



Published in final edited form as:

Circulation. 2019 January 22; 139(4): 559–561. doi:10.1161/CIRCULATIONAHA.117.033150.

C-Kit Cells Do Not Significantly Contribute To Cardiomyogenesis During Neonatal Heart Regeneration

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The neonatal mammalian heart has a remarkable regenerative ability for the first few days of life, mediated by the proliferation of preexisting cardiomyocytes (CM_S)¹. The contribution of preexisting cardiomyocytes to neonatal heart regeneration is supported by histological analysis of cardiomyocyte mitotic indices, as well as fate mapping studies using Myh6-driven inducible Cre. However, given that this fate mapping strategy labels approximately 70–80% of cardiomyocytes, it is plausible that there is additional contribution to the unlabeled population from non-cardiomyocytes.

To assess the contribution of C-kit⁺ cells to cardiomyogenesis in the neonatal heart after injury, we first crossed the inducible C-kit^{MerCreMer/+} mice² with the *Rosa26*^{tdTomato/tdTomato} reporter mice (Fig. 1A). One day postnatal mice (P1) were then injected with a single subcutaneous dose of 4-OH tamoxifen (100 µg/pup) to irreversibly label C-kit⁺ cells and their progeny. Subsequently, apical resection was performed at P2, and the hearts were harvested at P21 (Fig. 1B). All protocols were approved by the Institutional Animal Care and Use Committee of the University of Texas Southwestern Medical Center (UTSW). Aged-matched sham-operated animals were used as control. Hearts harvested at P2 showed an average of ~ 300 tdTomato⁺ cells per four-chamber view section (>90% of which were PECAM1⁺) (Fig. 1C). This pattern of colocalization with PECAM1 is consistent with previous observations which suggest that all C-kit⁺ cells represent a subpopulation of endothelial cells³. Similar to previous reports,¹ we found that the LV apex was regenerated by P21. Hearts were then cryosectioned in a four-chamber view, and stained for cardiac troponin T (Tnnt2) and tdTomato. We found that double positive cardiomyocytes (tdTomato

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Data Sharing:

The data, analytic methods, and study materials will be/have been made available to other researchers for purposes of reproducing the results or replicating the procedure by contacting the corresponding author.

Disclosures:

None

$^{+}/Tnnt2^{+}$) were rarely detected (Fig 1D), whether in sham or resected hearts, with an average number of 1.08 cell/section & 0.44 cell/section in the sham and resected hearts, respectively. These values indicate that less than 0.0015% of cardiomyocytes originated from a C-kit⁺ cells (Fig 1E). The percentage of tdTomato⁺/Tnnt2⁺ CM_s showed no statistically significant difference between sham and resected hearts, demonstrating that cardiac C-kit⁺ cells do not significantly contribute to cardiomyogenesis after neonatal cardiac injury. Of note, tdTomato⁺/Tnnt2⁺ CM_s seemed to be preferentially located in the base of the LV rather than the apex in both the sham or the resected group.

Next, C-kit^{Cre/+} mice, where Cre is knocked into the C-kit locus², were crossed with *Rosa26*^{tdTomato/tdTomato} reporter (Fig. 1F) and used to continuously label C-kit positive cells and their progeny. This strategy not only allows for detection of C-kit-derived cells during mid gestation onwards, but also labels cells that potentially express C-kit in response to injury irrespective of the presence of tamoxifen. As outlined above, apical resection was performed at P2, and the hearts were harvested at P21 (Fig. 1G). Aged-matched sham-operated animals were used as a control. Hearts harvested at P2 showed widespread tdTomato⁺ labeling (>90% of which were PECAM1⁺) (Fig. 1H). This extensive tdTomato⁺ labeling suggests that the constitutive cre-induced recombination labeled an early endothelial progenitor cell. Moreover, the high rate of co-localization with PECAM1 confirms the endothelial identity of cardiac C-kit⁺ cells, similar to what was observed in earlier reports³. While there was extensive labeling of tdTomato⁺ cells throughout the heart sections, less than 0.1% of cardiomyocytes (89 cells and 90 cells/section in sham and resected hearts respectively) were tdTomato⁺/Tnnt2 CM_s (Figs 1I,J). This low percentage of tdTomato⁺/Tnnt2⁺ CM_s further negates a significant role of c-kit⁺ cells in cardiomyogenesis. Furthermore, the lack of statistical significance between sham and resected hearts indicates that the neonatal injury does not induce further contribution of C-kit cells to myocardial lineage following neonatal injury (Fig 1J). Of note, tdTomato⁺/Tnnt2⁺ CM_s were found in colonies with a location preference towards the base of the LV and less frequently in the mid-heart section, and rarely in the apex (Fig 1J).

To determine whether the tdTomato⁺ cardiomyocytes were generated denovo from C-kit⁺ cells, we crossed the C-kit^{Cre/+} mice with the membrane-targeted tandem dimer Tomato/membrane-targeted green fluorescent protein (mT/mG) mice to obtain the compound heterozygotes (C-kit^{Cre/+}; membrane-targeted tandem dimer Tomato/membrane-targeted green fluorescent protein (mT/mG)) (Fig 1K). P21 hearts were then harvested, sectioned and examined by immunofluorescence. The percentages of tdTomato⁻/GFP⁺ novel cardiomyocytes and tdTomato⁺/GFP⁺ fusion-derived cardiomyocytes were about 0.01% and 0.005% respectively (Fig 1L,M). Expectedly, the average number of both the tdTomato⁻/GFP⁺ cells (novel cardiomyocytes) together with the tdTomato⁺/GFP⁺ cells (fusion events) was modestly lower than the number of tdTomato⁺/Tnnt2⁺ CM_s observed using the tdTomato reporter (Fig 1a) as previously reported³.

While a recent report suggested that C-kit cells may significantly contribute to neonatal heart regeneration⁴, the genetic model used in that study did not allow for lineage tracing, and the cryoinjury model used may have played a role in the discrepant results since cryoinjury is known to result in incomplete regeneration⁵. Taken together, our results

indicate that C-kit cells do not significantly contribute to cardiomyogenesis during early postnatal cardiac growth, or during neonatal heart regeneration following apical resection in mice.

Acknowledgments

Funding Sources:

H.S. is supported by grants from the NIH (1R01HL115275 and 5R01H2131778), National Aeronautics and Space Administration (NNX-15AE06G), American Heart Association (16EIA27740034), Cancer Prevention and Research Institute of Texas (RP160520), Hamon Center for Regenerative Science and Medicine, and Fondation Leducq.

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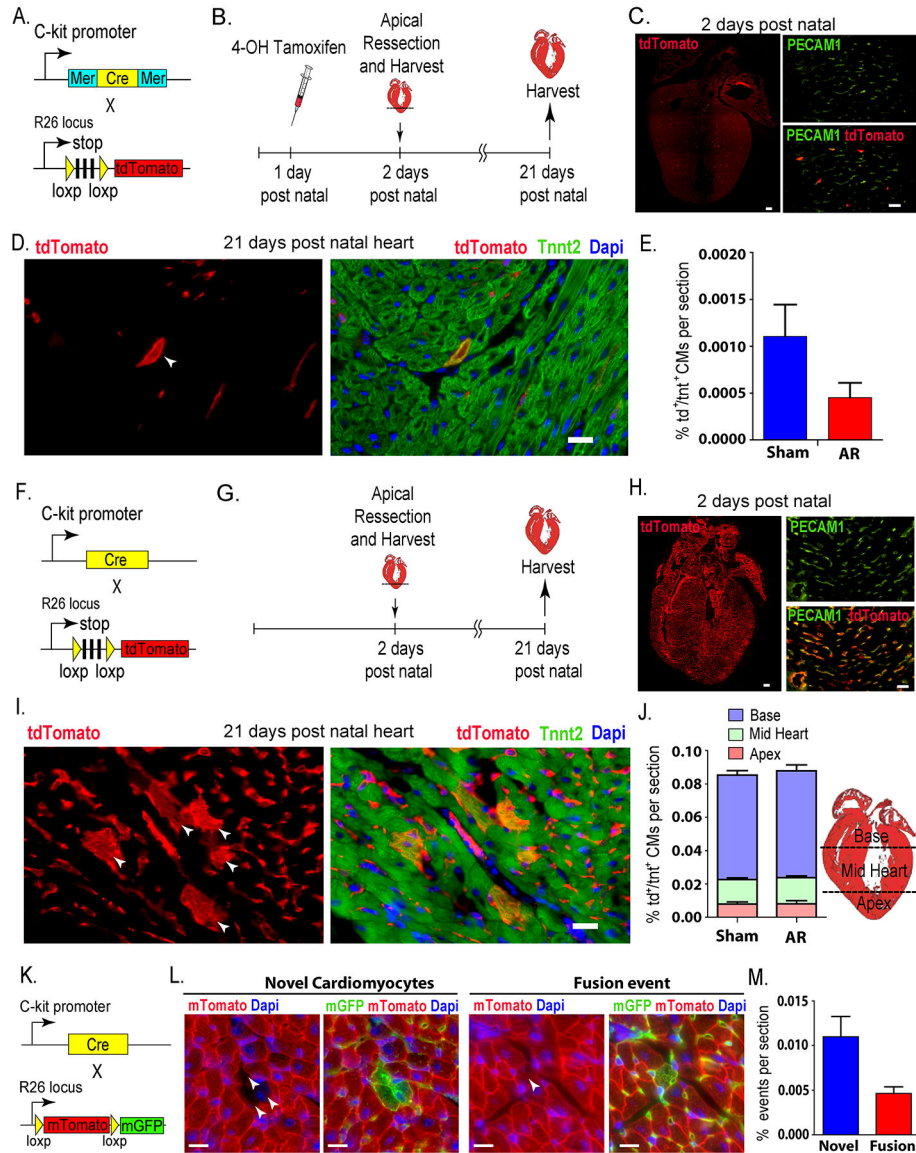


Figure 1: Role of C-kit cells in cardiomyogenesis in the neonatal mouse heart

(A) Schematic representing genetic mouse model of tamoxifen-dependent irreversible labeling of C-kit cells with tdTomato. (B) Time course of fate-mapping experiment. (C) **Left panel:** P2 whole heart showing ~ 300 tdTomato⁺ cell/section. Scale bar, 100 μ m (n=3). **Right panel:** more than 90% colocalization of tdTomato⁺ cells with PECAM1 staining. Scale bar, 20 μ m (D) **Left panel:** p21 heart (white arrowhead) showing a single tdTomato⁺ cardiomyocyte, **right panel:** showing the same cell to be tdTomato⁺/Tnnt2⁺. Scale bar, 20 μ m. (E) Graph showing no statistically significant difference in the percentage of be tdTomato⁺/Tnnt2⁺ CMs between sham and resected hearts (Sham: n=5 / AR: n=3). (F) Schematic representing genetic mouse model of continuous labelling of C-kit cells with tdTomato. (G) Time course of fate-mapping experiment. (H) **Left panel:** P2 whole heart scan showing widespread tdTomato⁺ signal. Scale bar, 100 μ m (n=10). **Right panel:** more than 90% colocalization of tdTomato with PECAM1 staining. Scale bar, 20 μ m. (I) **Left**

panel: p21 heart showing several arrowheads pointing to several tdTomato⁺ cardiomyocyte in colony, **right panel:** showing same cells to be tdTomato⁺/Tnnt2⁺. Scale bar, 20μm. **(J)** Graph showing no statistically significant difference in the percentage of tdTomato⁺/Tnnt2⁺ CM_s between sham and resected hearts, also differential percentage of the location of the cells in heart compartments (Sham: n=10 /AR: n=10). **(K)** Schematic representing the genetic cross of constitutive C-kit^{cre/+} with the reporter mTmG mice to enable detection of fusion events (n=10). **(L) Left panels:** p21 heart showing novel cardiomyocytes (tdTomato⁻/GFP⁺). **Right panels:** p21 heart showing fusion-derived cardiomyocytes (tdTomato⁻/GFP⁺). Scale bars, 20μm. **(M)** Graph showing percentage of real (novel cardiomyocytes) and fusion events.