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Getting to the heart of myocardial stem cells and cell therapy

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Introduction

Heart disease is both common and deadly. Cardiovascular disease is a global epidemic as it is the number one cause of death worldwide and it is estimated that one in three adults in the United States have cardiovascular disease¹. While a number of pioneering initiatives have transformed our treatment of cardiovascular disease, new therapies are required to further address the growing incidence of this deadly disease. Intense interest has focused on regenerative medicine as an emerging strategy for chronic diseases such as cardiovascular disease.

A number of human tissues including skin², gut, liver^{3–6} and skeletal muscle^{3, 7} have a tremendous regenerative capacity. For example, skeletal muscle is able to completely restore its cellular architecture and function following an injury that destroys more than 80% of the muscle^{7, 8}. This regenerative response lacks a fibroproliferative response (i.e. formation of scar) and is associated with restoration of the vasculature, myofibers and extracellular matrix. Unlike skeletal muscle, the regenerative capacity of the adult heart is more limited.

Recent studies suggest that the adult heart is capable of cellular turnover and limited regeneration following injury although the networks that govern this process are ill defined. The use of genetic mouse models and molecular biological techniques are unveiling cell populations, pathways and extracellular cues that may direct cardiac regeneration and provide a platform for further investigation. The goal of this review is to examine the endogenous regenerative capacity of the adult heart and highlight new experimental regenerative therapies aimed at restoring myocardial architecture and function.

Endogenous repair and regeneration of the metazoan heart

Previous studies have demonstrated that metazoans such as the newt and zebrafish are capable of cardiac regeneration in response to a significant injury^{9–12}. This myocardial regenerative response is complex and occurs over a two month period. In response to a myocardial injury (amputation of 30–40% of the ventricular chamber), there is formation of a fibrin clot, subsequent dedifferentiation of cardiomyocytes and recruitment of specialized cell populations including epicardial and ventricular myocardial cell populations^{11–13}. Importantly, the regenerative response observed in both the newt and zebrafish lacks the formation of scar¹³. These results support the notion that there is an inverse relationship between scar formation and myocardial regeneration (Figure 1). Moreover, the studies in these regenerative models have defined the role of Notch^{14, 15}, fibroblast growth factor2^{15, 16} and retinoic acid¹⁷ signaling pathways in myocardial regeneration. Examination

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of these regenerative organisms as well as mammalian tissues that have an enhanced regenerative capacity are instructive regarding the mechanisms and pathways that govern this repair process in response to injury.

Endogenous repair and regeneration of mammalian tissues

Every tissue is a product of stem cells and evidence suggests that essentially every adult mammalian tissue harbors a stem cell or progenitor cell population that participates in the maintenance or regeneration of their host tissue(s) in response to injury (Table 1)³⁰. For example, the satellite cell population occupies a niche (satellite cells are sandwiched between the basal lamina and the plasmalemma in close association with the myofiber) and resides within adult skeletal muscle^{7, 31}. The satellite cells represent the myogenic stem cell population that is quiescent in unperturbed muscle. In response to a severe injury, the quiescent satellite cells become activated (reenter the cell cycle), they proliferate and in response to cellular and extracellular cues they differentiate to form centronucleated myofibers (the hallmark of regenerated skeletal myofibers) thus restoring the cellular architecture of the injured tissue³¹. Importantly, the satellite cells are capable of self-renewal and reestablish their quiescent pool of myogenic stem cells^{32, 33}. These studies emphasize the dynamic capacity of adult mammalian skeletal muscle to completely regenerate in response to injury. All striated muscle does not respond in a similar fashion to an injury.

Endogenous repair and regeneration of the mammalian heart

The neonatal mammalian heart is associated with considerable growth and cellular proliferation of myocardiocytes (Figure 2A). During the first week of life, the neonatal heart continues to proliferate and grow as measured by proliferative assays (BrdU pulse assays³⁴ and tritiated thymidine assays³⁵). This postnatal stage is also marked by increased apoptosis in the developing heart, which suggests an active modeling or sculpting process that is associated with growth and modulation by hemodynamic challenges, hormonal surges and changes occurring in extracardiac tissues. Following this postnatal period, the mammalian heart is associated with modest cellular turnover.

Recent studies utilizing labeling strategies and genetic mouse models suggest that the adult heart is capable of cellular replacement of cardiomyocytes, repair and limited regeneration in response to an ischemic or nonischemic injury. One study utilized radiocarbon cellular dating to examine cardiomyocyte turnover. Bergmann and colleagues relied upon the integration of carbon-14 into DNA as a measure of cardiomyocyte turnover in the adult heart³⁶. The DNA of all organisms incorporated high concentrations of carbon-14 that were generated from nuclear bomb testing (which persisted until the 1963 Limited Nuclear Test Ban Treaty) into DNA as a measure of cardiomyocyte cellular kinetics or turnover in the adult heart. Therefore, the nuclear bomb testing and subsequent increase in atmospheric carbon-14 provided a pulse such that postnatal cellular turnover could be estimated by comparing the age of the DNA of the cardiomyocytes to the patients' chronological age. This cellular dating technique and mathematical modeling support the notion that the adult heart is capable of cellular turnover (representing cardiomyocyte renewal) at the rate of 1% per year (although this rate decreased to 0.4% at age 75)³⁶. The results further suggested that approximately 45% of the cardiomyocytes were generated or renewed in the 50 year old heart since birth. These studies provide an innovative strategy to examine the cardiomyocyte turnover of the adult human heart. Issues of polyploidy and origin of the cardiomyocytes remain active issues of investigation and further studies using similar strategies to examine the cardiomyocyte turnover in distinct populations are warranted. Nevertheless, these results provide further support that the adult human heart has ongoing turnover of the

cardiomyocytes and support the notion that strategies to enhance this turnover may prevent the genesis of heart failure³⁶.

A second study utilized genetic mouse models to evaluate cellular kinetics in the unperturbed and post-injured heart. Utilizing an inducible cardiomyocyte-specific transgenic fate mapping (MerCreMer) strategy in the mouse, cardiomyocytes were irreversibly labeled with the GFP reporter following the pulse of tamoxifen³⁷. In contrast, cardiac stem cells or progenitors were not genetically labeled in response to tamoxifen as they did not express the cardiomyocyte specific marker (myosin heavy chain 6)³⁷. This fate mapping strategy allowed for the measure of cardiomyocyte turnover in the adult mouse heart. While little to no cardiomyocyte turnover was observed in the unperturbed heart, the genetic labeling strategy following myocardial injury was associated with approximately 15% of unlabeled cardiomyocytes had undergone cellular turnover (presumably from a progenitor or stem cell population). These genetic studies are further supported by BrdU incorporation studies, which further support the hypothesis that cardiomyocyte renewal occurs following myocardial injury (Figure 2B)³⁸.

Collectively, these and other studies support the notion that the postnatal mammalian heart is associated with cardiomyocyte renewal (cellular turnover) which is increased following injury. These results further suggest that the amplification of resident or recruited stem cells or progenitors and regenerative myocardial pathways could increase endogenous myocardial repair of the acutely injured heart.

Resident cardiac stem and progenitor cell populations

Previous studies have identified the contribution of stem cell and progenitor cell populations that are resident in the postnatal heart that are capable of generating cardiomyocytes (Figure 3^{39–50}). While a hierarchy of stem and progenitor cell populations have not been defined, a clonal c-kit⁺ cell population has been shown to generate all lineages of the heart, increases in number following myocardial injury, undergoes self renewal and generates cardiomyocytes in a number of mammalian models (rat, mouse, dog, etc.) including human^{42, 43}. These c-kit expressing cells have been shown to occupy a niche within the adult heart. Using immunohistochemical studies, subpopulations of c-kit expressing cells have been identified and include those that express c-kit only versus those that coexpress c-kit with cardiac transcription factors and sarcomeric proteins. This continuum of c-kit expressing cells (with and without cardiac progenitor cell (CPC) to the committed progenitor cell to the immature cardiomyocyte⁴³.

An additional cell population includes the cardiac side population (SP) cells that express multidrug resistance proteins (members of the ATP-binding cassette transporter family) and are isolated by FACS analysis based on their ability to efflux Hoechst 33342 dye^{46–48}. These SP cells populate the heart early during development, are resident in the adult heart and increase in number within three days of cardiac injury. Using immunohistochemical techniques, these cardiac SP cells, following injury, have been shown to coexpress cardiomyocyte specific sarcomeric proteins (suggesting that the cardiac SP cells are capable of differentiating to fetal cardiomyocytes)⁴⁷. Previous studies have defined that members of the ABC transporter family serve not only to mark the SP cells but also play an important cytoprotective role for these stem/progenitor cells in response to oxidative stress⁴⁶. Transcriptome analysis has been useful in defining the molecular signature of cardiac SP cells that are isolated from the adult heart compared to other embryonic and somatic stem cell populations^{46, 47}.

Studies have shown that Sca1⁺ cells can differentiate into cardiomyocytes⁴⁴. Resident Sca1⁺/CD31⁻ cardiac progenitors have been reported to increase in number and more than double fourteen days following acute myocardial infarction. These cardiac Sca1⁺/CD31⁻ cells were capable of differentiation to endothelial and cardiomyocyte lineages in vitro and in vivo following the delivery into the post-injured heart. The transdifferentiation of the engrafted Sca1⁺/CD31⁻ cells were accompanied by a significant improvement of left ventricular systolic function compared to controls⁴⁵.

Further studies are warranted using a genetic labeling strategy (Cre-*loxP* technology) to fate map the c-kit, SP and Sca-1 expressing cell population during development, aging and following perturbations (including coronary artery ligation and pressure overload) using conventional and inducible genetic technologies. These fate mapping techniques will enhance our understanding of the degree to which these cell populations contribute to the renewal of cardiomyocytes and the vasculature following injury or aging.

Further cell populations that may represent a cardiac stem cell pool, progenitors (such as transit amplifying cells) or other stem cell populations that are capable of generating cardiomyocytes include EPC (endothelial progenitors)^{51, 52}, MSC (mesenchymal stem cells)^{53, 54}, CD34⁺ cells^{55, 56}, myofibroblasts and others that have been reported to participate in cardiomyocyte renewal and regeneration. Studies support the notion that these cell populations form cardiomyocytes under permissive conditions.

Cardiac progenitors that are obtained from adult hearts (following an endomyocardial biopsy) coalesce in culture to form a three dimensional spherical structure termed a cardiosphere. Two independent laboratories have generated cardiospheres from both mouse and human biopsy specimens of adult hearts. These cardiospheres (up to 150 microns in size) have a tremendous proliferative capacity (generating more than one million cardiospheres in a one month period) and are capable of forming differentiated, contractile cardiomyocytes^{39, 40}. Cardiospheres also represent a heterogeneous cell population with a cortex of c-kit expressing, proliferating cells and a mantle of differentiated cardiomyocytes^{39,40}. Delivery of cardiospheres as a graft following myocardial injury resulted in improved cardiac function in rodent models and limited clinical trials are in progress to evaluate these autologous cell preparations in patients. It is unclear whether cardiospheres are derivatives of a resident cardiac stem/progenitor cell population or whether they represent reprogramming of cardiomyocytes (dedifferentiation) or progenitor cell populations. Moreover, a recent study demonstrated that transplantation of cardiosphere derived cells into the post-injured pig model further induced repair and regeneration by endogenous cardiac progenitors⁴¹.

A complementary cell population to the resident cardiac stem/progenitor cell population is the reprogrammed, induced pluripotent stem cell population (iPSC). These iPSCs have been derived from somatic cells such as skin fibroblasts through forced expression of gene expression (Oct3/4, Sox2, c-Myc and Klf4 vs. OCT4, SOX2, NANOG and LIN28) in mouse (2006)^{57, 58} and human (2008)^{59, 60}. These iPSCs have been shown to mirror embryonic stem cells with regards to their proliferative capacity, pluripotency, chimera formation, teratoma formation and capacity to differentiate to all germ layer derivatives. Recent studies suggest that reprogramming is possible without genetic alteration of the somatic cell. In addition to these initiatives, the use of chemical genetics and exposure of cells to small molecules may be sufficient to direct somatic or stem cells to a cardiovascular fate^{61, 62}. An alternative reprogramming strategy is to decipher the pathways or factors that will directly convert somatic cells (fibroblasts) to cardiomyocytes, therefore, bypassing the pluripotent state. An example of this strategy included the forced expression of three developmental transcription factors (Tbx5, Gata4 and Mef2c) to reprogram murine cardiac fibroblasts into

cardiomyocyte-like cells that were similar to neonatal cardiomyocytes⁶³. These results provide a proof of concept and rationale for future studies aimed at reprogramming human cardiac fibroblasts to a cardiomyocyte fate. This field is rapidly evolving and will provide a platform for disease specific stem cell populations, personalized stem cell populations and provide cell sources for pharmacogenetics studies.

Cell therapy for chronic diseases

The delivery of allogeneic or autologous cellular populations for the treatment of chronic diseases has been utilized for more than forty years. Since the world's first and second successful bone marrow transplantation in 1968 at the University of Minnesota, this cellular therapy has been used to treat an array of diseases including solid tumors, mucopolysacharidoses, and hematological cancers⁶⁴. In total more than 50,000 patients worldwide receive this life-saving therapy each year⁶⁵. Bone marrow transplantation provides the rationale and the feasibility for using cellular therapeutic strategies for the treatment of terminal diseases such as cardiovascular disease and heart failure.

Cell therapy for cardiovascular disease

While bone marrow transplantation has been used successfully to treat terminal diseases, certain challenges need to be overcome in order to translate the use of cellular therapy to other tissues such as the heart. For example, the heart is unlike the bone marrow in that it has a highly structured cellular architecture that is electrically and functionally synchronized to produce more than 2 billion heart beats in a lifetime. Further, the working load of the heart is constantly changing and responding to local and systemic stimuli. These hemodynamic challenges provide both permissive and repressive challenges for the use of cell therapy. Third, the geometrical shape of the heart adapts in response to injury, scar, hemodynamic demand, etc. as remodeling promotes the change from a prolate ellipse to a spherical shape due to hemodynamic load. Collectively, these challenges are balanced with increasing prevalence of cardiovascular disease, decreasing donors for heart transplantation (the only definitive therapy for advanced heart failure) and a need to develop new therapies for this patient population.

As bone marrow transplantation has been an effective therapy for human diseases, studies were undertaken to examine the capacity of unfractionated bone marrow mononuclear cells (which contain hematopoietic stem cells) to transdifferentiate to a cardiomyocyte fate and improve the functional performance of the injured heart. Using genetic labeling strategies, studies demonstrated significant, limited or an absence of labeled bone marrow mononuncleated cells or stem cells to generate cardiomyocytes following the delivery into the post-injured rodent heart^{66–69}. Despite differences in reported differentiation potential, several studies demonstrated a functional improvement in response to the delivery of hematopoietic stem cells, unfractionated bone marrow mononuclear cells and other cell populations (including fibroblasts, skeletal myoblasts, mesenchymal stem cells, endothelial progenitors, cord stem cells, etc.)^{69–71}. Despite the variability of the results, the preclinical studies demonstrating positive results fueled the design of clinical trials using cell therapy for the treatment of cardiovascular disease (Table 2).

A majority of the cardiovascular clinical cell therapy trials have utilized autologous cell populations (the exception is the use of allogeneic mesenchymal stem cells), which obviate immunological mediated cellular rejection. The mode of delivery has been primarily intracoronary but has also included intramyocardial and endocardial routes. The initial nonrandomized studies delivered bone marrow mononuclear cells (intracoronary) seven days following percutaneous revascularization and observed no significant change in function (LVEF) compared to controls three months following delivery. TOPCARE-AMI

was the first randomized clinical trial and compared two different cell populations (peripheral derived stem cells vs. mononucleated bone marrow cells) five days following PCI. At one year following cell delivery there was a decrease in infarct area in both patient populations⁷². The first randomized, controlled study (n = 60) was the BOOST trial which delivered bone marrow mononucleated cells (intracoronary) five days following PCI. While LVEF improved in those patients that received cell therapy at six months compared to controls⁷⁷, no significant differences were noted at eighteen months post-delivery⁷³. In contrast, the larger REPAIR-AMI trial (n = 204) was a randomized, controlled study and reported a modest improvement of LVEF as measured by ventriculography in patients receiving intracoronary delivery of bone marrow mononucleated cells five days post-PCI at four and twelve months follow-up compared to controls^{74, 78}. The ASTAMI trial (n = 100) also examined the efficacy of intracoronary delivery of autologous bone marrow mononuclear cells six days post-PCI and observed no significant change in LVEF between experimental and control groups at three years post-intervention⁷⁵.

Potential side effects associated with cell therapy include: arrythmyogenesis, tumorigenesis, myocardial injury, infection (bacterial or viral pathogens) or immunological responses. One example of the side effects associated with the delivery of autologous human skeletal myoblasts includes the genesis of ventricular arrhythmias and the need for implantable cardioverter defibrillator support^{76, 79}. Another possible side effect is the immunological response to the cardiac delivery of allogeneic cell sources. While these complications are possible, very few side effects have been observed in clinical trials performed in the US and in Europe.

Collectively, these clinical trials support the conclusion that cell therapy for cardiovascular disease is relatively safe; it may modulate remodeling and may have a modest improvement in cardiac function (Table 2). Many of these clinical studies utilized ejection fraction as the only endpoint for the efficacy of these cell transfer studies which may be insufficient. Other primary or secondary endpoints for future studies may include exercise tolerance, infarct size, five year survival rate or progression to heart failure (NYHA Stages I–IV). Furthermore, future studies aimed at a mechanistic understanding of cell therapy for heart disease will be important for the advancement of this field.

Paracrine hypothesis and myocardial regeneration

An increasing number of studies suggest that cell therapy may also be associated with a bystander effect through the release of cytokines, antiapoptotic factors or growth factors which may improve cardiac function. For example, studies suggest that transplanted cells release paracrine factors that decrease program cell death (thereby limiting the remodeling process), promote angiogenesis or enhance myocardial regeneration mediated by the endogenous cardiac stem/progenitors that are resident in the adult heart^{80, 81}. Alternatively, small molecules or growth factors such as neuregulin1 may induce cell cycle reentry and division of differentiated cardiomyocytes⁸². Increasing evidence supports the hypothesis that paracrine factors or the delivery of small molecules may promote myocardial repair and regeneration.

Gaps in knowledge and challenges for the future

To further examine these mechanistic questions and accelerate cell based therapies for cardiovascular disease, the National Heart, Lung and Blood Institute funded the Cardiovascular Cell Therapy Research Network (CCTRN) that includes investigators at five institutions across the US⁸³. This network and other ongoing trials will need to collaborate with bench investigators to define the patient population that benefits from cell therapy (i.e. ischemic vs. nonischemic dilated cardiomyopathy), the optimal cell population (autologous

vs. allogeneic vs. EPCs, skeletal muscle satellite cells, bone marrow mononuclear cells, CD34+ cells, MSCs, cardiospheres, etc.), mode of delivery (intracoronary, intravenous, intramyocardial, etc.), cell preparation (cultured and possibly reprogrammed vs. freshly isolated cells), numbers of cells delivered, site of delivery (infarct related artery, border region of injured myocardium, distant ventricular delivery, atrium, etc.), mechanisms of action of cell therapy (paracrine effect to limit apoptosis, promote neovascularization, promote myocardial regeneration, limit fibroproliferative response) and the role of multiple or serial interventions with cell delivery. These studies will further benefit from the design of FDA approved cell labeling strategies that will allow for the detection of single cells using imaging technologies. Moreover, cell therapy studies performed in combination with patches or scaffolds and ventricular assist devices used as a bridge to heart transplantation will allow histological analyses of the explanted heart at the time of transplant. These technologies provide new mechanistic insight regarding the use of cellular therapy for treatment of cardiac failure.

To complement these clinical trials, the National Heart, Lung and Blood Institute established the NHLBI Progenitor Cell Biology Consortium, which is a collaborative network for the exchange of reagents and acceleration of discoveries related to stem cell and progenitor cell biology⁸⁴. One of the goals of this consortium is to gain an understanding of the mechanisms that direct stem and progenitor cells to a cardiac fate. Importantly, this network will provide an infrastructure for the field and will address issues including but not limited to:

- The definition of a hierarchy of somatic stem and progenitor cells that reside in the adult heart.
- The definition of transcriptional networks, epigenetic networks and microRNA networks that direct stem cells toward a cardiac fate.
- To provide protocols for stem/progenitor cellular characterization and cardiomyocyte differentiation pathways.
- To establish nonviral strategies to reprogram somatic cells to cardiomyocytes or iPSCs.
- To establish fate mapping strategies to define the contribution of selected stem/ progenitor cell populations to the cardiac lineage during development and following myocardial injury.
- To compare specific cardiac and hematopoietic stem/progenitor cells using FACS, transcriptional, microRNA, functional or epigenetic analyses.

Together, these translational and basic science networks of investigators facilitate communication and collaborations. They further provide an infrastructure that supports ongoing and future discoveries that are intended to lead to new therapies for heart disease.

In summary, the field of cardiac regeneration has exploded with interest and opportunities. While significant advances have energized the field, further studies will be necessary to provide additional mechanisms and insights into the possibility of identifying the key(s) that will promote myocardial repair whether the strategy relies on the endogenous repair program of the heart or the use of a cell delivery program. While neither the endogenous repair program nor the cell therapy programs are ready for prime time, they will serve as a platform that will launch the field forward. Collaboration and exchange of data through professional networks should amplify and accelerate the science to move the field towards effective therapies.

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Figure 1.

An emerging hypothesis is that the regeneration potential is linked to the fibroproliferative response of the injured tissue. A) The development of scar following injury results in tissue (cardiac) dysfunction. The ability of an injured heart to regenerate results in improvement of cardiac function. B) Models suggest there is a balance between scar formation and regeneration. These models support the notion that scar formation impairs regeneration and a regenerating tissue lacks scar.



Figure 2.

The neonatal and adult mammalian hearts are capable of cellular proliferation. A) Relative cardiomyocyte proliferation from BrdU labeling and tritiated thymidine incorporation studies of neonatal and adolescent mice support the notion that the heart's proliferative capacity decreases with age (see references #34 and #35). B) BrdU labeling studies following injury reveal proliferation of murine cardiomyocytes in the border region of the adult mouse heart. Arrowheads mark BrdU labeled cardiomyocyte nuclei, which are green and are immunostained with α -actinin, which is red. Adapted from Naseem et al. Physiol Genomics, 2007.³⁸ Am Physiol Soc, used with permission.

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Figure 3.

The adult heart is capable of limited regeneration. A) Schematic highlighting the potential of stem cells for self renewal and their capacity to generate cardiac progenitors or cardiomyocytes. B) Schematic highlighting the possibility that extracardiac stem or progenitors can be recruited from the bone marrow, the endothelial lineage, skeletal muscle or other sources to further participate in myocardial regeneration and repair. C) Table outlining several cardiac stem or progenitors that have been identified and their characterization. E: embryonic expression; A: adult expression.

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Selected examples of somatic adult stem cell populations that reside in adult, mammalian tissues.

Adult stem cell population	Niche (Tissue)	Derivatives	Reference
Bronchioalveolar Stem Cells	Lung	clara- and AT2-like cells	18-19
Bulge stem cells	Hair Follicle	hair follicles, sebaceous glands, and epidermis	20–21
Epithelial stem cells	Lining of Digestive Tract	villus columnar, mucous, entero-endocrine, and paneth cells	22–23
Hematopoietic stem cells	Bone Marrow	lymphoid (B cell, T cell, natural killer cell, and lymphoid dendritic cell lineage) and myeloid (all other lineages) blood cells	24–25
Neural stem cells	Ependymal Cell Layer	neurons, astrocytes, and oligodendrocytes	26–27
Oval cells	Liver	hepatocytes and biliary epithelium	4–6
Satellite cell	Skeletal muscle	skeletal muscle	7, 28–29

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Table 2

Results of selected cell therapy trials in patients with ischemic cardiomyopathies. AMI: acute myocardial infarction, STEMI: ST elevation myocardial infarction, CABG: coronary artery bypass surgery, BM: bone marrow, IC: intracoronary, IM: intramuscular, LV: left ventricle, LVESV: left ventricular end systolic volume, LVEDV: left ventricular end diastolic volume, MRI: magnetic resonance imaging, ECHO: echocardiography

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Study	No of Subjects	Design	Patients	Type of Cells	Mode of Delivery	Follow -up	Parameters measured	Modality	Improvement	Reference
TOP CARE- AMI	59	Randomized	AMI	Circulating progenitor cells Vs. BM mononuclear cells	IC	12 months	LV Function LV ESV	LV Angiogram MRI	All parameters improved	72
BOOST	60	Randomized	STEMI	BM monouclear cells	IC	18 months	Infarct Size LVEF	Stress ECHO MRI	No change	73
REPAIR-AMI	57	Randomized	STEMI	BM monouclear cells	IC	12 months	LVEF LV EDV LV ESV	MRI	Improvement in LVEF, LV EDV and LV ESV	74
ASTAMI	100	Randomized	STEMI	BM monouclear cells	IC	3 years	LVEF Infarct Size Exercise Capacity	ECHO MRI	No change in LVEF or infarct size. Slight improvement in exercise capacity.	75
MAGIC	120	Randomized	CABG	Autologous skeletal myoblasts	IM	6 months	LVEF LVEDV LVESV	ECHO	No change in LVEF. Improvement in LV EDV and LV ESV with high dose.	76