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Heart Regeneration

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Preface

Heart failure plagues industrialized nations, killing more people than any other disease. Heart failure usually results from a deficiency of cardiomyocytes, and a robust therapy to regenerate lost myocardium could help millions of patients a year. Heart regeneration is well documented in lower vertebrates and in developing mammals. After we are born, however, human heart regeneration becomes limited. In this article, Laflamme and Murry review the innate barriers to heart regeneration, the evidence for cardiomyocyte turnover in humans, and current experimental strategies to remuscularize the injured heart using adult and pluripotent stem cells, cellular reprogramming and tissue engineering.

Introduction

Heart regeneration has been intensely investigated, and intensely controversial, for over 150 years¹. In pursuit of this question the heart has been stabbed, snipped, contused, cauterized, coagulated, frozen, injected with toxins, infected and infarcted, studying species ranging from marine invertebrates to horses^{2,3}. Why has this proven to be such a difficult challenge? The heart is clearly one of the least regenerative organs in the body, so if there is a regenerative response, it is small in comparison to that seen in many other tissues, like liver, skeletal muscle, lung, gut, bladder, bone or skin. For most investigators, the question really is about whether there is no regeneration (intrinsically difficult to prove) or regeneration at very low rates (difficult to detect, but possible with sensitive approaches).

This is more than an academic argument. Heart failure is a burgeoning public health problem, and some predict this will reach epidemic proportions as our population ages. Cardiomyocyte deficiency underlies most causes of heart failure. The human left ventricle has 2–4 billion cardiomyocytes, and a myocardial infarction can wipe out 25% of these in a few hours⁴. Disorders of cardiac overload like hypertension or valvular disease kill cardiomyocytes slowly over many years⁵, and aging is associated with loss of ~1 g of

myocardium per year in the absence of specific heart disease⁶. If the human heart has an innate regenerative response, even a small one, it may be possible to exploit this therapeutically and enhance the heart's function. This fundamental motivation has kept investigators pursuing rare events for more than a century.

Over the last 15 years, researchers have taken a more interventional approach to the injured heart, creating the field of cardiac repair. The ultimate goal of cardiac repair is to regenerate the myocardium after injury to prevent or treat heart failure. This interdisciplinary field draws from advances in stem cells, developmental biology, biomaterials and others areas, in an attempt to create new myocardium that is electrically and mechanically integrated into the heart. Cardiac repair has moved rapidly from studies in experimental animals into clinical trials involving thousands of patients. In the following sections, we will review the evidence for heart regeneration in animal models and humans. We discuss the status of research using adult and pluripotent stem cells for cardiac repair in experimental animals, and we explore the promises and problems of cellular reprogramming and tissue engineering. Clinical trials will be covered only briefly due to space limitations, so we refer interested readers to recent reviews on this topic^{7,8}.

Heart Regeneration in Lower Vertebrates

Unlike humans, many lower vertebrates readily regenerate limbs, appendages and internal organs after injury. There is a long history of research in amphibian heart regeneration⁹; but, more recently, the zebrafish has proven to be a particularly useful model, given its substantial regenerative capacity and amenability to genetic manipulation¹⁰. As was first shown by Poss and colleagues in 2002, the zebrafish heart fully regenerates following the surgical amputation of the cardiac apex, an injury which corresponds to a loss of approximately 20% of the total ventricular mass¹⁰. In the low pressure fish heart, this large wound is effectively sealed by an initial fibrin clot, which is gradually replaced by de novo regenerated heart tissue rather than scar¹⁰⁻¹³.

Not surprisingly, this regenerative response involves a substantial amount of cardiomyocyte proliferation. Even at baseline, zebrafish cardiomyocytes show a much higher degree of cell cycle activity than their mammalian counterparts, with approximately 3% of myocytes in the compact myocardium of uninjured adult zebrafish hearts incorporating the thymidine analogue bromodeoxyuridine (BrdU) during a 7-day pulse. At two weeks following amputation of the cardiac apex, the fraction of BrdU-positive myocytes was increased ten-fold, and this parameter remained as high as 20% as late as one month post-injury¹⁰.

Initial experiments suggested that undifferentiated progenitor cells were the principal source of regenerating cardiomyocytes¹¹, but a pair of recent genetic fate mapping studies unambiguously demonstrates that preexisting committed cardiomyocytes are instead the main source^{12,13} (Box 1). The Poss and Izpisua Belmonte groups independently generated transgenic zebrafish in which the cardiomyocyte-specific *cmcl2* promoter drives expression of tamoxifen-inducible Cre recombinase. These animals were then crossed with a reporter line, in which Cre-mediated excision of a loxP-flanked stop sequence induces constitutive GFP expression. In the offspring of this cross, all preexisting cardiomyocytes and their

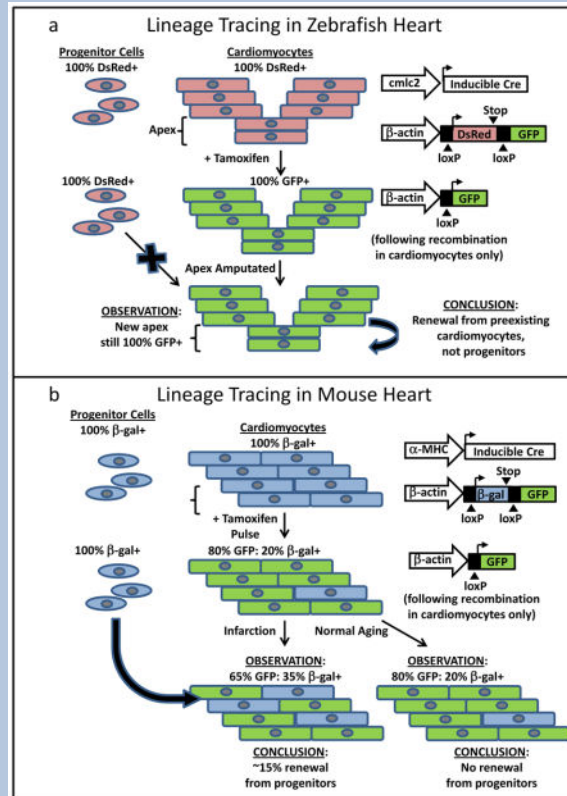
progeny can be induced to express GFP following treatment with tamoxifen. If the regenerated myocardium were derived from undifferentiated progenitor cells, the new ventricular apex should be GFP-null. Instead, both groups found the vast majority of the newly regenerated cardiomyocytes were GFP-positive^{12,13}. Thus, heart regeneration in zebrafish is principally mediated by the proliferation of preexisting cardiomyocytes, rather than generation of new cardiomyocytes from stem cells.

Box 1

Genetic Fate Mapping in Heart Regeneration

Genetic fate mapping has proven to be an invaluable tool for dissecting the mechanisms of endogenous cardiac repair in model organisms, and a number of labs have employed an elegant strategy based on the conditional Cre-lox system, which allows both temporal and cell-type specific control of reporter expression. **a**, Following amputation, the apex of the zebrafish heart can fully regenerate. To determine the source of the newly proliferating cardiomyocytes that underlie this regeneration, the Poss group created a zebrafish strain carrying two transgenes: one in which the cardiomyocyte-specific *cmcl2* promoter drives expression of tamoxifen-inducible Cre recombinase, and a second transgene in which Cre recombinase-induced excision of loxP-flanked stop sequences, causing a switch from constitutive DsRed to constitutive GFP expression¹². Thus, when this transgenic animal was pulsed with tamoxifen, all of its cardiomyocytes and their descendants became indelibly marked with GFP. In contrast, cardiomyogenic progenitor cells should have remained DsRed+/GFP-null, because the cardiomyocyte-specific *cmcl2* promoter would not be active in these undifferentiated cells. If progenitors later contributed to cardiomyocyte renewal following injury, one would expect those myocytes to also be DsRed+/GFP-null. Instead, the new apical myocardium was GFP+, indicating that heart regeneration in the zebrafish results from the expansion of preexisting cardiomyocytes, not the recruitment of cardiomyogenic precursors. The *Izpisúa Belmonte* group independently reached the same conclusions using a similar experimental design¹³. **b**, An analogous genetic fate mapping approach was employed by Hsieh et al²² to investigate the mechanisms of cardiac regeneration in mammalian hearts. These authors generated a double-transgenic mouse in which the cardiomyocyte-specific α -MHC promoter drives tamoxifen-inducible Cre recombinase, and Cre-mediated excision of loxP flanked stop sequences induces a switch from constitutive β -galactosidase to constitutive GFP expression. Following tamoxifen treatment, ~80% of the cardiomyocytes in this animal became GFP+, while 20% remained β -galactosidase+ but GFP-null. As in the analogous zebrafish experiment, any progenitor cells should remain GFP-null following the tamoxifen pulse. During normal aging for up to 1 year post-tamoxifen, the ratio of GFP+ to β -galactosidase+ cardiomyocytes remained fixed at 80:20%, implying no significant cardiomyocyte renewal by unlabeled progenitors. However, following infarction, the ratio of GFP+ to β -galactosidase+ shifted to ~65:35% in the peri-infarct zone, indicating that newly differentiated cardiomyocytes (β -galactosidase+ and GFP-null because they had not undergone Cre-mediated recombination) had been recruited from the progenitor pool. Hence, in the mouse, the

small amount of regeneration that occurs following injury involves the cardiac induction of progenitor cells.



Limited Regeneration in Rodent Hearts

While lacking the remarkable regenerative capacity of the zebrafish heart, postnatal mammalian hearts also undergo some degree of cardiomyocyte renewal during both normal aging and disease. Despite all of the recent attention by the field, this is not a new concept. Extremely low but detectable levels of cardiomyocyte cell cycle activity have been reported in rodent studies dating back to the 1960's^{3,14}. Capturing the rare dividing cardiomyocytes present in mammalian hearts is technically challenging, but recent work has taken advantage of the greater specificity and throughput afforded by transgenic mouse models. For example, the Field group used transgenic mice in which the cardiomyocyte-specific α -myosin heavy chain (α -MHC) promoter drives expression of nuclear localized expression of β -galactosidase¹⁵. This convenient readout allowed them to screen $>10^5$ cardiomyocyte nuclei for incorporation of radiolabeled thymidine and they found labeling indices of 0.0006% and 0.0083% in adult ventricular myocytes from intact and infarcted hearts^{14,15}.

While such proliferative indices are depressingly small, they beg the question whether such phenomena could be therapeutically augmented. Proof of concept for this approach has come from transgenic mice with cardiomyocyte-restricted overexpression of the cell-cycle activator cyclin D2, as these animals show reduced scar and improved mechanical function following myocardial infarction¹⁶. Other efforts to enhance the proliferation of adult

cardiomyocytes by manipulating oncogenes or cell cycle regulators have proven less consistent in improving outcomes post-infarction (for a comprehensive review see ref¹⁷). Pharmacological enhancement of cardiomyocyte cell cycle activity would be more practical clinically than gene therapy, and the signaling molecules periostin¹⁸, fibroblast growth factor-1¹⁹, and neuregulin-1²⁰ have all been reported to act as mitogens for adult ventricular myocytes and to exert beneficial effects on cardiac structure and function post-infarction. (It should be noted that a more recent study has called these effects into question in the case of periostin²¹.) The Kühn group's recent study of the mitogenic effects of neuregulin-1 particularly captured the attention of the field, as they reported simple systemic injection of this factor sufficed to enhance infarct scar shrinkage and improve mechanical function²⁰. The effect on cell cycle required intact ErbB4 receptors in cardiomyocytes and appears to stimulate mononucleated, but not binucleated, cardiomyocytes to divide. Although this intriguing result awaits independent confirmation, it suggests a straightforward approach to enhancing ventricular repair through administration of recombinant growth factors.

Most of the above studies focused on proliferation of existing cardiomyocytes, but they were not designed to detect cardiomyocytes formed from progenitor cells. To determine the whether such progenitors contribute to cardiomyocyte renewal, Hsieh et al²² performed an elegant genetic fate mapping experiment, akin to those previously described for the zebrafish model, in which cardiomyocytes were indelibly labeled following a tamoxifen pulse (Box 1). This system allowed the authors to distinguish between cardiomyocyte renewal from preexisting (and therefore fluorescently-labeled myocytes) and unlabeled progenitors. Interestingly, Hsieh et al found no significant contribution by such progenitors during normal aging (up to one year post-tamoxifen). However, they did observe a reduction in the fraction of labeled cardiomyocytes following infarction, indicating dilution by unlabeled progenitors. When combined with the findings that cardiomyocyte proliferation is vanishingly small in both normal and injured rodent hearts, these data suggest that the limited endogenous reparative mechanisms in the adult mammalian heart operate quite differently from those in zebrafish and depend more on replenishment by cardiomyogenic progenitors than myocyte proliferation.

An intriguing recent report by Sadek and colleagues²³ suggests that these differences between mammalian and fish hearts do not necessarily apply earlier in development. Borrowing approaches from the zebrafish model, these authors resected the left ventricular apex of 1-day-old neonatal mice and observed a brisk regenerative response reminiscent of that in the adult zebrafish. By 3 weeks post-injury, the defect had been replaced by normal myocardial tissue, which, by 8 weeks, showed normal contractile function. Genetic fate-mapping studies indicated that this regeneration was mediated by the proliferation of preexisting cardiomyocytes, again as in the zebrafish. Interestingly, this regenerative capacity was not observed in 7-day-old mice, suggesting that its loss may coincide with cardiomyocyte binucleation and reduced cell cycle activity. Nonetheless, in addition to representing a surgical *tour de force*, this study indicates that zebrafish-like regenerative mechanisms are latent in mammalian hearts. It also provides a new genetically tractable model for dissecting the blocks to these mechanisms in the adult.

The Evidence for Human Heart Regeneration

Before addressing whether new cardiomyocytes are generated in the human heart after injury, it is instructive to review a few points of normal cardiac growth and adaptation to workload. These are outlined in Box 2. In brief, most human cardiomyocyte nuclei are polyploid by the onset of puberty²⁴. In response to pathological workloads (hypertension, valve disease, post-infarction overload etc.), human cardiomyocytes commonly reinitiate DNA synthesis without nuclear division^{24,25}. This further increases their nuclear ploidy, reaching levels as high as 64C. Unlike rodent cardiomyocytes²⁶, most human cardiomyocytes appear to remain mononucleated throughout life²⁷. Thus, DNA synthesis is common in the adult human heart, but this cannot be equated with cardiomyocyte proliferation without accounting for the process of polyploidization.

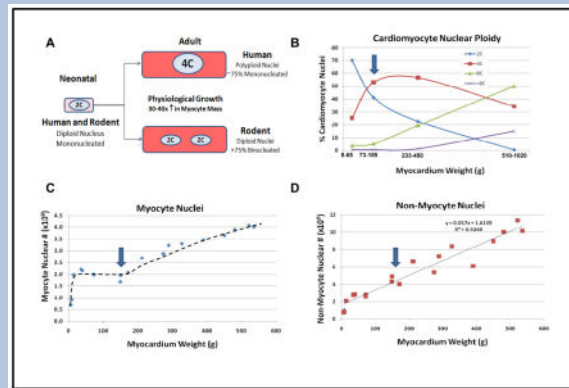
Box 2

Nuclear dynamics during human heart growth

A. Fetal hearts in humans and rodents grow through proliferation of mononucleated cardiomyocytes with diploid nuclei. After birth, rodent cardiomyocytes withdraw from the cell cycle within the first few days. Human cardiomyocytes appear to proliferate for the first few months after birth, after which replication slows markedly. The cells then undergo a period of physiological growth, increasing myocyte size by 30–40 fold. It seems that most myocytes cannot grow this much with a single diploid genome, but this problem has been solved differently in different species. Most rodent cardiomyocytes undergo a final round of DNA replication followed by nuclear division without cytokinesis, resulting in >75% binucleated cells with a normal diploid (2C) content of DNA²⁶. In humans and other primates, most cardiomyocytes undergo a final round of DNA replication, without nuclear division or cytokinesis, resulting in mononucleated cells²⁷ with tetraploid (4C) or higher DNA content³¹. (Aficionados will note that estimates of human cardiomyocyte binucleation rates range from 25% in enzymatically dispersed fresh tissue²⁷ to >60% in KOH-digested formaldehyde-fixed tissue⁹⁹. We favor the 25% value, because the harsh KOH digestion may selectively eliminate smaller mononucleated myocytes.) Interestingly, pacemaker cells of the sinoatrial and atrioventricular nodes remain small and diploid throughout life¹⁰⁰. **B.** Cytofluorometric analysis of human cardiomyocyte nuclear DNA content as a function of myocardial weight after carefully removing valves, vessels and fat. (Hearts were grouped by weight into 4 bins for the purpose of clarity.) The percentage of diploid cardiomyocyte nuclei falls steadily as myocardial mass increases. Studies of pediatric hearts indicate that polyploidization occurs in the pre-adolescent growth phase from 8–12 years of age. Tetraploid nuclei (4C) are most common in the adult heart. During cardiac hypertrophy octaploid nuclei (8C) become most common, with a substantial number of 16C or higher polyploidy. These data demonstrate that human cardiomyocytes have a substantial capacity for DNA replication. Blue arrow: upper limits of normal for human myocardial weight. **C.** Morphometric analysis of human cardiomyocyte nuclear number during physiological and pathological growth. The normal adult number of 2 billion cardiomyocyte nuclei is reached by ~2 months age. Cardiomyocyte nuclear number

remains steady during physiological hypertrophy, but when heart weight exceeds ~450 g (myocardial weight ~210 g) there appears to be a linear increase in cardiomyocyte nuclear number with increasing cardiac mass. Since human hearts do not change nuclear number with hypertrophy, this is evidence for generation of new cardiomyocytes, either from pre-existing cardiomyocytes or from stem cells. Blue arrow: upper limits of normal for human myocardial weight. **D.** Non-myocyte nuclei increase linearly with increasing myocardial mass, indicating proliferation of these cells accompanies all phases of cardiac growth. Blue arrow: upper limits of normal for human myocardial weight.

The data in panels B–D are derived from Adler and Costabel³¹. The trend line in panel C is hand-drawn and for illustration purposes only.



Historically, regenerative responses have been detected either by macroscopic regrowth of the tissue or by the microscopic presence of mitoses. Macroscopic regeneration of the heart clearly does not occur. Mitosis occupies only ~2% of the cell cycle, so one would predict that these would be difficult to quantify meaningfully. Experiments have borne this out, with some investigators reporting no mitotic figures after injury, and others reporting rare (and potentially abnormal) mitotic figures around the injured site^{28–30}. One factor contributing to these discrepancies has been the inherent difficulty in recognizing cardiomyocyte nuclei in conventional histological sections¹⁴. Many of the published images are persuasive for cardiomyocyte mitosis and provide important evidence that this can occur in humans. On the other hand, extrapolation to organ turnover rates from such low numbers is perilous.

A number of investigators have taken the seemingly straightforward approach of counting the number of cardiomyocytes in the human heart during normal and pathological growth. As anyone who has attempted a task like this knows, however, it is surprisingly difficult and requires multiple assumptions. Adler et al. have the most extensive body of work on this topic^{24,25,31,32}. Using a combination of meticulous dissection, histopathology, biochemical measurements of tissue DNA content, and fluorescent analysis of individual nuclear DNA content, they reported that from 50–350 g, cardiomyocyte nuclear number is steady at ~2 billion. Beyond that, there is a linear increase in nuclear number with increasing heart weight, reaching 4 billion nuclei in hypertrophied hearts weighing 700–900 g. Non-myocytes (e.g. fibroblasts and vascular cells) increase linearly with heart weight throughout life. If correct, these data imply cardiomyocyte renewal occurs during pathological

hypertrophy. An important caveat is that the assignment of cardiomyocyte vs. non-myocyte nuclear identity was based on size and morphology of isolated nuclei. Since we know that postnatal growth and pathological hypertrophy are accompanied by increases in nuclear ploidy (hence, size), it is possible that diploid myocyte nuclei in smaller hearts were mistakenly classified as non-myocyte nuclei. On the other hand, increases in cardiomyocyte nuclear number in pathological hypertrophy were also reported by others³³, using histological sections where cardiomyocyte nuclei can be more readily identified.

Two studies have attempted to use more direct means to measure cardiomyocyte DNA synthesis rates in human hearts. Bergmann et al³⁴ used a clever approach based on the worldwide pulse of ¹⁴C that occurred during atmospheric testing of nuclear weapons during the Cold War. The atmospheric ¹⁴C became incorporated into plants and entered the human food chain, labeling the DNA of dividing cells. After the Limited Nuclear Test Ban Treaty, atmospheric ¹⁴C levels dropped rapidly. This provided pulse-chase conditions that can be used to “birth date” cells, simply by identifying when atmospheric ¹⁴C levels match those of the DNA. As expected, non-myocytes in the heart were substantially younger than the patient, with ~18% turnover per year and a mean age of only 4 years. Interestingly, DNA from isolated cardiomyocyte nuclei (sorted by nuclear troponin staining) also was somewhat younger than the patient, though not nearly as young as that from non-myocytes. As indicated above, before one can infer cell division, it is essential to rule out a contribution from polyploidization. To do this, the authors sorted cardiomyocyte nuclei by their DNA content, and analyzed only the subset with a diploid DNA content. The diploid cardiomyocyte nuclei also were younger than the patient, providing good evidence for cardiomyocyte division. Mathematical modeling suggested that cardiomyocyte renewal was age-dependent, with ~1% of cardiomyocytes being renewed per year at age 20, and 0.4% at age 75. Based on these kinetics, ~45% of cardiomyocytes would be predicted to be renewed over a normal human life span, while 55% would be cells persisting since birth.

Kajstura et al³⁵ took advantage of an interesting “natural” medical experiment to study cardiac DNA synthesis rates. They examined hearts from cancer patients who were treated with the thymidine analog, iododeoxyuridine (IdU). This agent is incorporated into nascent DNA, where it sensitizes cells to radiation therapy. The IdU was given as bolus injections or multi-week infusions, and the time between treatment and death ranged from 7 days to 4.3 years. Using immunohistochemistry to detect the IdU signal and identify cardiomyocytes, they found remarkably high cardiomyocyte DNA labeling rates, ranging from 2.5% to 46%. No IdU staining was found in control hearts not exposed to the radiosensitizer. Mathematical modeling suggested that cardiomyocytes turn over at a rate of 22% per year, compared to 20% for fibroblasts and 13% for endothelial cells. Furthermore, they reported that 83% of cardiomyocyte nuclei were diploid, suggesting that this turnover reflects cell division, not increased ploidy.

It is hard to reconcile these two studies, which differ by nearly 50-fold in their estimates of cardiomyocyte turnover. One important difference appears related to higher cardiomyocyte DNA synthesis activity in cancer patients treated with IdU. Kajstura et al reported 3-fold lower DNA synthesis rates (based on Ki-67 immunolabeling) in control hearts vs. the IdU-treated patients. Neither study adequately rules out a contribution from DNA repair, which

can masquerade as DNA replication in these assays. This is of particular concern in cancer patients receiving radiation plus a radiosensitizer. Kajstura et al.³⁵ suggested that only senescent cardiomyocyte nuclei contain troponin, which could bias Bergmann's turnover studies toward low proliferation. A follow up paper from Bergmann et al.³⁶ provided evidence that nearly all cardiomyocyte nuclei were identified by troponin staining, however. Additionally, Kajstura's findings contradict two well accepted principles. First, they suggest that >80% of cardiomyocyte nuclei are diploid, whereas virtually all other investigators report these nuclei are polyploid. If so, polyploidization could underlie their high estimates of DNA synthesis. Second, they conclude that cardiomyocytes are as proliferative as non-myocytes, whereas virtually all other investigators find orders of magnitude greater proliferation in non-myocytes. Indeed, they reported IdU incorporation by cardiomyocytes at rates (2.5–46%) approaching those reported for the sarcomas targeted by IdU (50–70%)³⁷. The heart does not proliferate like a sarcoma, so these cardiac IdU incorporation estimates must be too high.

Taken together, these human studies provide strong evidence for plasticity in the adult human heart. There is extensive morphometric evidence for DNA synthesis and an increase in cardiomyocyte number in diseased human hearts. Cardiomyocyte division or generation from progenitor cells probably occurs in the human heart, but it appears to be a very slow process.

We need better tools to study this process quantitatively, and better ways to model it, if we hope to exploit it therapeutically.

Stem Cells and Cell Therapy

Stem cell biology is one of the fastest moving areas of biomedical research, and amongst all solid organs, the heart has one of the most active regeneration research programs. The field can be conceptually organized into work involving endogenous and exogenous cells. The many exogenous cell types can be further divided into pluripotent (ESCs, iPSCs etc.) and "adult" cells of more limited potential (circulating progenitors, resident cardiac progenitors, and cells native to other tissues. Here, we will focus on cells closest to clinical trials and those for which there are the most reliable data.

Cardiac Progenitor Cells (CPCs)

Multiple investigators reported resident populations of CPCs in postnatal hearts, which were identified using a variety of approaches, including expression of surface markers like c-Kit or Sca-1 (note that Sca-1 apparently has no human ortholog), or physiological properties like the ability to efflux fluorescent dye or form multicellular spheroids (reviewed in^{38,39}). Initially, it seemed there was little overlap amongst CPCs identified by these different approaches, and some scientists suggested that multiple populations of CPCs exist. More recent studies indicate shared markers among once-distinct populations or different stages of maturation in the same line of cells^{40,41}, so the field may be converging.

Most extensively studied are CPCs expressing the receptor tyrosine kinase c-Kit. In the adult, c-kit is expressed by hematopoietic stem cells, mast cells and other mature circulating

cells, telocytes (formerly known as the interstitial cells of Cajal) and thymic epithelium. During development, c-kit is also expressed by immature endothelial cells and cardiomyocytes⁴². Small round cells expressing c-Kit have been identified in the perivascular compartment of the adult heart, and their abundance increases in human heart failure⁴³. Following isolation from rat and human hearts, c-Kit+ cells have been reported to give rise to cardiomyocytes, smooth muscle cells and endothelial cells. Some studies indicate that, when transplanted, c-Kit+ cells induce large-scale regeneration of myocardial infarcts and contribute to the formation of new myocardium and vessels⁴⁴, while others suggest smaller-scale regeneration⁴⁵. Based on these data, a clinical trial is currently underway testing the safety and feasibility of autologous c-Kit+ cells as an adjunctive treatment for patients undergoing coronary bypass surgery (NCT00474461).

Not all studies with c-Kit CPCs gave robustly positive results. In studies with genetic readouts for lineage tracing and differentiation state, c-Kit cells from the adult heart have not been found to differentiate into cardiomyocytes in vitro or following transplantation into infarcted hearts⁴⁶. Another study using transgenic reporter mice found no evidence that endogenous c-kit+ cells differentiate into cardiomyocytes, though re-expression of c-kit in pre-existing cardiomyocytes was identified⁴². Others point out that myocardium, like all solid tissues, contains mast cells. Mast cells are small, round cells that reside in clusters in the perivascular space, they strongly express c-Kit, and their numbers go up in failing hearts⁴⁷. Studies in humans suggest that 90–100% of all of the cardiac c-Kit+ cells are actually mast cells. On the other hand, expansion in culture appears to select for c-Kit+ cells that lack mast cell markers, indicating that freshly isolated cells and cultured cells are different populations⁴⁸.

The other CPC population in clinical trials is the so-called cardiosphere forming cell (CFCs). CFCs are isolated based on their ability to migrate out of cultured cardiac tissue fragments and form spheroids in suspension cultures^{49,50}. As one might predict, this yields a mixture of cells, some of which express stem cell markers like c-Kit, and others that appear to come from the stromal-vascular compartment. CPCs have been reported to give rise to cardiomyocytes in vitro and in vivo after transplantation and to enhance cardiac function post-infarction⁵⁰. Based on these data, a clinical trial of autologous CPCs has been initiated for patients with recent myocardial infarctions (NCT00893360). The “stemness” of CPCs has recently been questioned as well, with some suggesting that these cells are principally cardiac fibroblasts and that CPC-derived cardiomyocytes are actually contaminants derived from the original tissue⁵¹.

Thus, while the study of CPCs is one of the most exciting new areas in cardiac research, it is also one of the most controversial. Most of the work has focused on cell culture and transplantation, driven by the clinical need for cardiac repair. We know almost nothing about the *endogenous* behavior of CPCs, however. What is the role of these cells in development, homeostasis, aging and reaction to injury? The field needs models that permit unambiguous tracing of CPC lineage and phenotype without resorting to transplantation or cell culture (e.g. Box 1).

Bone Marrow Cells

Considerable interest in marrow-derived cells for cardiac repair was prompted by reports that hematopoietic stem cells transdifferentiated into cardiomyocytes⁵². Subsequent studies showed that hematopoietic stem cells do not form cardiomyocytes but rather become mature blood cells after transplantation^{53,54}. Nevertheless, animal studies show improvements in ventricular function when hematopoietic cells are administered post-infarction, implicating paracrine signaling as the major mechanism of action.

Work with marrow-derived stromal cells (MSCs) has followed a similar trajectory. MSCs were originally reported to transdifferentiate into cardiomyocytes⁵⁵ but are now thought to exert their main actions through paracrine release of cytokines⁵⁶. Interestingly, most MSCs die within days to weeks after transplantation into infarcts, yet beneficial effects can be seen long term, suggesting a critical window of time post-infarction for their action. MSCs likely operate through multiple mechanisms, but considerable evidence points toward their regulation of the Wnt pathway. MSCs secrete antagonists of canonical Wnt ligands, such as secreted frizzled related protein 2⁵⁶. Blocking production of Wnt antagonists limits the beneficial effects of MSCs. Hatzistergos et al.⁵⁷ recently reported that administration of MSCs to pig infarcts stimulated endogenous CPCs to contribute to repair of the infarct. Further identification paracrine mediators may allow the development of simpler, cell-free treatments based on proteins or small molecules.

Clinical trials have mostly focused on delivery of bone marrow mononuclear cells via the coronary circulation. It should be emphasized that >99.9% of bone marrow mononuclear cells are not stem cells but rather are committed, though immature, granulocytes or other hematopoietic lineages. These trials indicate that delivery of bone marrow derivatives through the coronaries is feasible and safe, but the benefits are currently modest. MSCs are also in clinical trials (e.g. NCT00587990). There are few published results with these cells, but interestingly, Chen et al. reported one of the strongest treatment effects seen to date in cardiac repair (a 14% improvement in ejection fraction) following the intracoronary administration of large numbers of autologous MSCs⁵⁸. Hare et al. administered allogeneic MSCs intravenously to patients within 10 days of infarction, and they reported that the cells were well tolerated and were associated with decreased arrhythmias and an improvement in some indices of contractile function⁵⁹.

Taken together, the best current evidence indicates that bone marrow cells do not work via directly differentiating into new cardiomyocytes. Rather, the preponderance of evidence indicates that the marrow cells elaborate signals that control the response of cells native to the myocardium, and thereby regulate healing. While many view this as a novel aspect of stem cell biology, students of pathology will recognize this that phenomenon also fits under a more familiar heading: inflammation. We find it useful to consider the participation of marrow derivatives in cardiac repair as part of the inflammatory response, which has long been known to regulate angiogenesis, cardiomyocyte survival, and left ventricular remodeling post-infarction.

Pluripotent Stem Cells

Many adult stem cell types are unable to generate large numbers of unambiguous cardiomyocytes. This limitation does not apply to embryonic stem cells (ESCs) or their more recently developed, “man-made” counterpart, induced pluripotent stem cells (iPSCs). Since both ESCs and iPSCs can be propagated indefinitely, while still retaining the capacity to differentiate into virtually all cell types, they represent a potentially inexhaustible supply of human cardiomyocytes. Our current thinking about how cardiomyocytes arise from ESCs is shown in Figure 1. Human ESC-derived cardiomyocytes express early cardiac transcription factors (e.g. Nkx2.5), as well as the expected sarcomeric proteins, ion channels, connexins and calcium-handling proteins (Figure 2). They exhibit functional properties similar to those reported for cardiomyocytes in the developing heart, and they undergo comparable mechanisms of excitation-contraction coupling and neurohormonal signaling^{60–63}. Human ESC-derived cardiomyocytes have been more intensively studied, but the best available data suggests that human iPSC-derived myocytes have a very similar phenotype^{64,65}. Importantly, cardiomyocytes from either pluripotent stem cell type are immature and so lack the expression profile, morphology and function of adult ventricular myocytes.

While the cardiac potential of ESCs and iPSCs is undisputable, their unique origin and pluripotency does present a new set of challenges. ESCs are derived from the inner cell mass of preimplantation stage blastocysts⁶⁶, and this contributes to the ethical controversy surrounding their use. Moreover, ESC-based therapies will be allogeneic and require immunosuppression. iPSCs were originally generated by the reprogramming of adult somatic cells (e.g. dermal fibroblasts) via the forced expression of up to four stem cell-related transcription factors^{67–69}. As such, their derivation does not involve the destruction of embryos, and they could potentially be used in autologous cell therapies. Nonetheless, “first-generation” iPSCs were problematic because the reprogramming factors were introduced using integrating viruses, raising concerns about neoplastic transformation. More recently, there have been a variety of refinements to iPSC generation that should reduce or eliminate this risk, including the use of episomal gene delivery, excisable transgenes, synthetic mRNA or cell-permeant recombinant proteins (see ⁷⁰ and references therein). Perhaps most exciting, a number of small molecules have been shown to greatly enhance the efficiency of reprogramming⁷¹, inviting speculation that iPSCs may be generated using such factors alone in the reasonably near future. Additional work will be required to more precisely define the phenotype and maturation potential of cardiomyocytes derived from iPSCs generated by these various methods.

Another concern related to the clinical application of pluripotent stem cells is their capacity to form teratomas following transplantation⁷². To overcome this, the field must develop methods to enrich ESC and iPSC derivatives for cardiomyocytes or other useful cell types (e.g. endothelial, smooth muscle, and stromal cells). Historically, ESCs have been differentiated via culture in three-dimensional aggregates, so-called embryoid bodies (EBs), in medium containing a high percentage of fetal calf serum. This method is poorly cardiogenic, and differentiated human EBs are typically comprised of <1% cardiomyocytes⁷³. More recently, our group and others have used insights from

developmental biology to devise better controlled approaches in which human ESCs and iPSCs are treated with defined factors, resulting in highly enriched populations of cardiomyocytes^{74–76}. A common theme with such protocols has been the manipulation of cardioinductive molecules belonging to the transforming growth factor- β superfamily, specifically, activin and the bone morphogenetic proteins (BMPs). For example, our group has reported a protocol involving the serial application of activin A and BMP-4, which reliably yields ~60–80% human ESC-derived cardiomyocytes in large-scale preparations (~ 10^8 – 10^9 total cells)^{74,77}. Additional refinements are possible by manipulating the Wnt/ β -catenin signaling pathway⁷⁸, which mediates biphasic effects on ESC cardiogenesis, promoting mesodermal induction early, but inhibiting cardiogenesis late⁷⁹.

A complementary approach involves the isolation of mesodermal progenitors with a more restricted potential (e.g. cardiovascular progenitors able to differentiate into cardiomyocytes, smooth muscle cells, and endothelial cells). Such multipotent progenitors have been identified in differentiating ESC cultures based on their expression of transcription factors including *Mesp1*⁸⁰, *Nkx2.5*⁸¹, and *Isl1*^{82,83}. Arguably more useful for eventual clinical application are progenitor populations that can be sorted based on their expression of a cell surface marker, such as the cardiovascular progenitors marked by expression of KDR (also known as Flk-1)⁷⁵. If such cells could be induced to self-renew, they would be potentially very useful for cardiac repair.

Human ESC-derived cardiomyocytes have been shown to engraft in infarcted mouse, rat, guinea pig and porcine hearts (Figure 3), forming islands of nascent, proliferating human myocardium within the scar zone^{74,84,85}. This partial remuscularization was accompanied by beneficial effects on regional and global cardiac function^{74,84}, although some investigators have questioned whether these effects are sustained at late timepoints⁸⁶. An important point is that the mechanism(s) underlying the observed improvements in contractile function remain unresolved. In the aforementioned rodent studies, most of the graft tissue was isolated from the host myocardium by scar tissue, which may prevent synchronous beating. Additionally, these human cells (which fire in vitro at ~50–150 beats per minute (bpm)⁷⁷) may not keep pace with the rapid rate of rats (~400 bpm) and mice (~600 bpm). If they cannot, then the observed salutary effects likely resulted from an indirect, “paracrine” mechanism, like those described above for adult cells. On the other hand, this also implies that an additional increment of benefit is possible following transplantation in a slower-rated recipient, such as a canine or porcine infarct model.

Reprogramming Fibroblasts to Cardiomyocytes

Fifteen years ago, researchers showed that fibroblasts could be “transdifferentiated” into skeletal muscle in vitro or in the injured heart by overexpressing the myogenic transcription factor, *MyoD*⁸⁷. Despite an intensive search by multiple groups, no comparable “master gene” for cardiac muscle was found, and interest in reprogramming waned. Spurred by the discovery of iPSCs, scientists have returned to this field, this time using combinations of transcription factors to reactivate core transcriptional networks of desired cell types. In an attempt to induce cardiac differentiation, Ieda, Srivastava et al⁸⁸ systematically screened 14 cardiac transcription factors for their ability to activate a cardiac-specific transgene (α -

myosin heavy chain promoter driving YFP) in cardiac fibroblasts. The full cocktail activated fluorescence in ~1% of cells. A systematic winnowing yielded three transcription factors (Mef2C, Gata4 and Tbx5) that activated the transgene in 20% of fibroblasts. About 4% of the cells expressed endogenous sarcomeric proteins like cardiac troponin T, and only ~1% showed functional properties like spontaneous beating. Thus, most of the YFP+ cells were only partially reprogrammed, although their global gene expression patterns had shifted markedly from fibroblast toward cardiomyocyte.

While this manuscript was under review, Efe et al⁸⁹ reported a different method of reprogramming mouse embryonic fibroblasts into cardiomyocytes. This group used the so-called Yamanaka factors (Oct-4, Sox2, Klf4 and c-Myc) to initiate reprogramming, but they blocked signaling through the JAK-STAT pathway, which is required for pluripotency in the mouse, and added the cardiogenic factor BMP-4. These modifications yielded minimal generation of iPSCs, but instead activated the cardiac progenitor program and, within 2 weeks, substantial numbers of beating colonies. They reported ~40% of cells expressing cardiac troponin T by 18 days post-induction. The authors attributed their increased efficiency versus that of Ieda et al to the generation of highly proliferative progenitor cells, as opposed to forming cardiomyocytes with low proliferative potential. It should also be noted that Efe et al used mouse embryonic fibroblasts, while Ieda et al principally studied postnatal cardiac fibroblasts.

Reprogramming the scar-forming fibroblast to a cardiomyocyte is intuitively appealing, particularly if it can be done directly in the infarct. In order to succeed clinically, we will need to know how normal these reprogrammed cardiomyocytes are, and the process will need to be much more efficient and transgene-free. Despite some challenges, this is an exciting new avenue of research and could be a game-changer.

Tissue engineering

Tissue engineering refers to growth of 3-dimensional tissues in vitro, with goals of building more biologically relevant models for in vitro study, or building tissues for in vivo regenerative therapy. Most commonly this involves the use of porous, biodegradable scaffolds onto which cells are seeded, but other approaches include casting cells into hydrogels or creating scaffold-free tissues composed only of cells and the matrix they secrete. Synthetic materials have big advantages in manufacturability, typically being easy, cheap and reproducible. Synthetics generally have worse biocompatibility, however, because they cause foreign body inflammatory reactions and, sometimes, release locally toxic degradation products. Bioreactors are often employed in tissue engineering, providing electrical and mechanical conditioning or for delivering nutrients to the tissue via perfusion systems

A major goal in bioengineering is to improve the host response to biomaterials, in essence, to make materials that heal⁹⁰. Surprisingly, the chemical composition of a material does not have a major influence on how the body responds to it. Whether materials are organic or metallic, hydrophobic or hydrophilic, or positively or negatively charged, they all cause similar foreign body reactions. Instead, what the body seems to sense is the surface

topography of a material⁹⁰. When surfaces are smooth, there is intense inflammation and scarring, creating a fibrotic capsule around the implant. If a surface is given a more complex topography, e.g. by creating pores or grooves, there is less inflammation, scarring diminishes and blood vessels grow into the implant. Systematic variation in the topology can “tune” this host response. For example, our tissue engineering group has developed scaffolds with two compartments: cylindrical channels to generate cables of cardiomyocytes, surrounded a network of smaller interconnected pores for stromal and vascular ingrowth⁹¹. Pores are optimally sized to maximize vascularization within the implant and minimize fibrosis around it.

The cardiomyocytes used in tissue engineering have been immature cells derived from young animals or stem cells. In order to take on an adult workload, these cells will need to organize into the cable-like structure of myocardium and hypertrophy more than ten-fold. There is an ongoing debate in tissue engineering whether this maturation should take place before or after transplantation. On the one hand, electrical⁹² and mechanical⁹³ stimulation in vitro enhance hypertrophy, alignment and electromechanical function of rat cardiomyocyte constructs. On the other hand, greater cell differentiation is associated with worsened survival after transplantation⁹⁴, so there is likely a point of diminishing returns. This needs to be explored more experimentally.

One of the big lessons from tissue engineering has come from studies comparing myocyte-only versus mixed-cell constructs (Figure 4). When myocyte-only constructs are transplanted, the tissue survives poorly. When vascular endothelial cells and a stromal cell population are included, the endothelial cells form networks resembling a primitive vascular plexus, and the stromal cells elaborate a provisional matrix that enhances mechanical integrity^{95,96}. Upon transplantation, the endothelial network organizes into a definitive vascular network that connects to the host circulation, bringing blood flow into the tissue several days sooner than would otherwise be seen. Indeed, our group and others have demonstrated markedly improved survivability of “pre-vascularized” human myocardial constructs incorporating vascular and stromal elements over those with cardiomyocytes alone^{95,97}. This indicates there is there is considerable synergy to including vessels and connective tissue elements when engineering tissue.

Tissue engineering has not been as extensively studied as cell transplantation in preclinical disease models, but initial studies are promising. Zimmermann et al.⁹³ prepared constructs of engineered heart tissue from neonatal cardiomyocytes and conditioned these for several days using a cyclic stretch system. The constructs were sutured to the surface of rat hearts that had been infarcted two weeks previously, and they were studied one month after implantation. Hearts receiving the engineered heart tissue had better contractile function, and interestingly, conduction velocities across the infarct were improved, likely because the grafts had electrically connected to the surrounding viable myocardium. Another group reported that patches generated from cardiosphere-derived CPCs can enhance heart function post-infarction⁹⁸, and there are hints that tissue engineering affords larger graft size as well.

Summary and Closing Comments

After more than a decade of furious activity, the science of stem cells appears to be catching up with its promise. Clinical scale preparations of the main cardiac cell types can now be generated, and we are learning the rules of building myocardium and keeping it alive after transplantation. Clinical trials have established techniques for cell delivery, and protocols for establishing feasibility, safety and early stage efficacy in humans are in place. The first patient trials have demonstrated safety with hints of efficacy. So far, so good.

That said, many short and long term challenges remain. In the near term, it will be important to derive the right subtype of cardiomyocyte, for example, ventricular myocytes for repair of an infarction that are free of pacemaker cells. The major challenge facing the adult CPC field is developing protocols with higher yields of definitive cardiomyocytes. Researchers studying pluripotent stem cells need to identify the optimal stage of differentiation and demonstrate that these cells can be used without tumorigenesis. The question of allogeneic vs. autologous cells remains open. While desirable, autologous cells will be more expensive, more variable, and the time needed to expand them precludes their use in any acute setting. Allogeneic cells will provide the only “off the shelf” product, but we need to learn how best to manage the immune response to prevent their rejection. All of these efforts will be advanced by improvements in integration of the graft, including control of vascularization (growth of both arterial conduits and microvasculature), inflammation and scarring.

Further ahead, in situ manipulation of cells within the heart may allow us to control their fates, thereby obviating transplantation. For example, can CPCs be controlled with small molecules or growth factors to enhance their regenerative abilities? Can fibroblasts in infarcts be reprogrammed directly into cardiomyocytes? Given our increasing ability to control the fates of cells and tissues, the debate over whether the heart is intrinsically “terminally differentiated” seems anachronistic, for our hearts do not exist apart from the people who know how to manipulate them. It is more useful to ask what we can do to best promote cardiac regeneration, and then do it.

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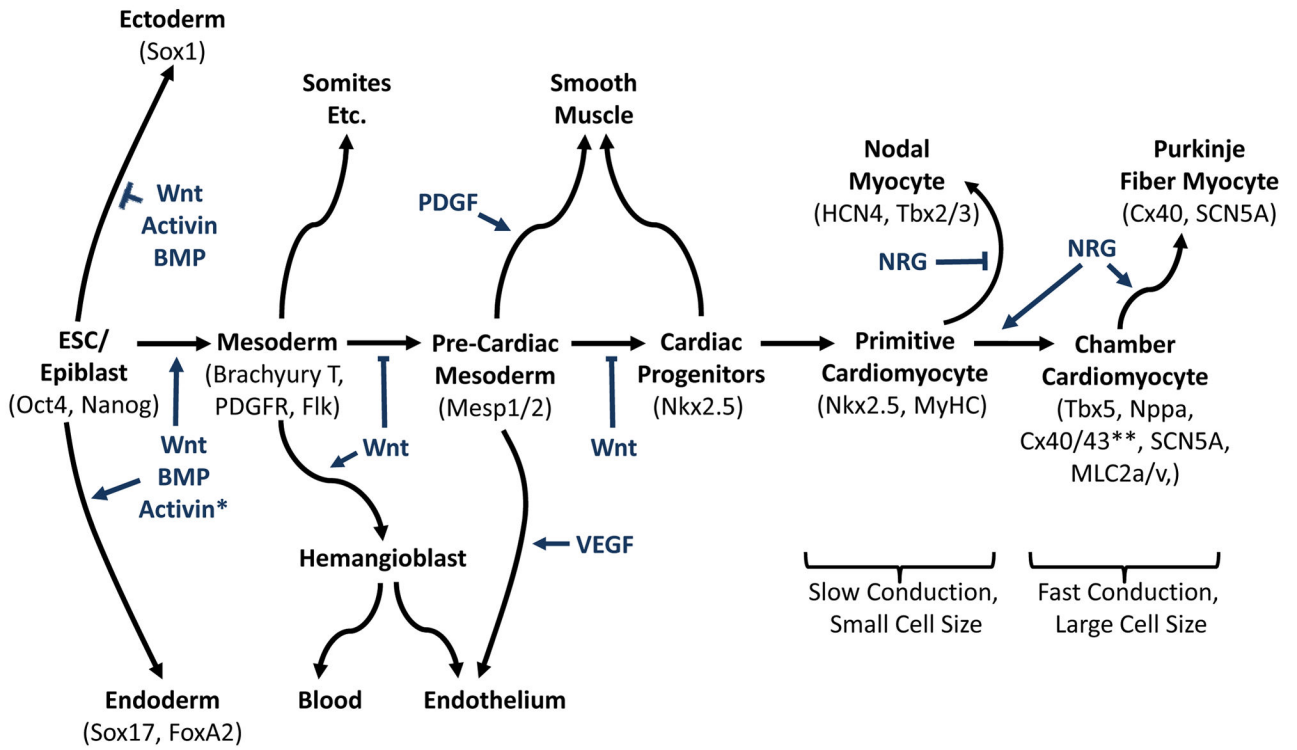


Figure 1. Cardiovascular lineages during embryonic development and embryonic stem cell differentiation

Cardiac differentiation from ESCs closely mimics cardiac development in the embryo. In either case, the specification of the cardiovascular lineages involves a transition through a sequence of increasingly restricted progenitors, proceeding from a pluripotent state to mesoderm and then to cells committed to cardiovascular fates. Growth factors that regulate fate choices are listed at branch points, and key transcription factors and surface markers for each cell state are listed under the cell types. These growth factors are useful for directing the differentiation of ESCs, while the markers are useful for purifying cells at defined developmental states. Primitive cardiomyocytes in the embryonic heart tube and nodal/pacemaker cells exhibit slow electrical propagation and a small cell size. In contrast, the eventual specification of working atrial and ventricular cardiomyocytes is accompanied by more rapid conduction, ion channel remodeling and increased cell size. While the field has made considerable progress toward the elucidation of early events during cardiogenesis, a better understanding of how pacemaker vs. chamber-specific cardiac subtypes are formed is required for clinical applications.

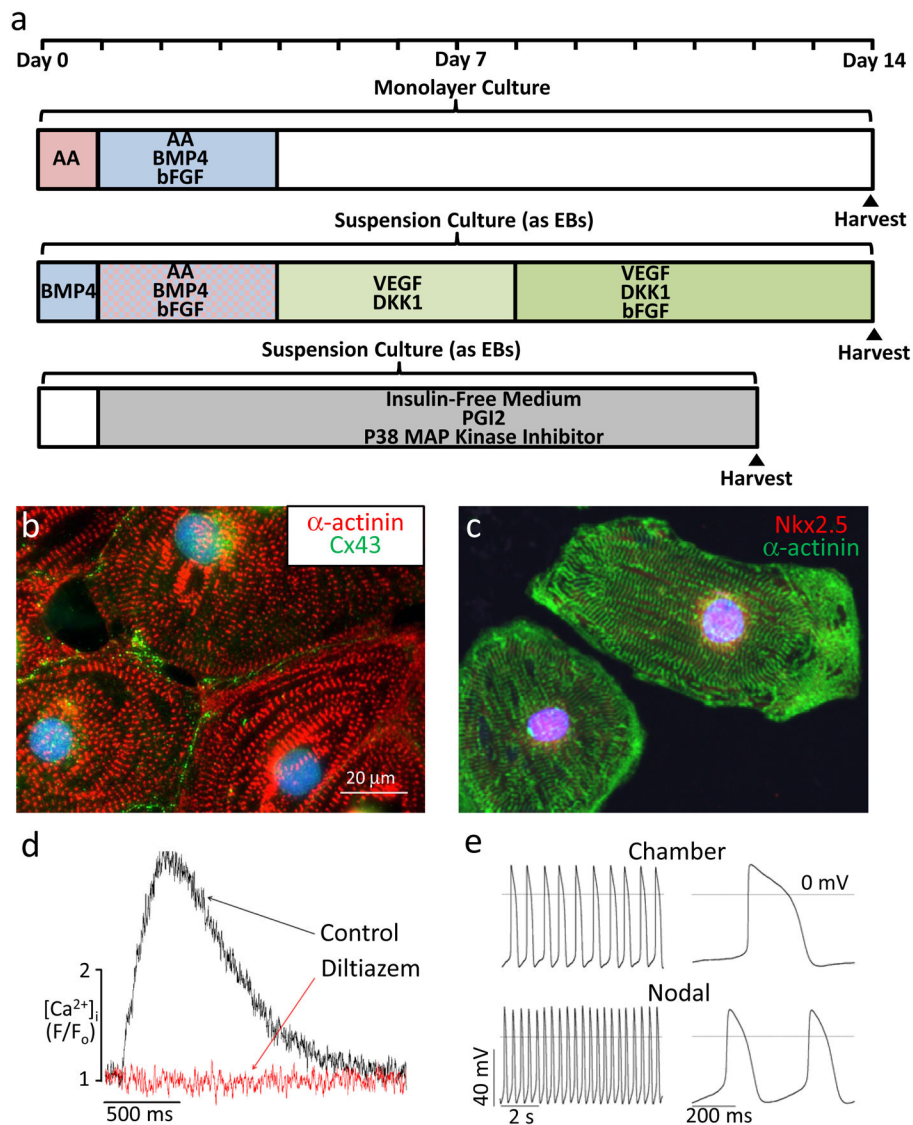


Figure 2. Guided differentiation and phenotype of cardiomyocytes from pluripotent stem cells
a, Selected protocols for the guided differentiation of human ESCs and iPSCs into cardiomyocytes using chemically-defined factors. The upper timeline shows a protocol from our group in which differentiating cells are serially pulsed with activin A (AA) and BMP4 under monolayer culture conditions⁷⁴. The middle timeline shows a protocol from the Keller lab that involves embryoid body (EB) formation in suspension cultures and the application of multiple signaling molecules including activin A, BMP4⁷⁵. The lower timeline shows a protocol from Davidson and colleagues in which EBs are continuously cultured in insulin-free medium supplemented with prostaglandin I₂ and an inhibitor of p38 MAP kinase⁷⁶. **b**, Representative human ESC-derived cardiomyocyte, differentiated using the monolayer protocol, immunostained for α -actinin (red) and connexin43 (Cx43; green). **c**, Representative human iPSC-derived cardiomyocyte, differentiated using the monolayer protocol, immunostained for α -actinin (green) and the transcription factor Nkx2.5 (red). **d**, Intracellular $[Ca^{2+}]_i$ transients in a human ESC-derived cardiomyocyte before (black) and

after (red) application of diltiazem, an L-type Ca^{2+} channel blocker. Absence of $[\text{Ca}^{2+}]$ transients after diltiazem treatment indicates extracellular $[\text{Ca}^{2+}]$ is required to initiate intracellular $[\text{Ca}^{2+}]$ release, just as in adult cardiomyocytes. *e*, Human ESC-derived cardiomyocytes show the characteristic action potential properties of either working chamber (upper) or nodal myocytes (lower), indicating early subtype specification.

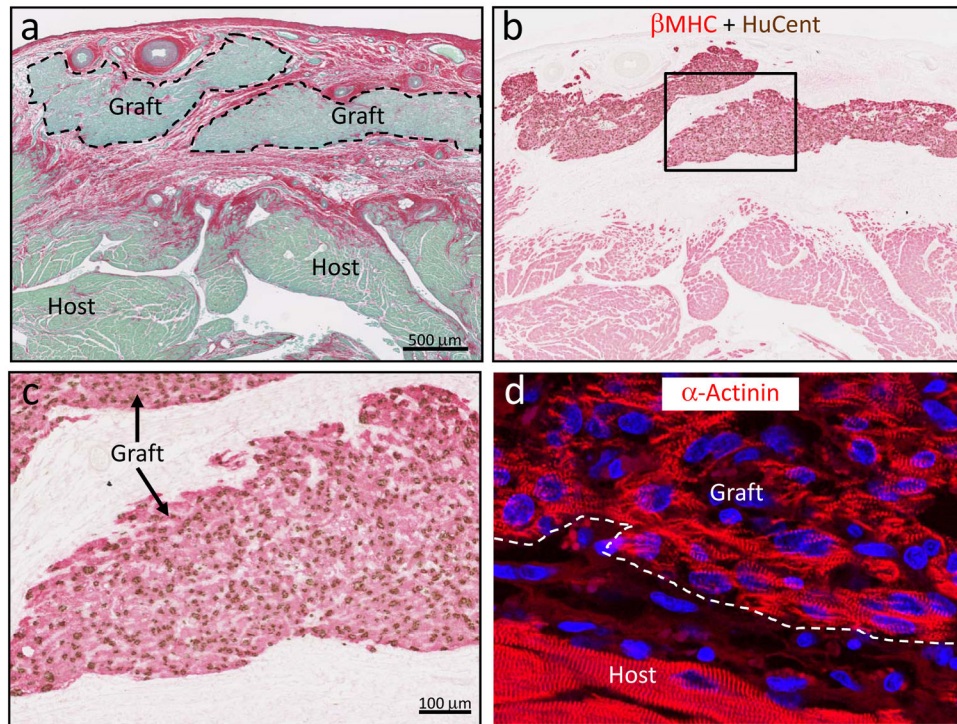


Figure 3. Grafts of human ESC-derived cardiomyocytes in the cryoinjured guinea pig heart
 Representative photomicrographs demonstrating substantial implants of human myocardium within the scar tissue. **a**, By picrosirius red stain, the scar appears red and viable tissue green. Scale bar=500 μm . **b**, The human origin of the graft myocardium was confirmed in an adjacent section by combined in situ hybridization with a human-specific pan-centromeric (HuCent, brown) probe and β -myosin heavy chain (βMHC , red) immunohistochemistry. **c**, Inset from panel b at higher magnification. Note the nuclear localization of the HuCent signal, confirming human origin of these cells. **d**, Immunostaining for α -actinin highlights the sarcomeric organization of the graft myocytes.