chemokines are potent mononuclear cell chemoattractants, whereas CXC chemokines that contain the ELR motif preceding the first aminoterminal cysteine mediate chemotactic recruitment of neutrophils (85).

Experimental studies in animal models of myocardial infarction demonstrated that several members of the chemokine family are rapidly and consistently upregulated in the infarcted heart and may play an important role in regulation of the post-infarction inflammatory response (23). Certain members of the family, including MCP-1/CCL2 and SDF-1/CXCL12 have shown promise as potential therapeutic targets. The findings of experimental studies targeting chemokine family members in myocardial infarction are summarized in Table 1.

CC chemokines

The CC chemokine MCP-1/CCL2 is rapidly upregulated in the infarcted myocardium and is predominantly expressed in vascular endothelial cells (86). Genetic disruption of MCP-1, or its receptor CCR2, attenuated adverse remodeling following myocardial infarction, inhibiting recruitment of pro-inflammatory monocytes and decreasing cytokine expression in the infarct (25),(87). In a model of non-reperfused infarction, anti-MCP-1 therapy exerted beneficial actions on the infarcted ventricle, reducing mortality, attenuating chamber dilation, and improving systolic function (88). Thus, experimental observations suggest that MCP-1/CCL2 may be a promising therapeutic target following myocardial infarction. However, a word of caution should be raised by observations suggesting that genetic disruption of MCP-1 results in impaired phagocytosis of dead cardiomyocytes, and delayed

formation of granulation tissue (89),(90). These defects may reflect decreased recruitment of monocytes and impaired macrophage maturation (25). The clinical consequences of impaired clearance of the infarct from dead cells cannot be predicted. Persistence of non-phagocytosed necrotic cardiomyocytes in the infarcted region may have adverse consequences on cardiac function and may be associated with dysrhythmic events in human patients with infarction.

A recent study suggested that inhibition of the CC chemokine CCL5/RANTES (regulated on activation, normal T cell expressed and secreted) may exert cardioprotective actions (91). RANTES neutralization decreased the size of the infarct and improved cardiac function 3 weeks after infarction, reducing expression of matrix metalloproteinase (MMP)-9. Whether RANTES mediates recruitment a specific subset of pro-inflammatory mononuclear cells, or promotes a matrix-degrading phenotype in leukocytes infiltrating the infarct remains unknown. Although MCP-1 and RANTES may be promising therapeutic targets, broad inhibition of CC chemokines may have detrimental actions. In a mouse model of reperfused infarction, genetic disruption of the CC chemokine receptor 5 (CCR5) was associated with accentuated dilative remodeling, presumed due to impaired recruitment of inhibitory monocyte subsets and of regulatory T cells (Tregs) (92). Thus, certain chemokine-chemokine receptor interactions may be important in repression and resolution of post-infarction inflammation through recruitment of anti-inflammatory leukocyte subpopulations.

ELR+ CXC chemokines

Although ELR+ CXC chemokines are markedly upregulated in the healing infarct and mediate neutrophil infiltration (24),(93), their potential therapeutic role has not been systematically studied. Early experiments suggested that IL-8/CXCL8 inhibition in rabbits undergoing ischemia/reperfusion protocols reduced infarct size; surprisingly the beneficial effects were not associated with attenuated neutrophil infiltration (94). Enthusiasm about neutrophil chemoattractant chemokines as therapeutic targets in myocardial infarction was dampened by the translational failures of anti-integrin approaches. However, a recent investigation in a mouse model of myocardial ischemia/reperfusion suggested that treatment with evasin-3, a protein that binds and neutralizes neutrophil chemoattractant CXC chemokines, reduces infarct size by attenuating leukocyte recruitment (95).

SDF-1

The ELR-negative CXC chemokine SDF-1/CXCL12 has potent angiogenic properties, activates pro-survival pathways in cardiomyocytes, and enhances the regenerative capacity of progenitor cells (96). Thus, it is not surprising that treatment with SDF-1 has been considered as a therapeutic strategy for patients with myocardial infarction. The effectiveness of SDF-1 therapy is supported by extensive experimental evidence. Treatment with SDF-1 reduced infarct size, promoting cardiomyocyte survival and accentuating angiogenesis in experimental models of myocardial infarction (97),(98). Several protective mechanisms have been suggested. First, SDF-1-induced angiogenesis in the infarct and in the border zone may improve the quality of the scar attenuating systolic dysfunction. Second, SDF-1 may exert direct anti-apoptotic actions on cardiomyocytes or may promote chemotaxis of a CXCR4+ myeloid cell subset that secretes cytoprotective mediators (99).

Third, SDF-1 may promote repair and regeneration through recruitment of progenitor cells. Extensive evidence suggests that SDF-1 is critically implicated in mobilization and trafficking of hematopoietic stem cells (100), and mediates homing of endothelial progenitor cells in ischemic tissues (80).

Recent studies have tested novel synthetic analogs of SDF-1 in both rodent and large animal models of myocardial infarction. Injection of a biomimetic hydrogel containing a combination of SDF-1 and angiogenic peptides reduced the size of the infarct and improved angiogenesis in a rat model of myocardial infarction (101). In both rat and ovine models, administration of a bioengineered SDF-1 analog in the infarct border zone induced chemotaxis of endothelial progenitor cells and preserved ventricular function, improving left ventricular mechanics (102, 103). Although these experimental studies are promising, it should be emphasized that SDF-1 may also exert pro-inflammatory actions. The pleiotropic, cell-specific and context-dependent actions of SDF-1 are highlighted by the conflicting observations reported in SDF-1 antagonism studies. Some investigations showed that SDF-1 inhibition accentuated dysfunction (supporting the protective actions of the chemokine revealed by the gain-of-function approaches) (104), whereas other studies suggested beneficial effects (105), (106).

Fractalkine/CX3CL1

The CX3C chemokine fractalkine exists in membrane-anchored and soluble forms and may regulate trafficking and function of immune cells. Its role in myocardial infarction remains poorly understood. In a mouse model of myocardial infarction, fractalkine expression was markedly increased in the viable remodeling myocardium (107); fractalkine inhibition delayed progression of chamber dilation, attenuating pro-inflammatory and matrix-degrading pathways (108). Thus, fractalkine may hold promise as a therapeutic target to attenuate adverse post-infarction remodeling.

THE CYTOKINES

Targeting the IL-1 system

The prototypical pro-inflammatory cytokine IL-1 plays a crucial role in stimulation of the post-infarction inflammatory response and is involved in the pathogenesis of cardiac remodeling. Thus, targeting the IL-1 signaling cascade may be a promising therapeutic target for patients with myocardial infarction. In experimental models of myocardial infarction, IL-1 α is released by dying cardiomyocytes (9), whereas IL-1 β synthesis is markedly upregulated following infarction (16), and is predominantly localized in leukocytes and vascular cells (109). Both genetic and pharmacologic strategies disrupting IL-1 signaling have been shown to protect the infarcted heart from adverse remodeling. Genetic loss of the type 1 IL-1 receptor (IL-1R1) attenuates dilation of the infarcted heart, reducing adverse remodeling (19). Pharmacologic approaches targeting the IL-1 system have also been tested in models of myocardial infarction. The availability of safe and effective pharmacologic strategies to inhibit IL-1 signaling in human patients provides promising therapeutic tools. Anakinra, a non-glycosylated recombinant form of interleukin-1 receptor antagonist (IL-1Ra), binds to the type 1 IL-1 receptor without activating a signaling

response, thus functioning as a competitive inhibitor for both IL-1 α and IL-1 β . Anakinra has been approved for treatment of patients with rheumatoid arthritis who fail to respond to disease modifying agents. Anti-IL-1 β antibodies (such as canakinumab) provide more selective options, specifically targeting IL- β signaling and have been approved for treatment of autoinflammatory diseases and certain forms of inflammatory arthritides. Treatment with anakinra reduced chamber dilation in a rat model of myocardial infarction (110); administration of an anti-IL-1 β antibody in a mouse model of non-reperfused infarction also exerted protective actions (111). The protective effects of IL-1 blockade may be only in part mediated through reduction in the size of the infarct. Attenuation of IL-1-driven protease activation in the cardiac interstitium may be implicated in protection from adverse remodeling.

Small clinical trials have tested the effectiveness of IL-1 inhibition in patients with myocardial infarction. Pilot studies have suggested that anakinra can be safely administered as a 2-week course in patients with STEMI and may attenuate adverse remodeling, while protecting from the development of post-infarction heart failure (112),(113),(114). Because IL-1 has been implicated in the pathogenesis of the vulnerable plaque, a large clinical trial is currently underway to examine the effectiveness of IL-1 β antibody inhibition in prevention of cardiovascular events in post-myocardial infarction patients with accentuated systemic inflammatory responses (115).

Although both experimental and early clinical findings suggest that IL-1 inhibition may hold promise for treatment of patients with myocardial infarction, a word of caution should be raised regarding cytokine inhibition in patients with heart disease. Cytokines are notoriously pleiotropic and multifunctional and are known to exert a wide range of context-dependent actions on all cell types involved in cardiac injury and repair. In the infarcted and remodeling heart, cytokines may exert both beneficial and detrimental effects; thus, prediction of the consequences of cytokine inhibition in the clinical context is challenging. The failure of anti-TNF strategies in patients with heart failure highlights the challenges in implementation of targeted anti-cytokine approaches in patients with cardiovascular disease. However, it should be noted that, in contrast to TNF- α , IL-1 is not known to exert protective actions on cardiomyocytes. Studies in patients with rheumatoid arthritis suggest protective actions of anakinra on myocardial function (116),(117).

Targeting the TGF- β cascade

Members of the TGF- β family are critically involved in regulation of inflammation and fibrosis in a wide range of pathophysiologic conditions (118). It has been suggested that, following myocardial infarction, TGF- β may serve as the “master switch” that de-activates inflammatory macrophages, while promoting fibrosis (119). Clearly, this concept represents an oversimplification. TGF- β modulates phenotype and function of all cell types involved in cardiac repair, activating both Smad-dependent and Smad-independent signaling (120), (121). The effects of TGF- β inhibition may be dependent on timing: early neutralization of TGF- β may prolong inflammation and enhance the incidence of cardiac rupture; late suppression may attenuate pro-fibrotic signaling improving diastolic function. Because TGF- β plays an important role in preservation of cardiovascular homeostasis, targeting the

TGF- β system in heart failure may carry significant risks, promoting aneurysmal rupture in vulnerable patients (122),(123),(124). Dissection of downstream signaling effectors and identification of specific TGF- β -activated pathways associated with post-infarction remodeling and dysfunction are needed to design safe and effective therapy for patients with myocardial infarction.

Do inflammatory mediators transduce cytoprotective and regenerative signals?

Identification of cytoprotective and regenerative actions of leukocyte subsets contributes an additional layer of complexity to the effects of inflammatory cells on the infarcted heart (125). Experiments in models of neonatal cardiac injury suggested that subpopulations of macrophages with unique phenotypic profiles may promote cardiomyocyte proliferation activating a regenerative program (126),(127). The signals that may drive macrophages towards a regenerative phenotype remain unknown. In adult mice, a recent investigation identified myeloid-derived growth factor (MYDGF), as a novel mediator released by a subset of CXCR4-expressing macrophages, that protects cardiomyocytes from ischemic death (99). These findings suggest that inflammatory cells recruited in the infarcted heart not only debride the wound and contribute to scar formation, but may also exert direct protective actions on cardiomyocytes, and may hold the key to cardiac regeneration.

CHALLENGES IN IMPLEMENTATION OF ANTI-INFLAMMATORY STRATEGIES IN PATIENTS WITH ACUTE MYOCARDIAL INFARCTION

Inflammatory mediators exert a wide range of diverse functions on the infarcted heart. The involvement of inflammatory cells and their secretory products in both injurious and protective effects complicates our efforts to design effective therapy for patients with myocardial infarction. Experimental studies in animal models of myocardial infarction have identified several promising therapeutic targets. However, the failures of the anti-integrin and complement inhibition approaches, despite strong experimental evidence supporting their effectiveness, have generated skepticism regarding our ability to translate promising animal findings into clinical applications.

It should be emphasized that investigations using animal models are essential for dissection of the pathophysiologic mechanisms, but have limited value in predicting success of a therapeutic intervention in the clinical context. As discussed in the previous section, the complexities of the clinical context cannot be simulated in an experimental model. In view of these challenges, how can we optimally use insights from animal models to design effective strategies targeting the inflammatory response in human patients with myocardial infarction? Considering the pathophysiologic heterogeneity of STEMI patients that may explain differences in susceptibility to adverse remodeling, there is a need to identify patients with overactive post-infarction inflammatory responses that may benefit from targeted anti-inflammatory strategies (37),(128). Certain patient subpopulations, such as diabetics and the elderly, may exhibit dysregulated inflammatory reactions following myocardial infarction that may be responsible for accentuated remodeling and worse dysfunction. For example, diabetics have an increased incidence of heart failure following myocardial infarction despite a smaller infarct size and comparable systolic dysfunction

(129). Development of post-infarction heart failure in diabetics is associated with diastolic dysfunction (130). In mice, diabetes and obesity are associated with cardiac fibrosis, hypertrophy and overactive myocardial TGF- β /Smad signaling (124),(131),(43). A link between diabetes-associated TGF- β activation and fibrotic remodeling of the infarcted heart is plausible; thus, in these patients targeting the TGF- β system may be a promising therapeutic strategy. On the other hand, persistently increased circulating levels of pro-inflammatory mediators (such as MCP-1/CCL2) are associated with worse prognosis in patients with acute coronary syndromes. Targeted inhibition of inflammation may be effective in patients with defective negative regulation of pro-inflammatory signaling that may exhibit evidence of prolonged inflammatory activation Biomarkers and imaging approaches may be used to obtain information on activation of inflammatory pathways in each patient, in order to personalize treatment options.

CONCLUDING REMARKS

Activation of inflammatory cascades in the infarcted heart stimulates a range of cellular responses that clear the wound from dead cells and promote repair, but may also extend injury and cause adverse remodeling of the ventricle. Progress in understanding the cellular effectors and molecular signals regulating post-infarction inflammation has not yet translated into effective therapy. Future research should dissect protective and detrimental inflammatory pathways in animal models, while expanding our understanding of the human pathophysiology. Identification and validation of biomarkers that may reflect specific perturbations of the inflammatory response in human patients may provide much-needed pathophysiologic guidance for implementation of personalized treatment approaches.

ACKNOWLEDGMENTS

Dr Frangogiannis' laboratory is supported by NIH grants R01 HL76246 and R01 HL85440. Ilaria Russo is supported by training grants from the Fondazione Cassa di Risparmio di Lucca and the Fondazione Banca del Monte di Lucca.

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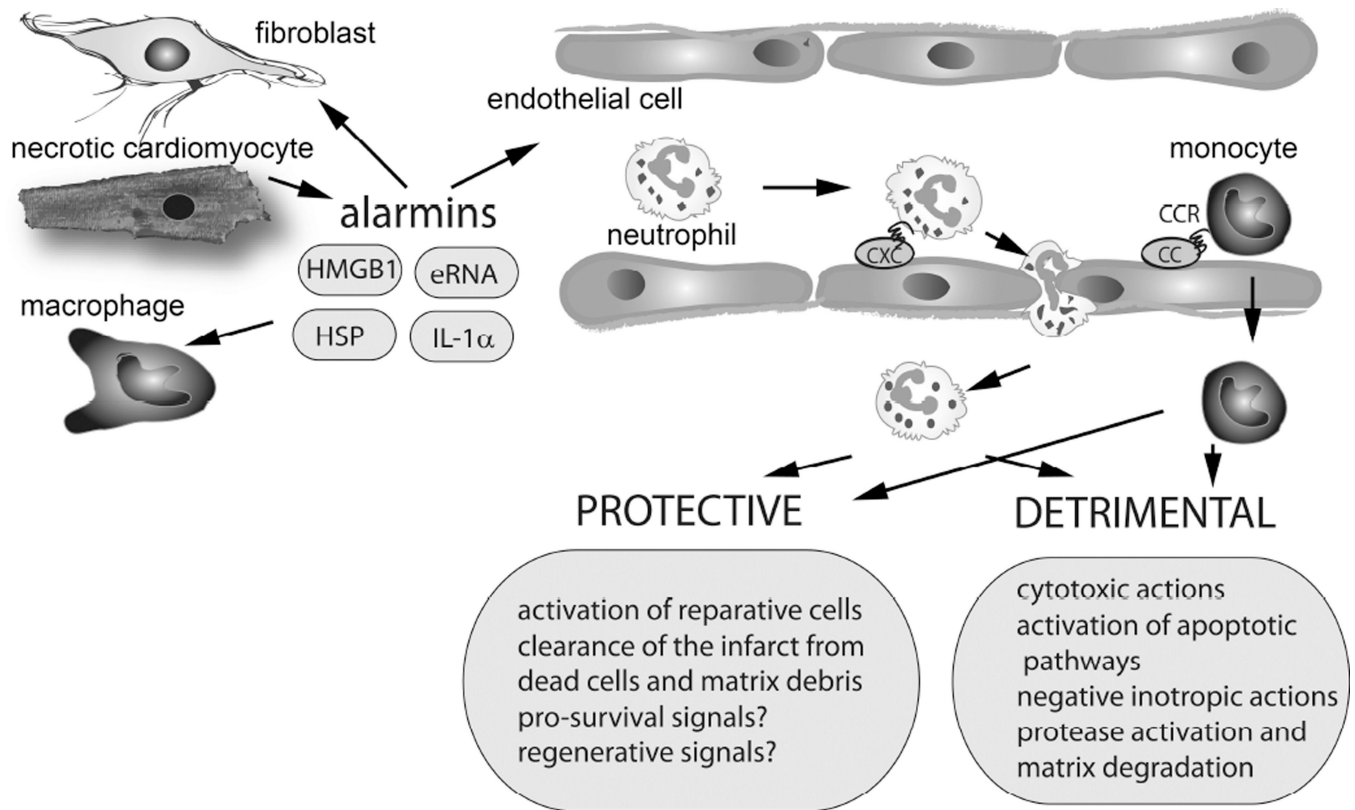


Figure 1.

The post-infarction inflammatory reaction exerts both protective and detrimental actions on the infarcted heart. Necrotic cardiomyocytes release danger signals that activate innate immune responses leading to induction of cytokines and chemokines. CXC and CC Chemokines bind to glycosaminoglycans on the endothelial surface and mediate recruitment of neutrophils and mononuclear cells in the infarcted area. Infiltrating leukocytes exert a wide range of protective and detrimental actions on the infarcted heart. It has been proposed that infiltrating neutrophils may extend ischemic injury exerting cytotoxic, and pro-apoptotic actions on cardiomyocytes. Moreover, leukocyte-derived secretory products may have negative inotropic effects and proteases may increase matrix degradation causing adverse remodeling of the infarcted heart. Other actions of leukocytes may be protective. Thus, leukocyte subsets may phagocytose dead cells and matrix debris and may promote repair by activating endothelial cells and fibroblasts. Recent studies have suggested that subsets of myeloid cells may also exert pro-survival and regenerative actions on cardiomyocytes.

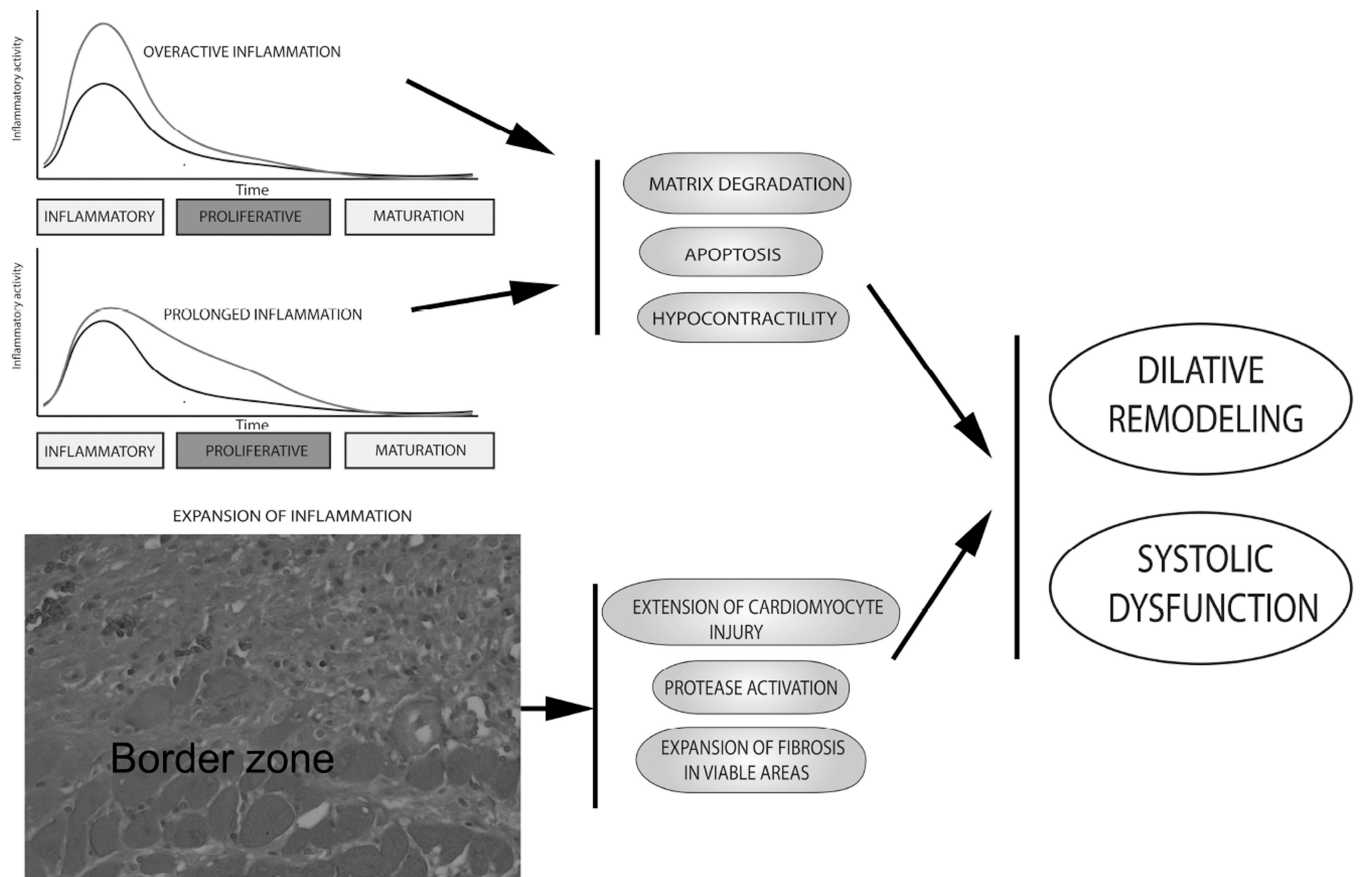


Figure 2.

The consequences of excessive, prolonged and expanded inflammatory responses on the remodeling infarcted heart. Overactive or prolonged inflammation may accentuate matrix degradation by stimulating protease activation. Increased cytokine expression may increase cardiomyocyte apoptosis and suppress contractility. Defects in spatial containment of the post-infarction inflammatory response may result in extension of inflammatory injury in the viable myocardial areas leading to expansion of fibrosis. Impaired negative regulation of post-infarction inflammation may cause adverse dilative remodeling and accentuate systolic dysfunction.

Table 1
Experimental studies targeting the chemokines in myocardial infarction: functional outcome and histopathologic consequences.

Target	Approach	Animal model	Timing of Intervention/Route	Outcome	Histopathologic/Molecular Consequences
MCP-1/CCL2(88)	Gene transfection	Mouse (PO)	3d before and 14d after PO/IM	Improved survival, attenuated LV dilation, reduced contractile dysfunction, comparable infarct size	Attenuated interstitial fibrosis, reduced macrophage recruitment, attenuated myocardial expression of TNF- α and TGF- β
MCP-1/CCL2(132)	Competitive inhibition	Mouse (I/R)	10' after I, 2h after R and daily up to 7 days/IP	Preserved LV function, reduced infarct size	Reduced monocyte infiltration, reduced collagen and myofibroblast content in the infarcted area
MCP-1/CCL2 (18)	Ab neutralization	Rat (I/R)	IV	Reduced infarct size	Decreased macrophage infiltration, reduced ICAM-1 mRNA expression
RANTES/CCL5(91)	Ab neutralization	Mouse (PO)	5' and 3d after PO/IV	Smaller infarct size, improved survival and LV function	Reduced neutrophil and macrophage infiltration, reduced cardiac MMP-9, lower collagen content
Fractalkine/CX3CL1(108)	Ab neutralization	Mouse (PO)	Daily from 7 to 14 days after PO/IP	Delayed progression of LV enlargement and improved LV function, smaller infarct size	Reduced ICAM-1 in vascular endothelium
IL-8/CXCL8(94)	Ab neutralization	Rabbit (I/R)	15' after R/IV	Reduced infarct size	Comparable neutrophil infiltration
SDF-1/CXCR4(103)	Hydrogel delivery system	Rat (PO)	At time of PO/into the borderzone	Preserved LV geometry, improved systolic function	Improved angiogenesis
SDF-1/CXCR4(102)	Hydrogel delivery system	Sheep (PO)	At time of PO/into the borderzone	Preserved systolic LV function, reduced infarct size	Reduced MMP-2 in the infarct border zone, elevated levels of TIMP-1 and elastin in the infarct, increased capillary and arteriolar density
SDF-1/CXCR4(133)	Hydrogel delivery system	Mouse (PO)	At time of PO/into the borderzone	Preserved cardiac function	Suppressed neutrophil infiltration, increased angiogenesis,

Target	Approach	Animal model	Timing of Intervention/Route	Outcome	Histopathologic/Molecular Consequences
SDF-1/CXCR4(134)	Administration of a small molecule CXCR4 antagonist	Mouse (I/R)	At reperfusion/SC	Smaller infarct size, improved LV function	reduced apoptosis Increased capillary density, mobilization of endothelial progenitors
SDF-1/CXCR4(105)	Administration of a small molecule CXCR4 antagonist	Rat (PO)	IP injection or oral administration 24h post-MI, for 6 days	Smaller scar size, improved LV systolic function	Attenuated atrial natriuretic peptide expression in remodelling myocardium
SDF-1/CXCR4(106)	Administration of a small molecule CXCR4 antagonist	Mouse (PO)	Acute treatment: single-dose, SC, 1h after infarction Chronic treatment: infusion over 2 weeks.	Acute: improved LV function and survival. Chronic: increased adverse remodeling	Acute: enhanced angiogenesis, attenuated inflammation, reduced incorporation of endothelial progenitor cells. Chronic: reduced capillary density

PO, permanent occlusion; I/R, ischemia/reperfusion; IM, intramuscularly; IP, intraperitoneally; SC, subcutaneously; LV, left ventricular.