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Cardiac Fibroblast in Development and Wound Healing

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Introduction

Cardiac fibroblasts are cells of mesenchymal nature that reside within the cardiac interstitium[1, 2]. They comprise the majority of the cells in the adult rodent and human heart and express proteins that maintain homeostasis of the extracellular matrix (ECM)[3, 4]. The interconnected network of cardiac fibroblasts forms the fibrous skeleton of the heart and serves as a scaffold for supporting all cardiac cell types. Aside from maintenance of the extracellular matrix, cardiac fibroblasts express a rich array of growth factors and cytokines and communicate with adjoining myocytes to facilitate electro-mechanical transduction[1, 5]. The fibroblast also lies at the heart of most cardiac pathologies[6]. Common congenital and adult cardiac pathologies are characterized by a loss of cardiac muscle and as the mammalian heart is unable to robustly regenerate, lost cardiac muscle is replaced by fibrosis. Fibrosis induces adverse changes in cardiac geometry and function that leads to progressive chamber enlargement, hypertrophy of viable myocytes, increasing wall tension and ultimately congestive cardiac failure[7–9]. Despite the enormous pathophysiological importance of fibrosis in cardiac diseases, the cardiac fibroblast remains an ill-defined cell

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Disclosures

None

and very few interventions effectively target the cardiac fibroblast and fibrosis[10, 11]. This review summarizes the role of the cardiac fibroblast in cardiac development and repair to emphasize the emerging but central role of this cell in regulating cardiac function in health and disease.

The Cardiac Fibroblast In Development

Developmental origin of cardiac fibroblasts

Cardiac fibroblasts are thought to be predominantly derived from the epicardium (Figure 1) [1]. Villous like projections protrude from the venous pole of the developing heart to form the proepicardium[12, 13]. Cells from the pro-epicardium detach and attach on the beating ventricular surface to form the epicardium[14]. Subsequently epithelial cells of the epicardium undergo epithelial-mesenchymal-transition (EMT) to form mesenchymal cells that invade the developing myocardium[15]. A subset of these mesenchymal cells after EMT acquire migratory properties and invade the developing myo-fascial planes to occupy an interstitial position in between cardiac myocytes to become resident cardiac fibroblasts[16, 17].

Cardiac fibroblasts are observed in the developing murine heart by E12.5 days post fertilization (dpf) and their numbers progressively increase throughout development[3]. Using flow cytometry, Banerjee et al estimated that cardiac fibroblasts comprise approximately 14% of all murine heart cells at E18.5 dpf. Fibroblast numbers progressively increase in the heart in post natal life comprising 27% of the total number of cells in the adult murine heart (Table 1)[3]. In the rat heart, cardiac fibroblasts constitute approximately 30% of the total number of heart cells on the first day of life and by the 15th day of post-natal life, they comprise approximately 2/3 of the total numbers of cells in the rat heart (Table 2)[3, 4]. In humans, non-myocyte cells comprise approximately 70% of the total number of cardiac cell types[18, 19]. The higher number of fibroblasts in rat and human hearts may be related to a larger heart size, greater wall tension and consequently a need for greater production of ECM[3].

In contrast to cardiac fibroblasts, valvular fibroblasts are thought to be derived from the endothelium overlying the region of the cardiac cushions (site of atrio-ventricular valve formation)[20–22]. The endothelium overlying the valve leaflets undergoes endothelial-mesenchymal-transition (EndMT) to generate cardiac fibroblasts that invade the valvular mesenchyme and contribute to the collagenous structure of the valve (Figure 1).

Signals regulating cardiac fibroblast formation during development

As cardiac fibroblasts are formed via EMT, signals regulating EMT affect fate specification of epicardial cells to form fibroblasts. EMT during development is tightly regulated and complex to analyze as different molecules exhibit species specific differences in the regulation of EMT. Fibroblast growth factors (FGFs) are thought to promote EMT. Members of the transforming growth factor β (TGF β) superfamily inhibit EMT in the developing avian heart to regulate the fraction of epicardial cells adopting a mesenchymal phenotype[23]. Notch signaling promotes EndMT and formation of valvular fibroblasts and disruption of Notch signaling in developing cardiac valves leads to formation of atrophic

valves[24]. Once the epicardial cells undergo EMT, platelet derived growth factor (PDGF) and TGF β are thought to promote differentiation into a myofibroblast or cardiac fibroblast phenotype. Platelet derived growth factor receptor (PDGFR α) in particular, is thought to be required for epicardial derived cells to adopt a cardiac fibroblast fate[25]. In addition to secreted molecules, several transcription factors such as T box transcription factor 18 (Tbx18), Wilms tumor 1 (Wt-1) and members of the snail family also regulate EMT during development[20]. Disruption of Wt-1 in the epicardium increases Snail1 levels that subsequently inhibits EMT and increases expression of epithelial markers[26].

Function of cardiac fibroblast during heart development

In contrast to the well described functions of the cardiac fibroblast in the adult heart, less is known about the role of the cardiac fibroblast in heart development. The number of cardiac fibroblasts progressively increases from E12.5 dpf and the increasing number of fibroblasts form a fibrous skeleton that likely provides structural support for the growing number of cardiac myocytes. Embryonic cardiac fibroblasts compared to adult cardiac fibroblasts express more growth factors and secrete in greater abundance extracellular matrix proteins such as periostin, fibronectin and tenascin C[27]. In co-culture experiments it was observed that fibronectin and type 3 collagen secreted by the embryonic cardiac fibroblast induced proliferation of embryonic cardiac myocytes via a β 1 integrin receptor dependent mechanism. These observations suggest that cardiac fibroblasts can regulate cardiomyocyte proliferation during development (Figure 2)[27]. Cardiac fibroblasts express angiogenic cytokines such as vascular endothelial growth factor (VEGF) and FGF. During cardiac development, FGF and VEGF drive mesenchymal cells originating from epicardial EMT to adopt an endothelial lineage[28]. Cardiac fibroblasts also express anti-angiogenic cytokines such as connective tissue growth factor (CTGF)[1]. More rigorous studies are needed to address the role of cardiac fibroblast in the regulation of angiogenesis in the embryonic heart.

The Cardiac Fibroblast In Wound Healing

Origins of cardiac fibroblasts after cardiac injury

Fibroblasts in the injured heart are thought to have diverse origins (Figure 1). Resident cardiac fibroblasts at the site of injury proliferate and are thought to be the predominant pool of cardiac fibroblasts contributing to cardiac fibrosis after injury, although rigorous fate mapping studies have not been done to confirm this[1]. Early studies using tritiated thymidine demonstrated a small pool of proliferating perivascular cardiac fibroblasts after cardiac injury[18, 29]. Fibrosis around the perivascular region is observed in the early stages of many cardiac pathologic conditions and pericytes and perivascular cells represent another population of cells contributing to fibrosis in the injured heart (Figure 1)[30, 31].

Bone marrow derived cells also contribute to cardiac fibrosis[32]. Several studies have demonstrated that between 3 and 24% of myofibroblasts in the injured region are of bone marrow origin[33, 34]. These circulating fibroblast precursors have been termed fibrocytes and express hematopoietic (CD45), monocytic (CD11b) and progenitor markers (CD34). The ability of bone marrow derived cells to contribute to cardiac fibrosis appears to be

physiologically important as inhibition of fibrocyte recruitment diminished fibrosis and had salutary effects on remodeling[33].

We and others have demonstrated that a subset of epicardial cells are activated after acute cardiac injury. Epicardial derived cells undergo EMT to generate cardiac fibroblasts, thus recapitulating a developmental program of epicardial EMT (Figure 1)[35, 36]. Following EMT, epicardial derived cardiac fibroblasts reside in the subepicardial space, express collagen and contribute to a pro-fibrotic repair response. Inhibition of epicardial EMT decreased the number of epicardial derived cardiac fibroblasts but led to cardiac chamber dilatation and worsening ejection fraction. These observations suggest that epicardial derived cardiac fibroblasts play an important physiological role in the cardiac repair response[35].

Endothelial cells undergo endothelial-mesenchymal transition (EndMT) and have been reported to contribute to 30% of the cardiac fibroblasts in a murine model of pressure overload injury[37], but the degree to which endothelial cells contribute to fibrosis in the acutely injured heart is less certain. The physiological significance of fibroblast diversity in the injured heart is unclear and whether different subsets of cardiac fibroblasts of diverse origins exert disparate effects to co-ordinate a repair process remains an unanswered question.

Function of the cardiac fibroblast in wound healing

The cardiac fibroblast plays a central role in wound healing after myocardial injury and affects various aspects of the wound healing response from deposition of extracellular matrix proteins to wound angiogenesis and scar maturation (Figure 2). Following acute myocardial infarction, cardiac fibroblasts in the heart become activated and rapidly proliferate. In rodent hearts, peak cardiac fibroblast numbers are achieved within 7–14 days after permanent ligation of the left anterior descending coronary artery[38] and within 3 days of ischemia-reperfusion injury[39]. Four weeks after ischemia-reperfusion injury, the number of cardiac fibroblasts decreases by 90% compared to its peak, demonstrating the dynamic nature of the fibroblast population in the region of injury. Activated fibroblasts at the site of injury express contractile proteins such as α -smooth muscle actin (myofibroblasts) and secrete ECM proteins (mainly collagens). This early fibrotic repair response is critical for maintenance of cardiac structural integrity and performance after cardiac injury. Disruption of cardiac fibroblast activation early after injury led to impaired wound healing and worsening cardiac performance[35].

In the later phases of wound healing (days to weeks), collagen fibers at the site of injury undergo cross-linking which increases the tensile strength of the scar[40]. The scar subsequently contracts undergoing a reduction in surface area and myofibroblasts expressing contractile proteins are thought to contribute to scar contraction (Figure 2). Scar contraction and thinning lead to adverse changes in ventricular chamber geometry and compliance. The hemodynamic burden on the remaining viable myocytes is increased, and over time, this leads to the development of congestive heart failure. In this section, we cover the various roles of the cardiac fibroblast in early and late wound healing, describe the molecular

activation of cardiac fibroblasts after cardiac injury and finally discuss how fibroblasts can be targeted for cardiac regeneration.

Production of extracellular matrix

Homeostasis of the extracellular matrix has been traditionally considered to be the main function of the cardiac fibroblast in the uninjured heart. The cardiac ECM consists of interstitial collagens (predominantly type 1 and type 3), proteoglycans, glycoproteins, cytokines, growth factors and proteases[2]. Cardiac fibroblasts not only synthesize new matrix proteins but also express various metalloproteinases (MMPs) that degrade extracellular matrix. In the healthy heart, synthesis and breakdown of extracellular matrix are tightly regulated, but in pathological states, increased MMP expression and activity can lead to excessive ECM degradation and turnover[2, 41]. For instance, in patients with dilated cardiomyopathy, characterized by progressive heart failure and fibrosis, several MMPs including MMP2 were found to be elevated in the myocardial interstitium and extracts of hearts exhibited higher degree of collagenase and proteolytic activity[42–44]. In pressure overload cardiac hypertrophy in humans, MMP expression in the heart increases with the onset of left ventricular failure. In rodent models of pressure overload cardiac hypertrophy, increased expression of MMPs is associated with transition from compensation to heart failure[45, 46]. These observations suggest that dynamic changes in collagen content and turnover are associated with and perhaps contribute to cardiac pathologic states.

Following acute myocardial infarction fibroblasts not only increase the synthesis of ECM proteins at the site of injury (replacement fibrosis) but also increase ECM protein synthesis remote from the injury [7, 47]. The deposition of fibrous tissue in areas remote from the area of injury has been termed reactive fibrosis[1, 48]. Reactive fibrosis contributes to the pathophysiologic process after acute myocardial injury by altering chamber compliance and increasing the stiffness of the ventricles. In the normal heart 85% of the collagen is type I, while early after injury the predominant type of collagen synthesized particularly in the injury area is type III[49]. Type III collagen is more susceptible to stretch than type I collagen and enables the infarcted ventricle to be more distensible particularly within the first two weeks. As the scar matures, the ratio changes with more of type I collagen predominating than type III [49, 50].

Production of cytokines and growth factors

Cardiac fibroblasts express a rich array of cytokines and growth factors that affects wound healing via autocrine or paracrine mechanisms[51]. The expression of cytokines is markedly altered in diseased conditions, often contributing to the pathophysiological process. After myocardial infarction, cardiac fibroblasts are the main source of expression of the pro-inflammatory cytokine IL-1 β [7, 52]. IL-1 β enhances fibroblast migration and promotes turnover of the cardiac ECM via inhibition of collagen type 1 and type 3 synthesis and increased secretion of MMPs[2, 53]. Cardiac hypertrophy is usually accompanied by cardiac fibrosis and cardiac fibroblasts express the pro-fibrotic cytokine IL-6 that promotes cardiac fibroblast proliferation and fibrosis[54]. It is thought that mechanical stretching of the cardiac fibroblasts, that potentially occurs secondary to cardiac dilatation or changes in

hemodynamic burden in the heart, induces expression of pro-fibrotic cytokines, expression of ECM proteins and ECM receptors on cardiac fibroblasts[55].

Role in wound angiogenesis

Endothelial cell proliferation accompanies fibroblast proliferation early after myocardial injury and neovascularization of the injury region promotes wound healing and cardiac repair, while excessive neovascularization beyond the region of injury can promote hemorrhage[56]. Fibroblasts exist in close association to endothelial cells and are known to express pro-angiogenic cytokines (FGF & VEGF) as well as anti-angiogenic cytokines (PDGF & CTGF)[51][57]. Fibroblasts also express Tissue inhibitor of metalloproteinases (TIMPs) and TIMPs are known to inhibit and promote angiogenesis in a context dependent manner[58][59]. It is thus possible that fibroblasts contribute to the regulation of wound angiogenesis via expression of angiogenic and anti-angiogenic molecules and more studies will be needed to delineate the role of fibroblasts in regulating post infarct angiogenesis.

Role in electro-mechanical transduction after injury

Cardiac fibroblasts have high membrane resistance, that makes them good conductors of electrical signals. They can physically communicate with adjoining cardiac myocytes *ex-vivo* through gap junctional proteins such as connexin 43, although the physiological relevance of such interactions *in vivo* has not been demonstrated[60, 61]. As fibrosis is associated with disruption of the normal arrangement of myocytes and fibroblasts, heterogeneity in conduction can occur around the area of fibrosis and predispose to life threatening re-entry tachycardias. How fibroblast to fibroblast and fibroblast to myocyte communication in the region of scar influence the genesis of arrhythmias is an active field of investigation in arrhythmia biology.

Role in scar maturation and contraction

As alluded earlier in this review, a subset of cardiac fibroblasts after cardiac injury, acquire new phenotypic characteristics, express α -smooth muscle actin and are known as myofibroblasts[62]. The uninjured heart except the valvular interstitium does not have myofibroblasts[63]. In changing their phenotype to a myofibroblast, cardiac fibroblasts are thought to transition via an intermediate stage called proto-myofibroblast, characterized by expression of early stress fibers such as β -actin and γ -actin[40, 64]. The proto-myofibroblasts subsequently transition into fully differentiated myofibroblasts in the presence of appropriate signals from the ECM as well as neighboring cells[40]. A fully differentiated myofibroblast expresses α -smooth muscle actin but can be distinguished from a smooth muscle cell by low expression of other smooth muscle proteins such as heavy chain smooth muscle myosin[2].

Myofibroblasts contribute to scar contraction and remodeling[63, 65]. Scar maturation and contraction results in decreased surface area of the scar and leads to a thinner scar with greater tensile strength. Renewal of matrix proteins plays a critical role in persistence of scars in other organs such as the liver [66] and myofibroblasts by continuing to turn over matrix proteins likely perpetuate the persistence of scars in the heart[6, 18]. It is interesting to note that in organs such as the skin that retain the ability to heal wounds without scar

formation, myofibroblasts undergo apoptosis in the later phases of wound healing[48, 67]. Myofibroblasts have been found to be embedded in mature cardiac scars in humans years after myocardial injury and it is not clear whether the persistence of myofibroblasts in chronic scars months or years after cardiac injury is secondary to these cells being resistant to apoptosis[48].

Signals regulating fibroblast activation

Modulation of signals mediating early fibroblast activation and late remodeling is an area of intense scientific interest for its obvious therapeutic implications. An exhaustive description of mediators and signaling pathways affecting fibroblast function after injury is beyond the scope of this review but in this section we briefly describe some of the major signaling pathways involved. The region of myocardial necrosis is first infiltrated with polymorphonuclear cells. Over the first 48 hours, neutrophils are replaced by a mononuclear/macrophage infiltrate. Macrophages in the region of injury release pro-inflammatory cytokines, such as $\text{TNF}\alpha$, $\text{IL-1}\beta$ and IL-6 that play an important role in fibroblast activation and proliferation (Figure 3)[39].

The $\text{TGF}\beta$ /SMAD axis is known to promote fibroblast activation and proliferation after cardiac injury (Figure 3)[68]. $\text{TGF}\beta$ is markedly up-regulated in the infarcted heart with different isoforms displaying different temporal patterns of expression[69]. $\text{TGF}\beta 1$ and $\text{TGF}\beta 2$ are induced early while $\text{TGF}\beta 3$ shows delayed and persistent upregulation[70]. $\text{TGF}\beta$ in the acutely injured heart is expressed by macrophages as well as fibroblasts in the injury area. Although the precise functions of the different isoforms are unclear, downstream mediators of $\text{TGF}\beta$ (SMAD 2,3,4) are significantly elevated in the infarct area and inhibition of the $\text{TGF}\beta$ /SMAD axis is associated with decreased fibrosis and amelioration of late infarct remodeling[69, 71].

The renin angiotensin system (RAS) is being increasingly recognized to exert pro-fibrotic effects[10, 72, 73]. Angiotensin II levels are elevated in the heart within days of injury and Angiotensin II is known to enhance cardiac fibroblast proliferation and expression of collagens[74]. The expression of angiotensin receptors is more pronounced in cardiac fibroblasts in the injured heart[75] than on cardiac myocytes and pharmacological inhibition of the angiotensin axis has been shown to ameliorate cardiac fibrosis[76]. Furthermore, Angiotensin II has been shown to act upstream of $\text{TGF}\beta$ and at least in part, $\text{TGF}\beta$ mediates effects of Angiotensin. Thus a complex interaction of the RAS and $\text{TGF}\beta$ /SMAD axis may affect post infarct remodeling (Figure 3)[75].

Endothelin-1 (ET-1) is a potent vasoconstrictor and secreted predominantly by endothelial cells but endothelin can be expressed by inflammatory cells and fibroblasts[77]. Endothelin-1 is a downstream mediator of $\text{TGF}\beta$ and some studies suggest that ET-1 can work with $\text{TGF}\beta$ to promote myofibroblast persistence (Figure 3)[77, 78]. Ang II can also exert its effects via ET-1 and these studies suggest that ET-1 can work downstream of $\text{TGF}\beta$ and Ang II to drive fibroblast activation. Endothelin antagonists are currently approved for the treatment of pulmonary hypertension and future studies will determine their benefit in cardiac fibrosis[79].

Signaling systems important for cardiovascular development have also recently been shown to regulate cardiac fibrosis and repair after acute injury. We demonstrated that epicardial cells expressed Wnt1 after acute ischemic cardiac injury and Wnt1 expressing epicardial cells, in a β -catenin dependent manner, underwent EMT to adopt a fibroblast phenotype[35]. A dynamic notch signaling injury response has been shown to activate the epicardium and contribute to epicardial derived cardiac fibroblasts[36]. Thymosin β 4 promotes epicardial cell migration and differentiation of epicardial progenitors to promote neovascularization[80]. Although currently, there is no specific therapy for the treatment of cardiac fibrosis, the detailed dissection of signaling pathways affecting various aspects of fibroblast activation provide fresh opportunity for targeting the cardiac fibroblast.

Fibroblasts and cardiac regeneration

More recently, reprogramming techniques have been applied to induce a fibroblast to change fate into a cardiomyocyte (Figure 4). Cardiac fibroblasts isolated from the skin can be transduced with a cocktail of transcription factors to create induced pluripotent stem (iPS) cells. Such iPS cells can be differentiated *in-vitro* to generate patient specific cardiomyocytes that could be transplanted into a failing or infarcted heart (Figure 4A). More recently, direct reprogramming approaches have been described where the introduction of a specific combination of factors can directly reprogram a cardiac fibroblast to a cardiomyocyte without having the fibroblast transition through a primitive undifferentiated state (Figure 4A). This technique is particularly appealing as introduction of such factors into fibroblasts in the region of scar can potentially change the fate of a fibroblast to a cardiac myocyte and reverse scarring (Figure 4B).

Ieda. et al demonstrated that a combination of transcription factors GATA4, Mef2C and TBX5 (GMT) when introduced into a cardiac fibroblast are sufficient to reprogram a cardiac fibroblast into a cardiomyocyte[81]. Moreover, the groups of Olson and Srivastava demonstrated that such direct reprogramming can also occur *in vivo*. The cocktail of transcription factors (GMT or GMT and Hand2) administered to mice hearts via retroviruses following myocardial infarction was sufficient to reprogram scar tissue into cardiac muscle cells that appeared to integrate with surrounding myocardium[82, 83]. Wada. et al have more recently demonstrated that the addition of Mesp1 and Myocardin to the GMT cocktail of transcription factors can induce human dermal fibroblasts to adopt a cardiomyogenic like fate[84]. Jayawardena. et al showed that delivery of a combination of microRNAs is sufficient to reprogram fibroblasts into a cardiomyogenic fate[85].

However with all the above reports, the fraction of fibroblasts undergoing successful direct reprogramming *in-vivo* into cardiac muscle cells remains small. More efficient delivery systems of transgenes or identification of novel factors that greatly increase the efficiency of direct fibroblast-myocyte reprogramming are needed for successful manipulation of large scars. Nevertheless, this method of directly reprogramming scar tissue into cardiomyocytes is undeniably attractive as it has the promise of changing chronic scar tissue into cardiac muscle cells and potentially reverse post infarct remodeling (Figure 4B).

Future Directions

There is mounting clinical evidence to suggest that fibrosis is an independent predictor of mortality and sudden death in patients with heart failure. Patients with a similar ejection fraction but having significantly greater cardiac fibrosis have a poorer prognosis[86]. There is no specific treatment for fibrosis and there is a dire need for major biological questions regarding cardiac fibrosis to be answered.

In this concluding section, we discuss some of the major questions that remain unanswered in the field of cardiac fibrosis. First, given the diversity of origin of fibroblasts contributing to cardiac injury, are there specific subsets of fibroblasts that need to be targeted both in the early and late phases of wound healing? Using fate mapping techniques, Kalluri's group recently demonstrated that after kidney injury, approximately 50% of fibroblasts in the injury area are of interstitial origin where as the remaining pool is derived from the bone marrow, endothelium (via endothelial-mesenchymal-transition) and epithelium (via EMT) [87]. Recent observations made in skin wound healing also suggest that different lineages of fibroblasts contribute to different aspects of wound healing in the skin[88]. Fibroblasts in the lower dermis are responsible for the deposition of extracellular matrix proteins and fibroblasts in the upper dermis are recruited only during re-epithelialization. Such concepts of specification of fibroblasts during wound healing can be tested in the heart after myocardial injury and can potentially identify whether specific subsets of fibroblasts should be targeted. In this regard, rigorous fate mapping studies should also be performed in the developing heart to identify other potential cell sources contributing to the cardiac fibroblast pool. Elucidation of mechanisms that promote cardiac fibroblast proliferation in the developing heart may identify new signaling pathways that regulate fibroblast activation after cardiac injury.

Second, identification of mechanisms perpetuating the persistence of scars long after injury are of paramount importance. In the liver, under certain circumstances, fibrosis is known to regress and regression of hepatic scars is thought to be secondary to inadequate ECM renewal rather than active catabolism and breakdown of collagen. Do scars in the heart persist secondary to continuous turnover of matrix proteins by myofibroblasts? What then are the factors driving myofibroblast turnover of ECM proteins long after cardiac injury?

Finally can cardiac fibrosis be reversed? Advances in direct reprogramming technology have provided proof of concept that fibroblasts in cardiac scar can be directly reprogrammed into cardiomyocytes. In theory, if a large number of myofibroblasts in the scarred region can be coaxed to adopt a cardiac muscle fate, there should be diminution in the amount of fibrosis. Moreover this strategy also generates new cardiac muscle in the region of scar formation and can potentially reverse deleterious ventricular remodeling. However, pockets of newly generated cardiomyocytes that do not integrate with surrounding cardiac muscle can lead to the formation of arrhythmogenic foci. This technology is also currently limited by the poor efficiency of fibroblast to cardiomyocyte reprogramming *in-vivo*. Optimization of transgene delivery as well as identification of factors that enhance the efficiency of direct reprogramming will be needed prior to translating these observations into clinical practice. Understanding the physiologic basis of fibroblast diversity, mechanisms regulating the

persistence of scars and manipulation of fibroblast plasticity in a spatiotemporal manner will be key for treating pathologic cardiac fibrosis.

List of abbreviations

CTGF	Connective tissue growth factor
ECM	Extracellular matrix
EMT	Epithelial-mesenchymal-transition
EndMT	Endothelial-mesenchymal-transition
FGF	Fibroblast growth factor
IL	Interleukin
MMP	Matrix metalloproteinase
TGF	Transforming growth factor
PDGF	Platelet derived growth factor
PDGFR	Platelet derived growth factor receptor
RAS	Renin-angiotensin system
TIMP	Tissue inhibitor of metalloproteinase
Wt-1	Wilms tumor 1
Tbx	T-box transcription factor
Mef	Myocyte enhancer factor

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HIGHLIGHTS

- Cardiac fibroblasts are the most abundant cardiac cell type
- Fibroblasts regulate cardiomyocyte proliferation during heart development
- Fibroblast activation is a critical early repair response after cardiac injury
- Persistence of fibroblasts in injury area leads to chronic scar and remodeling
- Reprogramming fibroblasts to myocytes can be a new strategy for treating cardiac scar

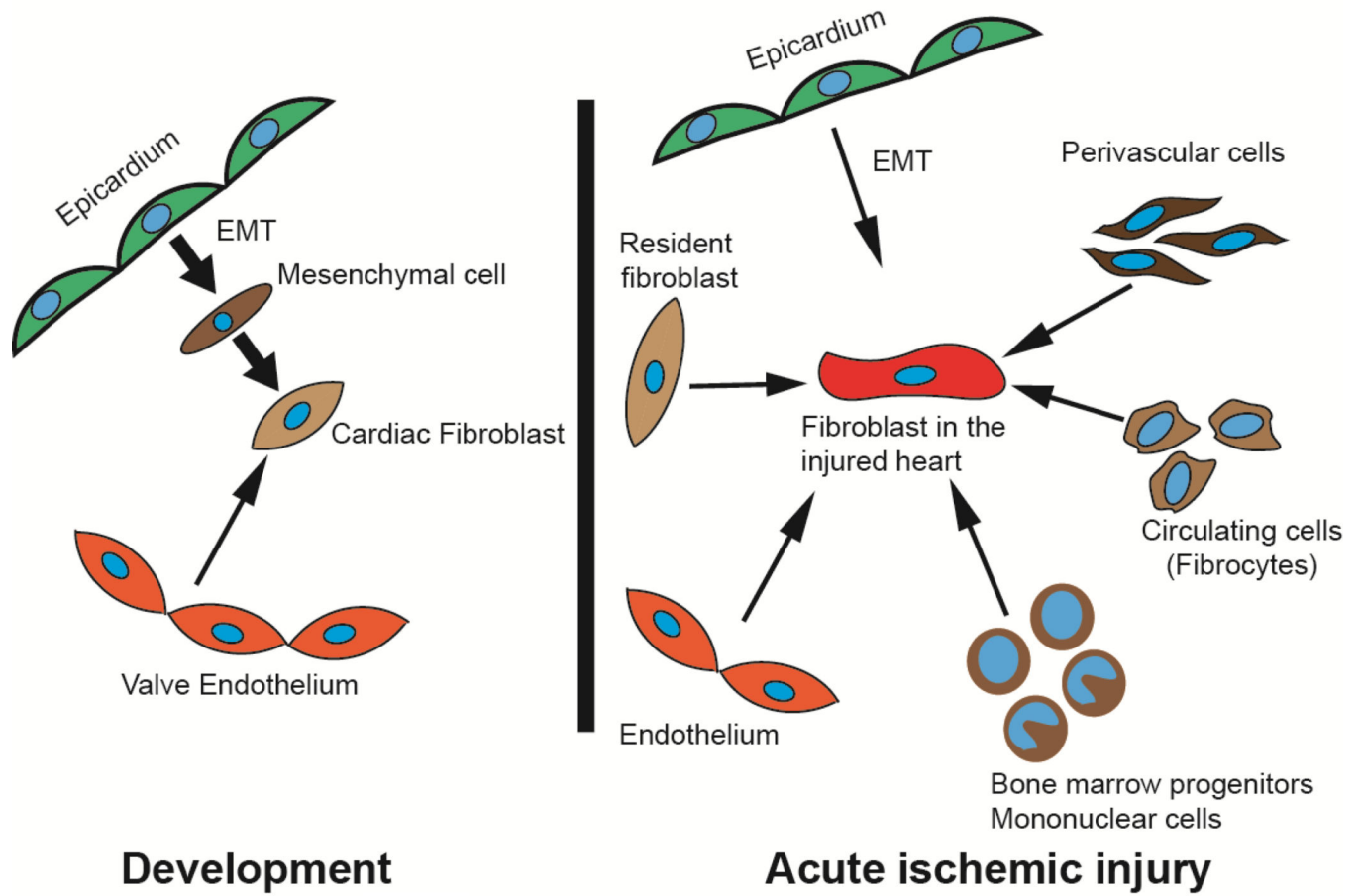


Figure 1. Origins of cardiac fibroblasts during cardiac development and following acute ischemic injury
 The cardiac fibroblast in the injured heart has diverse origins compared to the fibroblast in the developing heart.

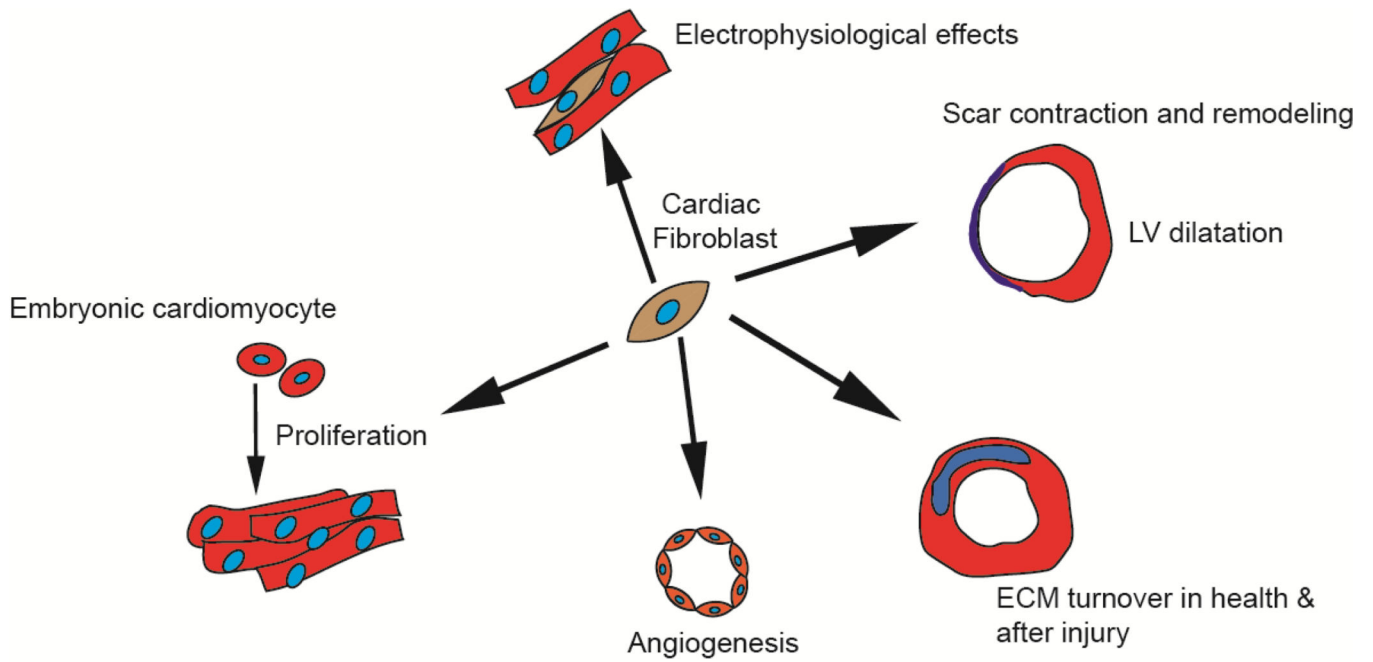


Figure 2. Major functions of the fibroblast during cardiac development and wound healing
 The fibroblast promotes proliferation of embryonic cardiomyocytes. It influences angiogenesis in the adult heart, regulates ECM turnover both in the adult uninjured heart and after acute injury. The fibroblast plays a pathophysiological role in scar contraction, adverse remodeling and ventricular dilatation and exerts electrophysiological effects.

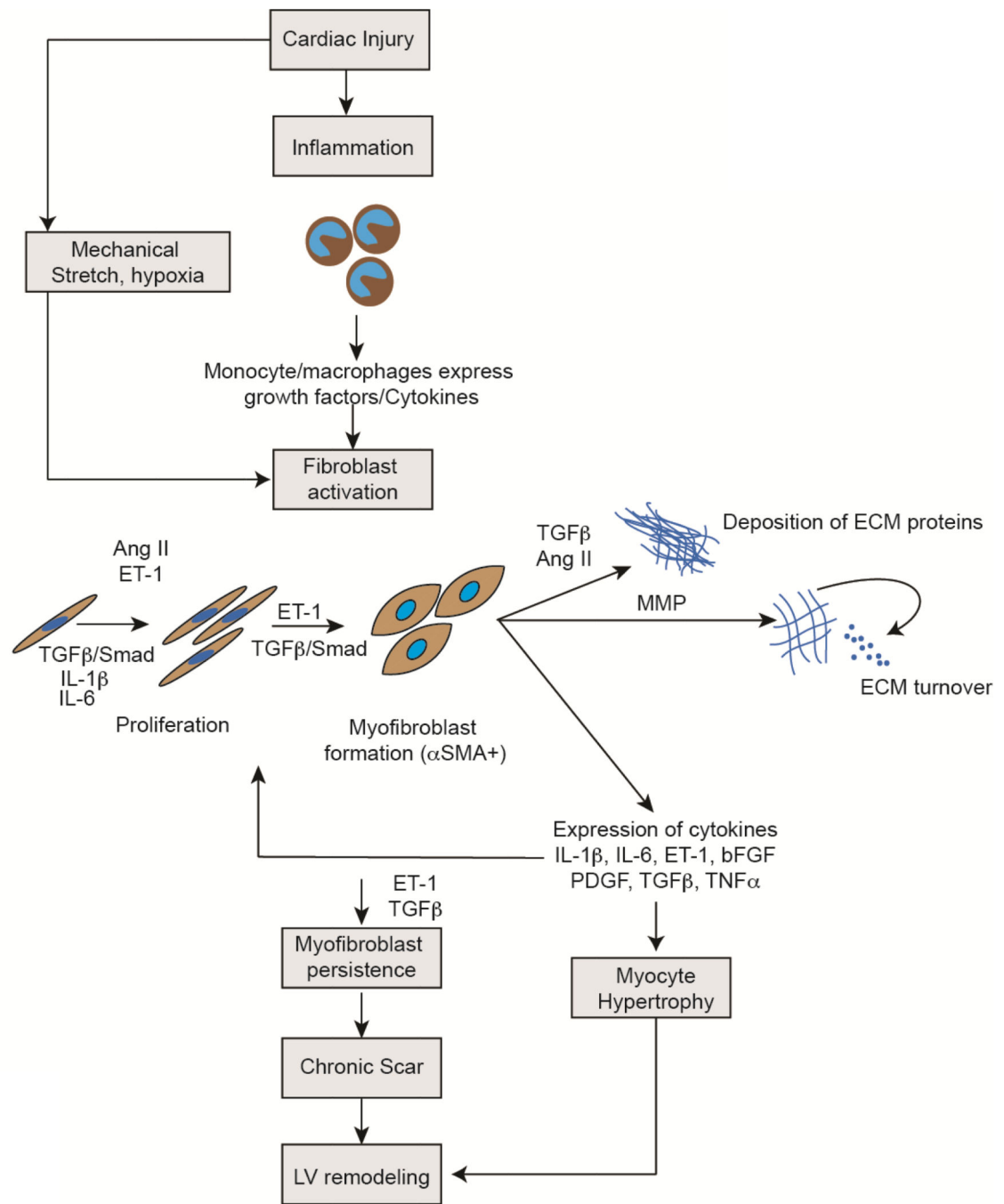


Figure 3. Signaling pathways regulating pathophysiological effects of cardiac fibroblasts after cardiac injury

Following cardiac injury, monocyte/macrophage infiltrate at the site of injury release growth factors and cytokines that activate cardiac fibroblasts. Mechanical stretch and hypoxia in addition are potent activators of fibroblasts. TGFβ, ET-1, Ang II and other inflammatory cytokines such as IL-1β and IL-6 induce cardiac fibroblast proliferation, formation of myofibroblast, deposition of extracellular matrix and matrix turnover. The fibroblast itself expresses many of these pro-fibrotic cytokines that exert effects via autocrine/paracrine mechanisms. Cytokines expressed by cardiac fibroblasts also induce cardiomyocyte

hypertrophy. Persistence of myofibroblasts in the injury region leads to continuing matrix turnover, formation of chronic scar and along with myocyte hypertrophy leads to adverse ventricular remodeling. (TGF β : Transforming Growth Factor β , IL: Interleukin, ET-1: Endothelin 1, Ang II: Angiotensin II, bFGF: Basic Fibroblast Growth Factor, PDGF: Platelet Derived Growth Factor, TNF α : Tumor Necrosis Factor α , ECM: Extracellular matrix)

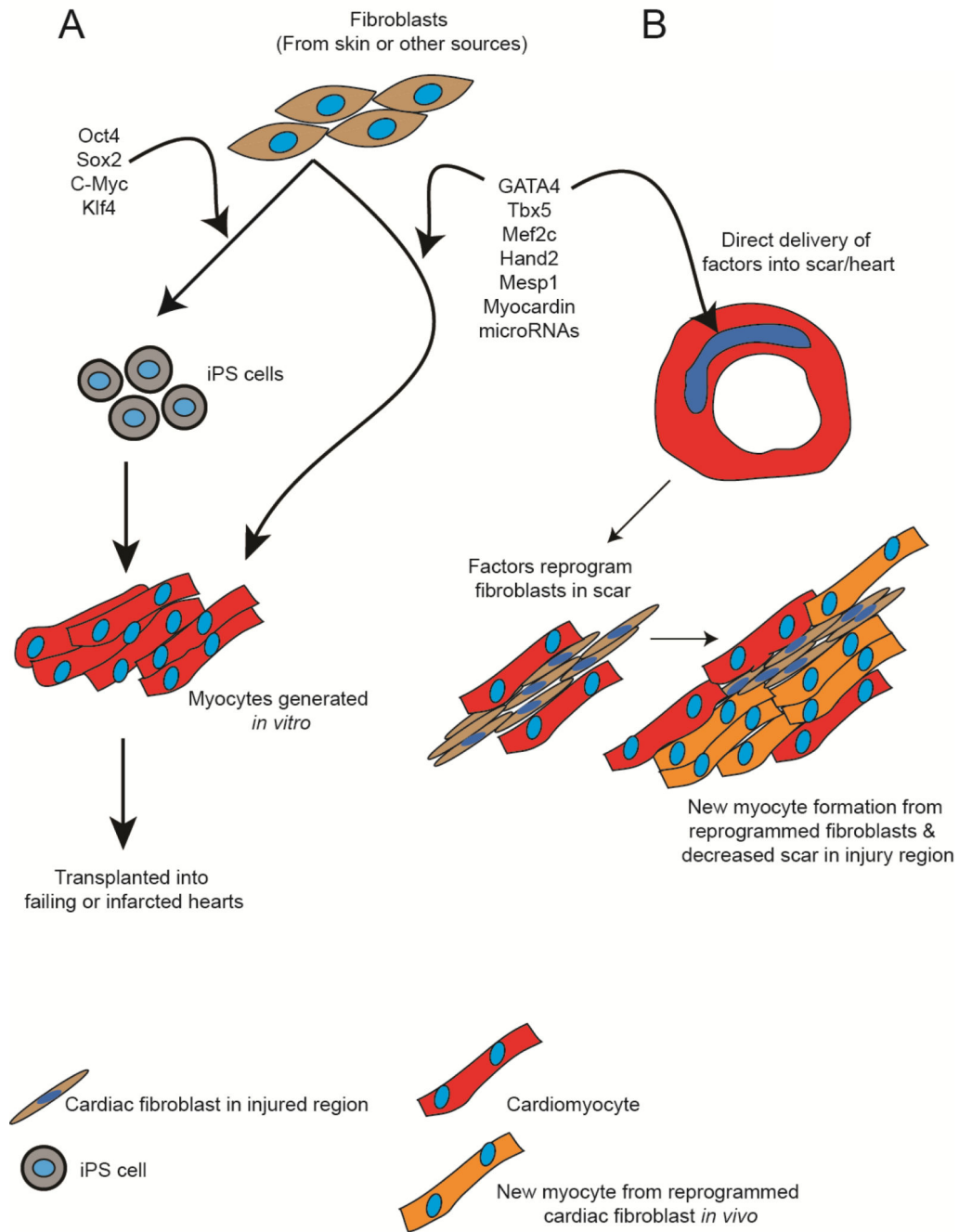


Figure 4. Targeting the fibroblast for cardiac regeneration

Fibroblasts can be reprogrammed into cardiomyocytes using cocktails of reprogramming factors. (A) In one approach, fibroblasts isolated from skin or other tissues of a patient can be transduced with cocktails of transcription factors to yield induced pluripotent stem cells (iPS cells). Subsequently iPS cells generated from these fibroblasts can be differentiated *in-vitro* into cardiomyocytes that could be subsequently transplanted into failing or injured hearts. Alternatively combinations of factors can be used to directly reprogram the fibroblast to a cardiomyocyte *in-vitro* bypassing the iPS step and then generated cardiomyocytes would be

transplanted in a similar manner. **(B)** In an alternative approach, combinations of factors can be directly delivered into the fibrotic region after cardiac injury. Cardiac fibroblasts in the injury region can directly reprogram into cardiomyocytes that potentially can integrate with existing cardiomyocytes to enhance cardiac function. By reprogramming fibroblasts in the injury region to form cardiomyocytes, this strategy would also decrease fibrosis and potentially ameliorate remodeling.

Table 1

Changes in cardiac fibroblasts numbers from late development to adulthood in the murine heart. Adapted from [3].

Developmental Stage	Fraction of all heart cells
E18.5	14%
Post natal Day 1	10%
Post natal Day 5	14%
Post natal Day 15	18%
Adult	27%

Table 2

Contribution of fibroblasts to the cellular composition of the adult murine and rat heart. (Numbers expressed as a fraction of all heart cells). Adapted from [3].

Cell type	Mouse	Rat
Cardiac fibroblasts	27%	64%
Myocytes	56%	30%
Non-myocyte/non cardiac fibroblasts	17%	6%