$p38\alpha$ MAPK regulates myocardial regeneration in zebrafish

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Although adult mammals are unable to significantly regenerate their heart, this is not the case for a number of other vertebrate species. In particular, zebrafish are able to fully regenerate their heart following amputation of up to 20% of the ventricle. Soon after amputation, cardiomyocytes dedifferentiate and proliferate to regenerate the missing tissue. More recently, identical results have also been obtained in neonatal mice. Ventricular amputation of neonates leads to a robust regenerative response driven by the proliferation of existing cardiomyocytes in a similar manner to zebrafish. However, this ability is progressively lost during the first week of birth. The fact that adult zebrafish retain the capacity to regenerate their heart suggests that they either possess a unique regenerative mechanism, or that adult mammals lose/ inhibit this process. p38a MAPK has previously been shown to negatively regulate the proliferation of adult mammalian cardiomyocytes. We sought to determine whether a similar mechanism exists in adult zebrafish, and whether this needs to be overcome to allow regeneration to proceed. To determine whether p38 α MAPK also regulates zebrafish cardiomyocytes in a similar manner, we generated conditional transgenic zebrafish in which either dominant-negative or active p38 α MAPK are specifically expressed in cardiomyocytes. We found that active p38 α MAPK but not dominantnegative p38a MAPK blocks proliferation of adult zebrafish cardiomyocytes and, consequently, heart regeneration as well. It appears that adult zebrafish cardiomyocytes share many characteristics with adult mammalian cardiomyocytes, including p38a MAPK-mediated cell cycle inhibition. These findings raise the possibility that zebrafish-like heart regeneration could be achieved in adult mammals.

Introduction

Adult zebrafish cardiomyocytes possess the ability to proliferate in vivo, which permits them to regenerate the heart following substantial amounts of damage.¹⁻³ However, the same is not true for adult mammalian cardiomyocytes, hence the inability of adult mammals to mount any significant regenerative response following cardiac injury. Nevertheless, adult mammalian cardiomyocytes can be induced to proliferate in vitro, which indicates that the necessary cellular machinery is present.⁴ This could mean that adult mammalian cardiomyocytes are not receiving the necessary proliferative signals, or that these are present but proliferation is being inhibited. Indeed, it is possible to induce adult mammalian cardiomyocytes to proliferate by either stimulating the cell cycle^{5,6} or by overriding inhibitory signals.⁷⁻⁹ Because adult zebrafish obviously possess the necessary proliferative signals, we sought to determine whether they are also regulated by known inhibitory mechanisms identified in mammals and hence begin to elucidate the difference in proliferative potential between adult mammalian and adult zebrafish cardiomyocytes. In mammals,

p38\alpha MAPK activity is inversely correlated with heart growth. During fetal development, when cardiomyocytes proliferate readily, p38\alpha MAPK activity is low; however, shortly after birth, p38a MAPK activity increases and is maintained throughout adult life.7 Inducing p38\alpha MAPK activity in fetal cardiomyocytes either in vitro or in vivo results in a substantial reduction of proliferation and, consequently, cardiogenesis is perturbed.^{7,10} To determine whether p38\alpha MAPK also regulates zebrafish cardiomyocytes in a similar manner, we generated conditional transgenic zebrafish in which either dominant-negative(dn) p38α MAPK (tg:cmlc2:LrL:dn p38α MAPK) or a constitutively active (ca)MKK6 (the upstream activator of p38a MAPK and a well-established technique for inducing and maintaining high levels of active p38α MAPK¹¹⁻¹³) (tg:cmlc2:LrL:caMKK6) are specifically expressed in cardiomyocytes following tamoxifen (4-HT)/Cre mediated recombination; both lines were created on our previously published tg-cmlc2a-Ert2-Cre-Ert2/tg-cmlc2a-LnL-GFP background,¹⁴ allowing us to monitor Cre activity via GFP expression and to concomitantly lineage trace recombined cardiomyocytes.

Cell Cycle

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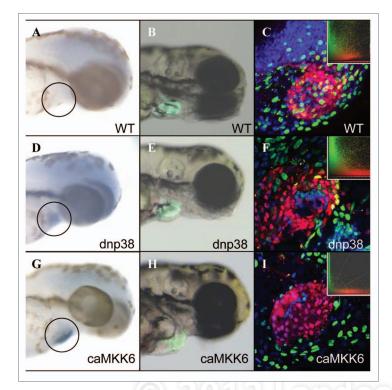


Figure 1. Cardiomyocyte specific expression of caMKK6 disrupts zebrafish cardiogenesis. In situ hybridization for either caMKK6 or dnp38 α MAPK in wild-type 3dpf embryos (A), dnp38 α MAPK-expressing embryos (D) or caMKK6-expressing embryos (G); the black circle indicates approximately where the heart is. Morphological analysis of cardiogenesis, (cardiomyocytes are expressing GFP under the control of the *cmlc2a* promoter) in wild-type 3dpf embryos (B), dnp38 α MAPK-expressing embryos (E) or caMKK6-expressing embryos (H). BrdU colocalization with cardiomyocytes (BrdU, green; cardiomyocytes, red) in wild-type 3dpf embryos (C), dnp38 α MAPK-expressing embryos (F) or caMKK6-expressing embryos (I); insets show a graphical representation of BrdU/cardiomyocyte colocalization.

Results

Cardiomyocyte-specific expression of caMKK6 disrupts zebrafish cardiogenesis. Previous results in mammals indicate that active p38\alpha MAPK inhibits the proliferation of fetal cardiomyocytes.7 To determine if this is also the case in zebrafish, we induced expression of either dnp38a MAPK or caMKK6 specifically in cardiomyocytes during embryogenesis. Transgenic (tg:cmlc2:LrL:dn p38α MAPK) or (tg:cmlc2:LrL:caMKK6) zebrafish embryos were treated at 24 hpf with 4-HT then allowed to develop for a further 48 hrs. In situ hybridization using antisense dnp38a MAPK or caMKK6 RNA probes revealed robust expression of both transgenes, specifically in the heart of developing zebrafish embryos obtained from their respective transgenic lines; no expression was observed in wild-type embryos (Fig. 1A, D and G). During zebrafish embryogenesis, the heart undergoes a series of distinct morphological changes. By 24 hpf, the heart tube has been established, then during the next 24 hrs, it performs an S-shaped loop, bringing the prospective atria and ventricle adjacent to one another.¹⁵ We could not detect any noticeable morphological defects in embryos expressing dnp38α MAPK compared with wild-type embryos (Fig. 1B and E), similar to results obtained from conditional, cardiomyocyte-specific p38a MAPK-knockout mice.¹⁶ However, embryos expressing caMKK6 exhibited severely disrupted cardiogenesis characterized by a failure of the heart tube to loop by 3dpf (Fig. 1H). Despite the severity of this phenotype, we could not detect any obvious cytotoxic effects, such as heart edema produced by expressing caMKK6. To determine whether cardiomyocyte proliferation had been affected by expression of caMKK6, we BrdU labeled developing embryos between 24 hpf and 48 hpf. Previous research has established that cardiomyocyte proliferation does occur during this developmental time window, albeit at a relatively low rate.¹⁷ We were able to detect BrdU labeling co-localized with RFP-positive cardiomyocytes in both wild-type and dnp38a MAPK embryos; however, this was not the case in caMKK6 embryos, which show virtually no co-localization of BrdU with cardiomyocytes (Fig. 1C, F and I, insets). These results indicate that caMKK6 inhibits cardiomyocyte proliferation during zebrafish cardiogenesis, and also that the small amount of proliferation that normally occurs in wild-type embryos is required for cardiac looping.

Mosaic caMMK6 expressing cardiomyocytes do not contribute to the regenerative process. Although adult mammalian cardiomyocytes are unable to proliferate, a concomitant inhibition of p38 α MAPK in conjunction with FGF stimulation is sufficient to induce cardiomyocyte proliferation both in vitro and in vivo.⁷ We next sought to determine whether expression of caMKK6 would inhibit the extensive proliferation of cardiomyocytes that occurs during zebrafish heart regeneration.^{1,14} To achieve this, we utilized a transgenic line in which the Cre transgene has been partially silenced, resulting in random mosaic expression of Cre throughout the heart. Consequently, treatment

with 4-HT induces Cre-mediated recombination only in a subset of cardiomyocytes, which begin to express GFP (Fig. 2A-D). This allowed us to express caMKK6 in a random subpopulation of cardiomyocytes and then determine whether these cells proliferate and contribute to the regenerative process (Fig. 2D). At 14 d post-amputation (dpa), we found substantial regeneration had occurred with new myocardial tissue clearly labeled for α -sarcomeric actin (Fig. 2E). Within the same heart, only a random subpopulation of cardiomyocytes express caMKK6 (Fig. 2F); however, no caMKK6 expressing cardiomyocytes are present in the newly regenerated tissue (Fig. 2E-H). BrdU labeling also clearly delineates the regenerated tissue with the most abundant BrdU labeling clearly overlapping the newly formed myocardium (Fig. 2E and G). At 30dpa, heart regeneration is almost complete, and again we found that the newly regenerated tissue was completely lacking any cardiomyocytes expressing caMKK6 (Fig. 2I-L). To determine whether any of the caMKK6 cardiomyocytes had proliferated, we first counted the number of BrdU-positive nuclei per section then subsequently determined what proportion of these were cardiomyocytes. In control 14dpa hearts, we found that 52.1% of the total amount of counted

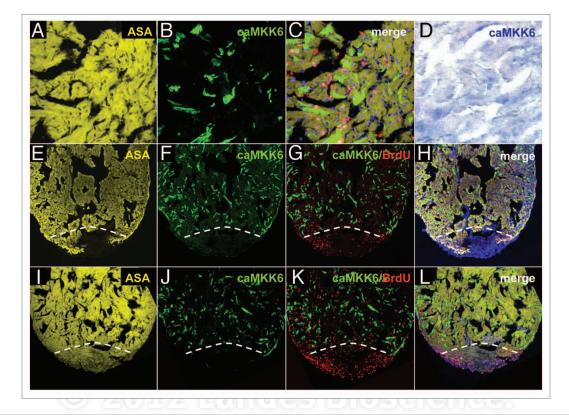


Figure 2. Mosaic caMMK6 expressing cardiomyocytes do not contribute to the regenerative process. An adult zebrafish heart immunolabelled for α *sarcomeric actin* (yellow) (A), GFP(caMKK6) (green) (B), merged image showing α *sarcomeric actin* (yellow), GFP(caMKK6) (green) and BrdU (red) (C). In situ hybridization for caMKK6 (blue) in a similar mosaic adult zebrafish heart (D). A 14dpa regenerating heart (white dashed line indicates the plane of amputation) immunolabelled for α *sarcomeric actin* (yellow) (E), GFP(caMKK6) (green) (F), GFP(caMKK6) (green) and BrdU (red) (G) and a merged image showing α *sarcomeric actin* (yellow), GFP(caMKK6) (green), BrdU (red) and DAPI (blue) (H). A 30dpa regenerating heart (white dashed line indicates the plane of amputation) immunolabelled for α *sarcomeric actin* (yellow) (I), GFP(caMKK6) (green) (J), GFP(caMKK6) (green) and BrdU (red) (K) and a merged image showing α *sarcomeric actin* (yellow), GFP(caMKK6) (green), BrdU (red) and DAPI (blue) (L).

cells labeled with BrdU (6,887) were cardiomyocytes (3,591). However, although considerable proliferation occurs in regenerating hearts mosaically expressing caMKK6, we could not detect any BrdU labeling in cardiomyocytes that expressed caMKK6 (BrdU-positive nuclei counted = 4,657, BrdU positive/caMKK6positive nuclei = 0). These results indicate that cardiomyocytes expressing caMKK6 are unable to proliferate and do not contribute to the regenerative process.

Expression of caMKK6 in all cardiomyocytes effectively blocks zebrafish heart regeneration. Although we could not observe any proliferation of caMKK6-expressing cardiomyocytes, this does not necessarily mean that caMKK6 is completely blocking this process. Another possible explanation is that cardiomyocytes not expressing caMKK6 preferentially proliferate rather than the caMKK6 expressing cells. To determine to what extent expression of caMKK6 blocks cardiomyocyte proliferation, we performed identical amputation experiments on transgenic zebrafish that express caMKK6 in all cardiomyocytes using a Cre transgenic line which is not silenced. As we have previously shown, at 14dpa, a substantial amount of myocardial regeneration normally occurs (Fig. 3A and B). We could not detect any observable differences in regenerative capacity between controls hearts and those expressing dnp38α MAPK (Fig. 3A-H). However, in hearts expressing caMKK6 in all cardiomyocytes, regeneration, if not completely blocked, was severely inhibited (n = 8). Few if any cardiomyocytes had been produced below the plane of amputation, and in a significant number of hearts (n = 3/8), a large clot remained next to the plane of amputation, something we have never observed in wild-type regenerating hearts at this stage (Fig. 3I and J). Similarly, at 30dpa, when regeneration is virtually complete in wild-type hearts (Fig. 3C and D), this process has been severely inhibited in zebrafish expressing caMKK6 (n = 10) (Fig. 3K and L). These results indicate that the expression of caMKK6 effectively blocks myocardial regeneration in zebrafish. Furthermore, we could not detect any obvious GFP-negative myocardium (Fig. 3I–L), indicating that if stem/progenitor cells reside in the zebrafish heart, then they are unable to significantly regenerate the myocardium when cardiomyocyte proliferation is blocked.

Adult zebrafish cardiomyocytes express active $p38\alpha$ MAPK which is switched off upon entry into mitosis. Although active $p38\alpha$ MAPK naturally inhibits the proliferation of adult mammalian cardiomyocytes, one possible explanation for our observed results is that this mechanism does not exist in wildtype zebrafish, which would go some way to explaining why zebrafish are capable of extensive heart regeneration and mammals are not. To determine whether or not this is the case, we first examined if active $p38\alpha$ MAPK is present in cultured adult

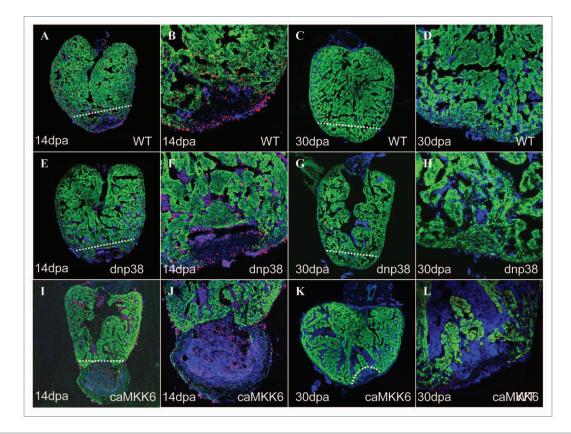


Figure 3. Expression of caMKK6 in all cardiomyocytes effectively blocks zebrafish heart regeneration. A 14dpa wild-type regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (A). The same heart at higher magnification (B). A 30 dpa wild-type regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (C). The same heart at higher magnification (D). A 14dpa dnp38α MAPK regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (E). The same heart at higher magnification (F). A 30dpa dnp38α MAPK regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (H). A 14dpa caMKK6 regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (I). The same heart at higher magnification (J). A 30dpa caMKK6 regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (J). A 30dpa caMKK6 regenerating heart immunolabeled for GFP (green), BrdU (red) and DAPI (blue); the white dashed line indicates the plane of amputation (L).

zebrafish cardiomyocytes. Using a specific active p38a MAPK antibody that had been verified via western blot, we found that all adult zebrafish cardiomyocytes contain active p38a MAPK in their nucleus (Fig. 4A and B). To determine whether active $p38\alpha$ MAPK needs to be switched off during cardiomyocyte proliferation, we first analyzed BrdU-labeled cardiomyocytes. Although the majority of BrdU labeled cardiomyocytes retain a clear active p38a MAPK signal, we were able to find individuals in which the signal was absent, indicating that active p38α MAPK may need to be switched off at a specific point during the cell cycle (Fig. 4A-D). p38a MAPK is known to induce arrest at various points in the cell cycle, including the G₂/M phase.¹⁸ To establish if p38α MAPK activity must be switched off in order for cardiomyocytes to enter mitosis, we immunolabelled them with a phospho histone H3 (PHH3) antibody that is only present during the M phase of the cell cycle.¹⁹ We found numerous cardiomyocytes labeled with PHH3; however, the active p38a MAPK signal was completely absent from these cells (51 cells PHH3 positive/active p38a MAPK negative) (Fig. 4E-H). We next sought to determine if this was also the case in vivo during heart regeneration. Adult zebrafish hearts were amputated and allowed to regenerate for 10 d; these were subsequently analyzed via immunohistochemistry using PHH3 and active p38 α MAPK antibodies. We were unable to detect any PHH3 signal co-localized with active p38 α MAPK (n = 10 PHH-positive cardiomyocytes analyzed from two different hearts) (Fig. 4I–L). This shows that in order for adult zebrafish cardiomyocytes to enter into mitosis, they must first switch off p38 α MAPK activity.

Discussion

Here we have shown that adult zebrafish and mammalian cardiomyocytes share a common molecular mechanism that regulates their proliferation. Our findings indicate that active p38 α MAPK negatively regulates the proliferation of adult zebrafish cardiomycoytes in a similar manner to previous reports in mammals.⁷ p38 α MAPK activity is inversely correlated with heart growth in mammals, low during hyperplastic fetal stages and high in adults.⁷ Consequently, inducing p38 α MAPK activity in mammalian fetal cardiomyocytes blocks proliferation.⁷ Similarly in embryonic zebrafish, induction of p38 α MAPK activity blocks cardiomyocyte proliferation, resulting in severely disrupted cardiogenesis. Conversely, expression of dnp38a MAPK has no observable effects during embryonic cardiogenesis. These findings mirror previous reports in mammals and indicate that during zebrafish embryonic development, p38\alpha MAPK activity must be switched off (or be absent) for normal cardiomyocyte proliferation to occur. During heart regeneration in adult zebrafish, we found that induction of p38a MAPK activity effectively blocks this process; however, inhibiting p38a MAPK activity had no observable effects. This highlights the fact that p38a MAPK activity must be switched off in order for adult zebrafish cardiomyocytes to proliferate and, ultimately, regenerate the heart. Conversely, expression of dnp38a MAPK does not lead to unrestricted proliferation as one might have expected, indicating that other mechanisms are also involved in regulating cardiomyocyte proliferation. Interestingly, we also found that when cardiomyocyte proliferation is blocked, this essentially also blocks myocardial regeneration. It would appear then, that there is no back up mechanism for heart regeneration in zebrafish. If stem/progenitor cells are present in the zebrafish heart, then they are incapable of producing any substantial regenerative response when cardiomyocyte proliferation is blocked. We also found that active p38\alpha MAPK is present in adult zebrafish cardiomyocytes both in vitro and in vivo and only disappears when the cells enter mitosis. This suggests that p38a MAPK seems to be involved in entry into the M phase of the cell cycle. In adult mammals, active p38\alpha MAPK is known to induce arrest at various points of the cell cycle, including the G₂/M checkpoint.¹⁸ It appears then that active p38\alpha MAPK arrests both adult mammalian and zebrafish cardiomyocytes in the cell cycle, and once this inhibition is released, either pharmacologically in mammals or naturally in zebrafish, then cardiomyocytes are able to enter mitosis and, ultimately, proliferate. Our findings indicate that adult zebrafish and mammalian cardiomyocytes share a common p38a MAPKmediated molecular mechanism that regulates their proliferation. Although adult mammalian cardiomyocytes do not naturally proliferate, numerous reports have shown that they can be

induced to achieve this, which indicates that the proliferative machinery is present in these cells.⁴ While one could argue that mammals lack some extrinsic signal(s) necessary for inducing heart regeneration, results from neonatal mice indicate that these are present early in life.²⁰ If adult mammals lose/inhibit heart regeneration, then understanding how adult zebrafish regenerate their heart will, hopefully, one day allow us to reactivate this latent ability in adult mammals.

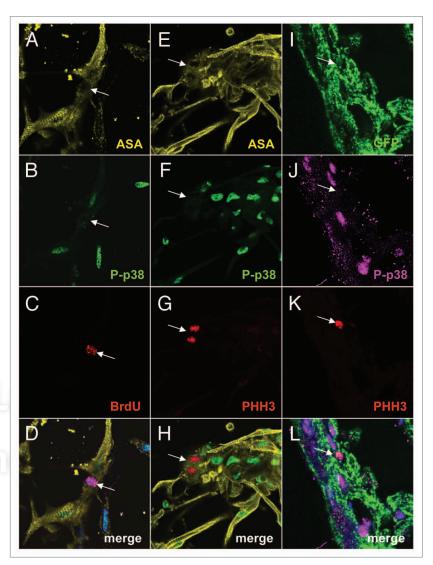


Figure 4. Adult zebrafish cardiomyocytes express active p38α MAPK, which is switched off upon entry into mitosis. Cultured adult zebrafish cardiomyocytes were immunolabeled for α *sarcomeric actin* (yellow) (A), phospho-p38α MAPK (green) (B), BrdU (red) (C) and a merged image with α *sarcomeric actin* (yellow), phospho-p38α MAPK (green), BrdU (red) and DAPI (blue) (D); the white arrow indicates the BrdU positive cardiomyocyte in (A–D). Cultured adult zebrafish cardiomyocytes were immunolabeled for α *sarcomeric actin* (yellow) (E), phosphor-p38α MAPK (green) (F), phospho histone H3 (PHH3) (red) (G) and a merged image with α *sarcomeric actin* (yellow), phospho-p38α MAPK (green), BrdU (red) and DAPI (blue) (H); the white arrow indicates the PHH3 positive cardiomyocytes in (E–H). A 10dpa *cmlc2a*-GFP regenerating heart immunolabeled for GFP (green) (I), phospho-p38α MAPK (magenta) (J) and phospho histone H3 (PHH3) (red) (K) and a merged image with GFP (green), phospho-p38α MAPK (magenta), phospho histone H3 (PHH3) (red) and DAPI (blue) (L); the white arrow indicates the PHH3 positive cardiomyocytes in (E–H). A 10dpa *cmlc2a*-GFP regenerating heart immunolabeled for GFP (green) (I), phospho-p38α MAPK (magenta) (J) and phospho histone H3 (PHH3) (red) (K) and a merged image with GFP (green), phospho-p38α MAPK (magenta), phospho histone H3 (PHH3) (red) and DAPI (blue) (L); the white arrow indicates the PHH3 positive cardiomyocytes in (I–L).

Materials and Methods

Constructs and zebrafish lines. All constructs and transgenic lines were generated using the Tol2 Kit as described in references 14 and 21. For the tg:cmlc2:LrL:dnp38 α MAPK construct, the 5' entry clone *cmlc2a* contained a 1 kb PCR fragment of the *cmcl2a* promoter;²² the middle entry clone contained a floxed RFP stop cassette amplified from pBOB-LRL-CBReGFPpA

(a kind gift from Geoff Whal), and the 3' entry clone contained murine dominant-negative p38 α MAPK (86.2% protein homology to zebrafish p38 α MAPK) (a kind gift from Roger Davis). The tg:cmlc2:LrL:caMKK6 was identical, except the 3' entry clone contained human constitutively active MKK6 (86.9% protein homology to zebrafish MKK6) (a kind gift from Roger Davis).

Cre/tamoxifen induction and heart amputation. Transgenes were expressed by inducing Cre-mediated recombination with tamoxifen as described in reference 14. All amputations were performed as described in reference 14.

BrdU labeling. BrdU treatment was performed essentially as described in references 17 and 23. Metamorph[®] software (Molecular Devices) was used to count BrdU labeled cardiomyocytes.

Cardiomyocyte isolation. Fish were sacrificed in tricaine. Hearts were collected and put into PBS with penicillin and streptomycin and 10 U/ml heparin. The outflow tracts were then removed, and ventricles and atriums were opened to get rid of the blood. They were then washed three times in perfusion buffer [PBS, 10 mM HEPES, 30 mM taurine, 5.5 mM glucose and 10 mM 2,3-butanedione monoxime (BDM)] and placed into digestion buffer [perfusion buffer plus 12.5 µM calcium chloride and collagenase, II and IV, 5 mg each (Gibco)] to digest for 2 h at 32°C in a thermomixer at 800 rpm. Next, an equal volume of stop buffer 1 (perfusion buffer plus 12.5 µM calcium chloride and 10% FBS) was added, and cells were mechanically separated. Undigested material was left to sediment, and cells suspended in the supernatant were pelleted by centrifugation at 250 g for 5 min. The cells were then resuspended in stop buffer 2 (perfusion buffer plus 12.5 µM and 5% FBS), and calcium reintroduction was performed by gradually raising the concentration to 62, 112, 212, 500 and 1,000 µM, respectively. Cells were then pelleted again and resuspended in plating medium (MEM, 5% FBS, 10 mM BDM, 2 mM Glutamax, 100 U/ml penicillin and 100 µg/ml streptomycin). Cell preparations were plated onto collagen type I

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(BD Biosciences, ref. 354236) treated chamber slides (Nunc, ref. 177445), and allowed to attach overnight. Immunofluorescence was performed the following day.

Immunohistochemistry. Immunohistochemistry was performed on 10 μ m cryo-sections as previously described in reference 1. The The antibodies used in this manuscript are anti-GFP (AVES GFP-1020), anti-BrdU (Accuratechemical OBT0030), anti-phopsho histone H3 (Upstate 06-570), anti-PCNA (Sigma P8825), anti- α sarcomeric actin (Sigma A2172), anti-RFP (Abcam ab34771), anti-Tropomyosin (Sigma T2780), anti-p-p38 (Thr180/Tyr182) (Cell Signaling #4511), anti-Mef2c (Abcam ab79436).

In situ hybridization. In situ hybridization was performed as described previously in reference 23. The *dnp38α MAPK* probe was generated by subcloning dominant-negative dnp38α MAPK into the *Hind111* and *BamH1* sites of pBSK⁻. The *caMKK6* probe was generated by subcloning constitutively active MKK6 into the *Hind111* and *Xba1* sites of pBSK⁻.

Confocal microscopy. Confocal microscopy was performed using a Leica SP5. For co-localization analysis the formula used to calculate the axial resolution is as follows.

 $\begin{array}{l} Dz = \sqrt{((l \ge n/NA^2) + (AU \ge n \ge \sqrt{2} \ge 1.22 \ge 1/NA^2)^2)} \\ Emission (l) = 500 nm \\ Refractive index 1.518 \\ NA = 1.4 \\ Airy Units: 1 \\ This results in a section thickness of z = 0.773 \ \mu m. \end{array}$

This results in a section three less of $z = 0.775 \,\mu \text{m}$

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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