The regenerative capacity of zebrafish reverses cardiac failure caused by genetic cardiomyocyte depletion


SUMMARY

Natural models of heart regeneration in lower vertebrates such as zebrafish are based on invasive surgeries causing mechanical injuries that are limited in size. Here, we created a genetic cell ablation model in zebrafish that facilitates inducible destruction of a high percentage of cardiomyocytes. Cell-specific depletion of over 60% of the ventricular myocardium triggered signs of cardiac failure that were not observed after partial ventricular resection, including reduced animal exercise tolerance and sudden death in the setting of stressors. Massive myocardial loss activated robust cellular and molecular responses by endocardial, immune, epicardial and vascular cells. Destroyed cardiomyocytes fully regenerated within several days, restoring cardiac anatomy, physiology and performance. Regenerated muscle originated from spared cardiomyocytes that acquired ultrastructural and electrophysiological characteristics of de-differentiation and underwent vigorous proliferation. Our study indicates that genetic depletion of cardiomyocytes, even at levels so extreme as to elicit signs of cardiac failure, can be reversed by natural regenerative capacity in lower vertebrates such as zebrafish.

KEY WORDS: Cardiomyocyte, Genetic ablation, Heart regeneration, Zebrafish, Epicardium, Endocardium, Heart failure

INTRODUCTION

Heart failure is a disabling, costly human condition of insufficient cardiac output, characterized by exercise intolerance, shortness of breath, edema and increased risk of premature death from arrhythmia or contractile failure. Ischemic, and many non-ischemic, cardiomyopathies commonly involve major cardiac muscle loss and fibrosis, and are primary causes of heart failure (Braunwald and Bonow, 2010; McMurray, 2010). Although the injured adult mammalian heart has a very poor capacity to regenerate cardiac muscle, several directed approaches aim to stimulate this capacity as a means to prevent or treat heart failure. These include identification and manipulation of stem or progenitor cells (Bertini et al., 2003; Lautz et al., 2005; Oh et al., 2003; Winter et al., 2007), reprogramming of non-muscle cells such as fibroblasts into cardiomyocytes (Ieda et al., 2010), or factor delivery to stimulate the limited proliferative ability of cardiomyocytes (Bersell et al., 2009). Although these approaches are promising, they do not yet achieve sufficient cardiac muscle regeneration that would restore function to a failing heart.

Examples of successful heart regeneration exist among vertebrates, and their underlying cellular and molecular mechanisms have the potential to inform future therapies. Cardiac regeneration appears to be an evolutionarily conserved response among adult non-mammalian vertebrate species (Flink, 2002; Kikuchi et al., 2011; Laube et al., 2006; Oberpriller and Oberpriller, 1974; Poss et al., 2002; Rumyantsev, 1973), and regenerative capacity was recently demonstrated in growing hearts of fetal and neonatal mice (Drenckhahn et al., 2008; Porrello et al., 2011). Adult zebrafish are particularly well-studied for their ability to fully regenerate cardiac muscle after surgical resection of ~20% of the ventricle, at or near the maximum amount that can be removed without exsanguination (Poss et al., 2002). Zebrafish hearts also regenerate well after surgery and direct cryoinjury to the ventricle (Chablais et al., 2011; Gonzalez-Rosa et al., 2011; Schnabel et al., 2011). An important mechanistic feature of zebrafish heart regeneration was revealed by recent genetic fate-mapping experiments, which indicated that existing cardiomyocytes are the major source of new muscle after resection of the ventricular apex. Regenerating cardiomyocytes at the injury site display reduced sarcomeric organization, consistent with a mechanism involving some degree of de-differentiation (Jopling et al., 2010; Kikuchi et al., 2010).

Mechanical injury models are not only more tedious and imprecise than desirable, but are limited in their ability to address certain important questions about cardiac regenerative capacity. For example, it is unclear whether larger injuries that might cause signs of acute cardiac failure are subject to similar regenerative capacity and/or mechanisms. Additionally, it is unknown whether internally isolated areas of destroyed myocardium can be effectively replaced with new cardiomyocytes in an adult animal. Finally, whereas mechanical injury removes or destroys many different cardiac cell types and leads to clot formation, cardiac cell type-specific injury models are preferred for certain mechanistic questions; for example, whether cardiomyocyte death on its own, without tissue disruption and clotting, can signal for regeneration. Thus, investigations into new cardiac injury models and potential regenerative responses are crucial for the field.

Here, to address key outstanding issues, we generated a transgenic system to inflict extreme myocardial injury of much greater specificity and severity than has been achieved in previous...
studies. Injuries achieving destruction of >60% of the ventricular myocardium disrupted electric conduction and elicited signs of severe cardiac failure. Yet, an injury response involving all major cardiac cell types was stimulated that rapidly replaced lost myocardium and ameliorated symptoms. This regenerative process featured widespread proliferation by a high percentage of spared cardiomyocytes, cells that displayed multiple indicators of de-differentiation. Our findings demonstrate that adult zebrafish possess a striking capacity for activating regeneration from spared cardiomyocytes after genetically induced muscle loss.

MATERIALS AND METHODS

Zebrafish and heart injuries

Outbred EK or EK/AB mixed background zebrafish (4-10 months of age) were used for ventricular resection surgeries as described previously (Poss et al., 2002), or for cardiomyocyte ablation. Animal density was maintained at approximately four fish per liter in all experiments. To ablate cardiomyocytes, anesthetized Z-CAT fish were injected intraperitoneally with 0.5 mg/ml 4-hydroxytamoxifen (4-HT) in 10% ethanol as described previously (Kikuchi et al., 2010). For bath treatment, animals were treated for 12 hours in 0.02 μM, 0.1 μM or 0.5 μM 4-HT in fish water. Other transgenic strains used were: cmlc2:actinin3-EGFP (Y.-F.L. and D.Y., unpublished), Rl1:EGFP (Lawson and Weinstein, 2002) and bactin2:loxp-DsRed-STOP-loxp-EGFP (Kikuchi et al., 2010). Newly constructed strains are described in the section below. All transgenic strains were analyzed as hemizygotes. All animal procedures were performed in accordance with Duke University guidelines.

Heat stress experiments

Animals with resection injuries, or Z-CAT fish injected with 0.5 mg/ml 4-HT or 10% vehicle, were exposed at 7 days post-injection (dpi) to a transient, automated increase in temperature from 26°C to 38°C, as described (Lee et al., 2005). With this protocol, animals remained at 38°C for one hour before gradual return to 26°C.

Maximum swimming speed and endurance experiments

To evaluate swimming performance, male animals with resection injuries, or male Z-CAT fish that had been treated previously with 0.1 μM 4-HT or vehicle for 12 hours, were placed in a swim tunnel respirometer (Mini Swim-170, Loligo Systems, Denmark) and allowed to acclimatize for 20 minutes at a low flow velocity of 3 cm/s. Swimming speed was then assessed per animal. For resection injuries, we imaged the three ventricular sections with the largest injuries, and quantified Mef2+ and Mef2+PCNA+ cardiomyocytes after genetically induced muscle loss.

Histological methods

In situ hybridization (ISH) on 10 μm cryosections of paraformaldehyde-fixed hearts was performed using digoxigenin-labeled cRNA probes as described (Poss et al., 2002) with the aid of an InSituPro robot (Intavis). Acid Fuchsin-Orange G staining was performed as described (Poss et al., 2002). Primary antibodies used in this study were: anti-Mef2 (rabbit; Santa Cruz Biotechnology), anti-Myosin heavy chain (MHC; F59, mouse; Developmental Studies Hybridoma Bank), anti-GFP (rabbit; Invitrogen), anti-DeRed (rabbit; Clontech), anti-Brdu (rat; AbD) and anti-PCNA (mouse; Sigma). Secondary antibodies (Invitrogen) used in this study were: Alexa Fluor 488 goat anti-rabbit; Alexa Fluor 594 goat anti-rabbit, goat anti-rat and goat anti-mouse; and Alexa 633 goat anti-mouse. Myeloperoxidase staining was performed using a kit from Sigma-Aldrich (390A-1KT) following manufacturer’s instructions. FITC-labeled phallolidin was obtained from Sigma. Immunofluorescence and terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) staining on cryosections were performed as described (Kikuchi et al., 2011).

Myofiber area was measured by imaging three medial, longitudinal F59-stained ventricular sections for each animal, quantifying the fluorescent and total ventricular areas using Openlab and ImageJ software, and calculating an average number of fluorescent pixels per mm² of ventricular tissue. To calculate cardiomyocyte proliferation indices, ventricular (or atrial) sections were stained with antibodies against Mef2 and PCNA. For ablation injuries, three medial, longitudinal sections of each ventricle were selected for analysis, and high-magnification images were taken of different ventricular regions and merged using Photoshop software to represent the entire ventricular section. Mef2⁺ and Mef2⁺PCNA⁺ cells were counted manually with the aid of ImageJ software, with 400-3000 cardiomyocytes assessed per animal. For resection injuries, we imaged the three ventricular sections with the largest injuries, and quantified Mef2⁺ and Mef2⁺PCNA⁺ cardiomyocytes.
cells within a defined region, typically including almost all Me2’PCNA+ cells near the injury (216 pixels in vertical). The percentages of Me2’PCNA+ cells from the three selected sections were averaged to determine a proliferation index for each animal.

For transmission electron microscopy (TEM), injured hearts were fixed in 2% glutaraldehyde/4% formaldehyde. Tissue was embedded using the EMBed 812 resin kit (with DMP-30) (EMS) and sectioned to 70-90 nm thickness using Ultracut or Ultracut E ultra-microtomes (Leica). Microscopy was performed on a FEI Tecnai G2 Twin (200kv) transmission electron microscope and images were obtained using a FEI Eagle camera.

4-HT labeling and quantification of EGFP fluorescence in Z-CAT; bactin2:loxp-DsRed-STOP-loxp-EGFP animals

To quantify Cre-mediated release of EGFP expression in lineage-tracing experiments with Z-CAT; bactin2:loxp-DsRed-STOP-loxp-EGFP animals, three medial, longitudinal sections were selected from each heart and stained with F59. Images of single optical slices of the ventricle were taken using a 20× objective (1024×1024 pixels) by adjusting gain to detect EGFP or MHC signals above background level. To obtain an entire single slice image, we merged images of different ventricular regions using Photoshop software. EGFP+ and MHC+ areas were quantified in pixels by ImageJ software, and the percentage of EGFP/MHC versus MHC area was calculated for each heart.

Optical mapping and data analysis

Detailed methods have been previously described (Kikuchi et al., 2010). Briefly, hearts from anesthetized fish were isolated in Tyrode’s solution and stained with 56 μM di-4-ANEPPS (D-1199, Invitrogen) for 10 minutes. For data acquisition, hearts were transferred to Tyrode’s solution with 30 μM of blebbistatin (EMD Chemicals) and point stimulated with a fine platinum electrode at the base of the ventricle at 80 beats/minute. Optical action potentials were recorded from the epicardial surface of the ventricle. Acquired fluorescence data were analyzed with custom-written software (MatLab, Mathworks) as previously described (Kikuchi et al., 2010; Panakova et al., 2010) Signals were processed with temporal (200 Hz cutoff) and spatial (4-pixel weighted average) filters. Epicardial surface conduction velocities were estimated from regions of interest (ROIs; 156×156 μm) near the apex of each heart. Action potential durations were estimated as the time interval between 20% depolarization and 80% repolarization (APD80). Derived maximum action potential upstroke velocities were calculated as the maximum of the first time-derivative of the action potential upstroke. The data were processed and analyzed using GraphPad Prism5.

Construction of transgenic animals

bactin2:loxp-mCherry-STOP-loxp-DTA

A cassette containing a floxed mCherry followed by five repeats of polyadenylation (pA) sites was subcloned behind the zebrafish β-actin2 promoter (Traver et al., 2003). The pA repeats were derived from sox10-GFP-DTA (provided by W. Richardson, London, UK) and pBigT (Srinivas et al., 2001). The mutated version of DTA (G383A) (provided by M. Capecchi, Salt Lake City, UT, USA) was amplified by PCR using the following primers: 5’-CCGGATCGATACGCGTCCCTAGCCATGGG-3’ and 5’-CCGGCTCGAGTTA-3’. The mutated version of DTA was co-injected with I-Sce1 into one-cell-stage zebrafish embryos. The entire cassette was flanked with 2× core insulator elements and I-Sce1 sites, and co-injected with I-Sce1 into one-cell-stage zebrafish embryos. The full name of this transgenic line is Tg(bactin2:loxp-mCherry-STOP-loxp-DTA176).246.

tcf21:nucEGFP

The translational start codon of tcf21 in the BAC clone DKEYP-79F12 was replaced with the nucEGFP cassette by Red/ET recombineering technology (GeneBridges). The 5’ and 3’ homologous arms for recombination were a 50-base-pair (bp) fragment upstream of the start codon, and a 50-bp fragment downstream, respectively, and were included in PCR primers to flank the nucEGFP cassette. To avoid potential mis-recombination between the nucEGFP cassette and an endogenous loxp site in the BAC vector, we replaced the vector-derived loxp site with an I-Sce1 site using the same technology. The final BAC was purified with Nucleobond BAC 100 kit (Clontech), and co-injected with I-Sce1 into one-cell-stage zebrafish embryos. The full name of this transgenic line is Tg(tcf21:nucEGFP).40.

RESULTS

Genetic ablation of adult zebrafish cardiomyocytes

Modeled on studies using mice (Akazawa et al., 2004; Breitman et al., 1987; Brockschneider et al., 2004; Lee et al., 1998; Saito et al., 2001; Stanger et al., 2007), we generated a double transgenic system to facilitate cell type-specific ablation in zebrafish. The first transgenic line has a 4-hydroxytamoxifen (4-HT)-inducible Cre recombinase (CreER) restricted to cardiomyocytes by the cmle2 (myl7 – Zebrafish Information Network) promoter, as described recently (cmle2:CreER) (Kikuchi et al., 2010). We created a second line (bactin2:loxp-mCherry-STOP-loxp-DTA) that targets cytotoxic DTA (diptheria toxin A chain) expression to CreER-expressing cells upon 4-HT injection (Fig. 1A). The β-actin2 promoter fragment drives expression strongly in myocytes, with little or no expression in epicardium and endocardium (Kikuchi et al., 2010). We intentionally selected a line that displayed moderate mCherry expression in what appeared to be most, if not all, cardiomyocytes (see Fig. S1A in supplementary material). To test the efficiency of myocyte ablation, we intercrossed these lines and injected 6-month-old cmle2:CreER; bactin2:loxp-mCherry-STOP-loxp-DTA animals intraperitoneally with 0.5 mg/ml 4-HT. This treatment was...
The text describes experimental procedures and results regarding cardiomyocyte ablation and regeneration. It mentions the use of multiple assays to confirm and quantify cardiomyocyte ablation, with a focus on histology and digital quantification. The text also discusses the role of apoptosis in myocardial death and the use of TUNEL staining to assess cell death. The rapid regeneration of ventricular cardiomyocytes after ablation is highlighted, with a focus on the time course of cardiomyocyte proliferation assessed by Mef2 and PCNA staining. The text includes statistical analysis and figures to support the findings.
exercise tolerance in the same animals that had underperformed at 7 dpi, we found substantial improvements to the levels of control animals (Fig. 2C). Swimming endurance, the average time during which animals could maintain orientation in a high velocity current, was not different in vehicle- and 4-HT-injected animals at 45 dpi (data not shown). Thus, massive ablation of ventricular muscle dramatically reduced the performance of adult zebrafish during stress and exercise, but these abilities recovered within several days.

We next assessed histological evidence that might explain this recovery. Quantifications of myofiber area and Mef2c+ cardiomyocyte nuclei at 14 dpi indicated that animals restored 39% of lost myofiber area (to 63% of that of vehicle-treated animals) and 44% of lost myocyte nuclei (to 60%) within one week. These measurements suggested that zebrafish require between ~40 and 60% of their cardiomyocytes for normal performance in the exercise assay that we used. By 30 dpi, ventricles were fully muscularized (Fig. 3A-C). Measurements of ventricular section surface area at 30 dpi indicated no increases in cardiac size with this nuclear recovery (see Fig. S1E in the supplementary material), suggesting that cellular hypertrophy is not a major component of muscle restoration. To determine whether muscle restoration occurred as a result of cell proliferation, we assessed the percentage of myocytes expressing the marker proliferating cell nuclear antigen (PCNA), which compares well with BrdU incorporation assays (see Fig. S3 in the supplementary material) (Wills et al., 2008). Whereas 3% of myocytes were PCNA+ in vehicle-treated Z-CAT animals, 42% of myocytes displayed nuclear PCNA at 7 days after 4-HT injection, and 34% at 14 dpi. These proliferating cardiomyocytes were distributed throughout the ventricle, suggesting that most cardiomyocytes can participate in the regenerative response. Cardiomyocyte proliferation returned to near baseline levels by 30 dpi, concomitant with muscularization (Fig. 3D,E). We also confirmed that scarring, the predominant response of mammals to cardiac injury, was not a significant response to myocardial ablation. Slight increases in cardiac collagen deposition were occasionally observed at 14 dpi and 30 dpi compared with control animals. However, large deposits were generally not detected, even at 180 dpi (see Fig. S4 in the supplementary material; data not shown). Our data indicate rapid and complete regeneration after ablation-induced muscle damage that was severe enough to elicit signs of cardiac failure.

Robust injury responses by ventricular non-muscle cells and atrial myocytes after genetic cardiomyocyte ablation

Prior to this study, it had not been known whether diffuse, extensive adult cardiomyocyte loss could be replaced through regeneration. In addressing this, our results indicated considerably more efficient regeneration in this context than from the injury zone cardiomyocytes that are spared by resection of the zebrafish ventricle. In that case, regeneration is typically completed by 1-2

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**Fig. 4. Effects of cardiomyocyte ablation on non-muscle cells.** (A) Visualization of endocardial cells (green) and muscle (red) in sections from vehicle-or 4-HT-injected Z-CAT fish. The endocardial layer appears intact one week after massive cardiomyocyte loss. Single confocal slices are shown. (B) In situ hybridization for pu.1-expressing cells, which are prominent by 3-5 dpi. (C) Whole-mount visualization of epicardial nuclei in vehicle- or 4-HT-injected Z-CAT fish at 7 dpi, indicating higher epicardial cell density after myocardial injury. Inset shows enlarged view of boxed area. (D) Histological visualization of epicardial cells after myocardial ablation. Epicardial cells proliferate by 7 dpi and are incorporated into the regenerating myocardial wall by 14 dpi. Brackets indicate area containing tcf21-expressing epicardial cells. (E) Visualization of coronary vascular endothelial cells (green) in sections of the ventricular wall (outlined by dotted lines). Coronary vasculature is reduced in 4-HT-injected Z-CAT animals by 7 dpi, but restored by 14 dpi. (F) Sections from Z-CAT ventricles stained for the retinoic acid (RA)-synthesizing enzyme Raldh2. RA synthesis appears low in vehicle-injected tissue, but is strongly induced in epicardial (arrowheads) and endocardial (arrows) structures after cardiomyocyte ablation. Single confocal slices are shown. Scale bars: 50 μm.
months post-injury, with only 10-15% of cardiomyocytes at the resection plane proliferating (PCNA+) at 7 dpa (Kikuchi et al., 2011; Poss et al., 2002). Interestingly, in mammalian skeletal muscle, regeneration is poor when large areas of muscle and connective tissue are removed, e.g. by amputation; yet, injuries that preserve the connective tissue scaffolding permit vigorous regeneration (Tajbakhsh, 2009). Thus, we surmised that the presence and activity of analogous, non-muscle components like endocardial, epicardial and vascular cells contribute to regenerative efficacy after cardiomyocyte ablation.

To visualize endocardial cells that line the inner, trabecular myocardium, we crossed the fli1:EGFP transgene into the Z-CAT background. We found that endocardial architecture was grossly intact at 7 dpi despite cardiomyocyte depletion (Fig. 4A). We examined injured ventricles for macrophage presence within and around these endocardial structures. In situ hybridization with a pu.1 (spi1 – Zebrafish Information Network) probe showed little staining in vehicle-treated Z-CAT vehicles. By contrast, upon cardiomyocyte ablation there was a massive enrichment of injured trabecular myofibers as well as the compact myocardial layer with pu.1 cells, most prominently at 3 and 5 dpi, periods of peak cell death (Fig. 4B). This inflammatory response to cardiomyocyte ablation also included neutrophil recruitment, as assessed by myeloperoxidase staining (see Fig. S5 in the supplementary material). Thus, the endocardium is spared after severe myocyte ablation, and endocardial-coated pockets of ablated myocardium are targets of macrophage and neutrophil infiltration.

The peripheral compact muscle is surrounded and penetrated by epicardial cells rather than endocardium. To visualize epicardium, we generated a new transgenic strain reporting nuclear-localized EGFP in virtually all epicardial cells (tcf21:nucEGFP). Epicardial cells surrounding the ventricle were stimulated by ablation injury to proliferate and form a multilayered structure by 7 dpi (Fig. 4C,D). By 14 dpi, these augmented epicardial cells became incorporated into the underlying myocardial wall, events suggestive of epithelial-mesenchymal transition (Fig. 4D). The epicardium has been proposed to support neovascularization during regeneration, analogous to its known embryonic roles (Kim et al., 2010; Lepilina et al., 2006; Kikuchi et al., 2011). We observed an initial reduction in the density of vessels within the ventricular wall at 7 dpi. However, one week later and concomitant with muscle regeneration, the ventricular wall was well-vascularized, indicative of a neovascularization process (Fig. 4E).

Endocardial and epicardial cells are likely to impact cardiac muscle regeneration in several ways. Soon after resection injury, retinoic acid (RA) synthesis, as visualized by expression of the enzyme Raldh2 (Aldh1a2 – Zebrafish Information Network), is increased in endocardial and epicardial cells throughout the injured ventricle and uninjured atrium (Kikuchi et al., 2011; Lepilina et al., 2006; Wills et al., 2008). During regeneration, RA production localizes to injuries, where RA signaling is essential for cardiomyocyte proliferation (Kikuchi et al., 2011). Raldh2 was induced and maintained strongly and throughout ventricular endocardial and epicardial tissue at 7 and 14 days, adjacent to regions of ablated myocardium (Fig. 4F). We expect that this response to ablation injury is likely to have the same function as after resection injury. In total, these analyses indicate that widespread cardiomyocyte-specific cell ablation elicits strong injury responses from non-myocardial cells, including endocardial and epicardial RA production, epicardial proliferation and neovascularization, all of which are likely to impact rapid regeneration from extreme cardiac damage.

Finally, we also took advantage of this ablation system to assess atrial myocyte regeneration. The zebrafish atrium is difficult to access surgically, and its regenerative capacity has not previously been assessed. Because atrial muscle is also subject to genetic
ablation via the cmlc2 promoter, we examined apoptosis and proliferation in this chamber after an injection of 0.5 mg/ml 4-HT. Atrial myocyte apoptosis showed similar dynamics as ventricular myocytes, with evenly distributed TUNEL-positive myocytes by 5 dpi (Fig. 5A). Atrial myocyte proliferation indices were 43% at 7 days and 23% at 14 dpi, versus 4% in vehicle-injected Z-CAT fish, indicative of a strong regenerative response (Fig. 5B,C). Thus, the zebrafish atrium, like the ventricle, is highly regenerative after major ablation injury.

Ultrastructural and physiological de-differentiation in source myocytes during regeneration
To determine the source of new cardiomyocytes regenerated after ablation injury, we crossed a lineage-tracing reporter transgene (bactin2:loxP-DsRed-STOP-loxP-EGFP) into Z-CAT. Recombination between loxP sites releasing EGFP expression is more efficient than events releasing DTA expression, and resulted in 96% of spared ventricular muscle being labeled with EGFP at 7 dpi after myocardial ablation (Fig. 6A-C). Assessment of ventricles at 30 dpi indicated that, similarly, 95% of regenerated muscle retained the EGFP label (Fig. 6B,C). To determine whether cardiomyocytes dissembled sarcomeres during ablation-induced regeneration, we crossed a Z-line-labeling cmlc2:actinin3-EGFP transgene into Z-CAT. At 7 dpi, and more so at 14 dpi, there were many cardiomyocytes with visibly disordered sarcomere structure, a phenotype indicative of a more primitive differentiation state and reduced contractile properties (Fig. 6D). These observations were corroborated by transmission electron microscopy, which also revealed disorganized myocyte sarcomere structure at 7 and 14 dpi (Fig. 6E). Sarcomere disorganization was coincident with cardiomyocyte proliferation and subsequent to cardiomyocyte apoptosis, making it unlikely that they reflect dying DTA-expressing cells. Also, the sudden appearance of disorganized sarcomeres in a high percentage of myocardium throughout the ventricle (~23 times the frequency of that observed in uninjured ventricles; data not shown) makes it unlikely that they arose from rapid expansion of an existing population of poorly differentiated cardiomyocytes.

**Fig. 6. Lineage-tracing and ultrastructural analysis of regenerating muscle.** (A) Schematic representation of transgenes used for myocyte ablation and lineage tracing. (B) Triple transgenic (Z-CAT; bactin2:loxP-DsRed-STOP-loxP-EGFP) zebrafish were injected once with vehicle (left) or 4-HT. In 4-HT-injected animals, EGFP labeled the majority of spared cardiomyocytes by 7 dpi (middle) and the majority of regenerated cardiomyocytes by 30 days post-injection (dpi; right). Arrowheads indicate examples of MHC* tissue expressing EGFP. (C) Quantification of EGFP* myocardium as a percentage of MHC* ventricular tissue. At 7 dpi, 96% of myocardium was EGFP-labeled, and 95% at 30 dpi. For each group, 4-10 animals were assessed. Insets show enlarged views of the boxed area. (D) Assessment of sarcomere structure in Z-CAT fish crossed to a transgenic strain that labels Z-lines (green). Cardiomyocytes of control animals show well-organized sarcomeres with clear Z-lines. By contrast, many myocytes of 4-HT-injected animals display disorganized sarcomeres and loss of Z-lines (arrowheads) by 7 dpi and 14 dpi. By 30 dpi, sarcomeric structure and Z-lines are typically restored. Scale bars: 50 μm. (E) Transmission electron micrographs of ventricular myocytes from vehicle- or 4-HT-injected Z-CAT animals. Control myocytes have prominent sarcomeric structure and normal mitochondria (m), whereas 7 and 14 dpi myocytes appear less organized with fewer Z-lines (arrows) and swollen mitochondria (arrowheads). Scale bar: 0.5 μm.
myocytes. By 30 dpi, sarcomere structure was similar to uninjured ventricles (Fig. 6D). Thus, our results indicate that regenerating cardiomyocytes are primarily derived from spared, mature cardiomyocytes that transiently de-differentiate throughout the ventricle after ablation injury.

Finally, we examined whether cardiomyocytes show electrophysiological evidence of reduced differentiation during regeneration. Optical mapping of action potentials in ventricles from 4-HT-injected Z-CAT fish at 7 dpi revealed a sharp increase in the density of isochrones across the entire myocardium, indicating a slowing of conduction (Fig. 7A,D). This slowing was accompanied by fractionation of the depolarization wavefront, with areas of patchy surface breakthrough that were rarely observed in control hearts. In addition, we observed significant prolongation of action potential duration in the hearts of 4-HT-injected animals at 7 dpi, as well as substantial slowing of action potential upstroke velocities (Fig. 7B,C,E). The conduction slowing and wavefront fractionation observed at 7 dpi suggests markedly reduced coupling between myocytes throughout the regenerating myocardium. The wavefront fractionation might also represent intercalation of non-coupled epicardial cells into the myocardium with resultant conduction block and delayed subepicardial impulse propagation. In addition to the conduction findings, the prolongation of action potential duration and diminution of peak action potential upstroke velocities recapitulate the electrical phenotypes of less differentiated cardiomyocytes (Milan et al., 2009; Panakova et al., 2010). Such changes in action potential upstroke velocity probably represent an increased dependence of depolarization on calcium channel conductance and a reduced role for sodium channel conductance (Chopra et al., 2010; Milan et al., 2009; Panakova et al., 2010). Global myocardial surface conduction progressed towards normal velocities at 14 dpi and 30 dpi, with recovery observed at 45 dpi. Similarly, wavefront fractionation was considerably reduced at 14 dpi and 30 dpi, and had nearly resolved by 45 dpi (Fig. 7A,D). Taken together, these observations indicate that functional de-differentiation and partial uncoupling occur in myocytes regenerating after ablation-induced injury, followed by ultrastructural and electrical recovery.

**DISCUSSION**

Here, we introduced a model of cardiac injury in zebrafish that facilitates ablation of a high proportion of adult cardiac myocytes. We found that zebrafish survive after depletion of >60% of their ventricular myocardium, but exhibit signs that they are experiencing cardiac failure. In this condition, the cardiac chambers presented as dilated, engorged sacs, that were sparsely muscularized and structurally buoyed by epicardium and endocardium. Despite the extreme effects of these injuries, cardiomyocyte proliferation was robustly initiated and muscle regeneration was rapidly completed. Because this injury model stimulated regenerative events throughout the heart, transient changes in cardiomyocyte sarcomere organization and electrophysiology were clear and widespread, each observation being indicative of de-differentiation. Most impressively, all signs of cardiac failure were reversed concomitantly with these regenerative events. Although observed in a non-mammalian model system, we suggest that these findings provide strong support to the concept of regenerative therapy for certain forms of human heart failure.

Our study reveals other important features of cardiac regenerative capacity. First, it is clear that large areas of internal, destroyed myocardium can be effectively replaced with new cardiomyocytes. Thus, regeneration is not purely a characteristic of cardiomyocytes with greater access to components of a blood clot or that are not encased within spared muscle. As injury is widespread in the cardiac chambers of Z-CAT animals, there is greater participation in regeneration by the inner trabecular form of cardiac muscle than after a ventricular resection injury, illuminating...
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Supplementary material
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Akazawa, H., Komazaki, S., Shimomura, H., Terasaki, F., Zou, Y., Takano, H., Nagai, T. and Komuro, I. (2004). Diphtheria toxin-induced autophagic responses, such as cardiomyocyte death on its own, as opposed to signals emanating from sheared tissue or a blood clot, is a potent signal for regeneration. How myocyte death instructs regeneration will be an important topic for further investigations. Finally, our data indicate that regeneration occurs with two- to fourfold higher cardiomyocyte proliferation indices after ablation than after resection, an unexpected boost in regenerative efficiency that we suspect reflects structural, vascularizing and/or paracrine influences of non–muscle cells. Whereas cardiomyocyte ablation injury spares and enhances endocardial and epicardial tissues, respectively, a ventricular apex injured by resection, with all major cell types lost in the resected portion, requires repair of the endocardial scaffold and recruitment of a new epicardial covering before these tissues can optimally support muscle regeneration. We also find here that cardiomyocyte death is an efficient attractant for macrophages to injured myofibers in the absence of a large clot, recruitment that might be key to regeneration through tissue debris clearance and/or intercellular signaling.

Regeneration in non-mammalian vertebrate model systems has historically been limited by discovery tools. We suggest that Z-CAT has several advantages as a cardiac injury and regeneration system. Ablation injuries are easy to perform and noninvasive, and extensive injuries can be inflicted without affecting animal viability. Certain questions can also be addressed more sensitively; for example, how cardiomyocyte death triggers developmental responses, such as raldh2 induction, in epicardial and endocardial tissue. Finally, this system, like nitroreductase ablation systems previously described for larval zebrafish regeneration studies (Curado et al., 2007; Pisharath et al., 2007), has high potential for use in screens to identify new regulators of regeneration. Many adult animals with major cardiac injuries can be generated daily by 4-HT injection or incubation, and the impact of small molecule ablation/regeneration system described here will enhance efforts to understand how and why heart regeneration occurs in zebrafish. This information, in turn, can inform approaches to musclerizing bioengineered scaffolds, or to reversing ischemic or non-ischemic muscle loss in mammals through regeneration.


