RESEARCH HIGHLIGHT



Circulating macrophages as the mechanistic link between mosaic loss of Y-chromosome and cardiac disease

Xuehong Xu^{1*}, Rong Zhou¹⁺, Qinchun Duan¹⁺, Yuanlin Miao¹, Tingting Zhang¹, Mofei Wang¹, Odell D. Jones² and MengMeng Xu^{3*}

Abstract

Background Genetics evidences have long linked mosaic loss of Y-chromosome (mLOY) in peripheral leukocytes with a wide range of male age-associated diseases. However, a lack of cellular and molecular mechanistic explanations for this link has limited further investigation into the relationship between mLOY and male age-related disease. Excitingly, Sano et al. have provided the first piece of evidence directly linking mLOY to cardiac fibrosis through mLOY enriched profibrotic transforming growth factor $\beta 1$ (TGF- $\beta 1$) regulons in hematopoietic macrophages along with suppressed interleukin-1 β (IL-1 β) proinflammatory regulons. The results of this novel finding can be extrapolated to other disease related to mLOY, such as cancer, cardiac disease, and age-related macular degeneration.

Results Sano et al. used a CRISPR-Cas9 gRNAs gene editing induced Y-chromosome ablation mouse model to assess results of a UK biobank prospective analysis implicating the Y-chromosome in male age-related disease. Using this in vivo model, Sano et al. showed that hematopoietic mLOY accelerated cardiac fibrosis and heart failure in male mice through profibrotic pathways. This process was linked to monocyte-macrophage differentiation during hematopoietic development. Mice confirmed to have mLOY in leukocytes, by loss of Y-chromosome genes Kdm5d, Uty, Eif2s3y, and Ddx3y, at similar percentages to the human population were shown to have accelerated rates of interstitial and perivascular fibrosis and abnormal echocardiograms. These mice also recovered poorly from the transverse aortic constriction (TAC) model of heart failure and developed left ventricular dysfunction at higher rates. This was attributed to aberrant proliferation of cardiac MEF-SK4 + fibroblasts promoted by mLOY macrophages localized to heart and eventually resulted in cardiac fibrosis via enhanced TGF- β 1 and suppressed IL-1 β signaling. Furthermore, treatment of mLOY mice with TGF β 1 neutralizing antibody was able to improve their cardiac function. This study by Sano et al. was able to provide a causative link between the known association between mLOY and male cardiac disease morbidity and mortality for the first time, and thereby provide a new target for improving human health.

Conclusions Using a CRISPR-Cas9 induced Y-chromosome ablation mouse model, Sano et al. has proven mosaic loss of Y-chromosome in peripheral myeloid cells to have a causative effect on male mobility and mortality due

[†]Rong Zhou and Qinchun Duan are contributed equally to this work.

*Correspondence: Xuehong Xu xhx0708@snnu.edu.cn; xhx070862@163.com MengMeng Xu mex9002@nyp.org Full list of author information is available at the end of the article



© The Author(s) 2023. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

to male age-related cardiac disease. They traced the mechanism of this effect to hyper-expression of the profibrotic TGF-β1 and reduced pro-inflammatory IL-1β signaling, attenuation of which could provide another potential strategy in improving outcomes against age-related diseases in men.

Keywords Mosaic loss of Y-chromosome (mLOY), Cardiac fibrosis, Hematopoietic macrophage, Profibrotic regulons, Proliferation of cardiac fibroblast, Age-related cardiac disease

A correlation between peripheral leukocyte mosaic loss of Y-chromosome (mLOY) and many life-threatening diseases *i.e.* age-related diseases and oncologic diseases including colorectal, bladder, lung, and androgen associated cancers such as prostate and testicular germ cell tumors in men has been revealed for over a decade [1-7]. These diseases cover a wide range of organs, from agerelated macular degeneration to multiple oncogenic tissues [1-7]. MLOY had also been implicated in hypertension and cardiovascular disease, the leading cause of death in men [8, 9]. Cardiac fibrosis alone causes more than 800,000 deaths annually and plays a key role in all cardiovascular disease (CVD), which is responsible for over 17.5 million annual deaths worldwide [10]. Although significant resources have been directed at CVD over the past four decades, there is still a lack of effective treatments targeting the fibrotic process and a lack of understanding behind the mechanism of cardiac fibrosis. Here we highlight a recent study identifying a mechanism behind accelerated cardiac fibrosis in men with leukocyte mLOY.

Using a mouse model of Y-chromosome deletion to confirm UK Biobank prospective study findings of increased all-cause and cardiac morbidity and mortality in men with a higher proportion of mLOY (UK Biobank resource/application no. 18,623, 21,552 and 61,272) [11, 12], an international collaborative revealed the causal and mechanistic relationships behind mLOY and cardiac fibrosis and mortality [11, 13]. Using CRISPR-Cas9 to target repeat sequences around the Y chromosome centromere using the Y chromosome specific guide RNAs (gRNAs- empty control, LOY-gRNA1- Y specific guide one and LOY-gRNA2-Y specific guide two) and tomato red fluorescent protein (tRFP) marker, Sano et al. generated hematopoietic Y chromosome ablation mice generated by lentivirus transduction. Bone marrow cells successfully transduced by the lentivirus were selected by their tRFP positivity and collected from the reconstituted mice [13, 14]. Using this approach, the efficiency of Y chromosome ablation reached 95% for LOY-gRNA1 and 80% for LOY-gRNA2 as detected by tRFP. This model was able to match the percentage of mLOY white blood cells (as) found in the human population of 49-81% [11, 13]. The validity of their model was confirmed by a lack of expression of Y-chromosome genes Kdm5d, Uty, Eif2s3y, and Ddx3y in tRFP+leukocytes. They were then able to show that mLOY mice had accelerated age-related cardiomyopathy and echocardiogram findings consistent with heart failure. These mLOY mice also had poorer outcomes following the transverse aortic constriction (TAC) technique, a commonly used model of non-ischemic heart failure.

Compared to wildtype mice after TAC, TAC mLOY mice had significantly increased heart weight to tibia length ratio, atrial natriuretic peptide expression, and interstitial and perivascular fibrosis, all indicating advanced heart failure. TAC mLOY mice were found to have aberrant proliferation of cardiac MEF-SK4+fibroblasts. This along with no change in the number or size of cardiomyocytes, indicted that mLOY related heart failure is through increased cardiac fibrosis. These findings were consistent with UK Biobank data indicating an increased risk of death due to cardiovascular disease in men with a high proportional of mLOY leukocytes [11, 13]. Several studies have previously linked LOY to coronary artery disease (CAD). In early 2012, an inheritance cohort study proved Y chromosome function to be strongly linked to CAD [15]. Using 11 genetic markers specific to the Y chromosome, Charchar et al. investigated the correlation between CAD and the Y chromosome in 3233 British men included in the British Heart Foundation Family Heart Study (BHF-FHS), West of Scotland Coronary Prevention Study (WOSCOPS), and Cardiogenics Study. Using the data from these three cohorts, they were able to unveil Y chromosome polymorphisms strongly associated with CAD [15]. Even accounting for conventional CAD risk factors such as BMI, hypertension, and hypercholesterolemia, male-specific region of Y chromosome (MSY haplogroup I) has been associated with ~ 50% increase in CAD risk compared to other Y chromosome lineages [15]. Furthermore, through the Cardiogenics cohort, they identified this relationship to be mediated through dysregulation of immunity and inflammation facilitated by circulating hematogenous macrophages. Their transcriptome analysis isolated 19 pathways with strong differential expression including down-regulated pathway related to haemopoietic cell lineage and up-regulated pathways strongly associated with high risk of heart dysfunctions such as arrhythmogenic right ventriclar cardiomyopathy, dilated cardiomyopathy

and hypertrophic cardiomyopathy [15]. These findings were later confirmed by Eales et al., which analyzed over 129,000 men from the UK Biobank and confirmed that MSY haplogroup I had a 11% increased risk of CAD in comparison to all other haplotypes combined. They also found expression of the Y chromosome gene, *UTY*, as the only hematogenous gene expression associated with MSY haplogroup I. Macrophage involvement was confirmed when experimental reduction of *UTY* expression in macrophages to significantly reduce the immune costimulatory signal important in both early and late stage atherosclerosis development [16].

Building off these findings, using gene network analysis on single cell regulatory network inference and clustering (SCENIC), Sano et al. were able to provide exact mechanistic evidence to how mLOY in leukocytes were able to affect cardiac function [11, 13]. During development of the hematopoietic macrophages-monocyte lineage, they found mLOY leukocytes to be simultaneously enriched in profibrotic regulons of TGF- β 1 signaling along with suppressed IL-1 β proinflammatory regulons. This results in circulating pro-fibrotic macrophages which localized to the heart during stress (natural aging process) and injury (TAC) and predisposing organ repair down the pro-fibrotic pathway.

These findings by Sano et al. can be added to by Hulsmans et al. who showed hematopoietic macrophage can localized to the heart and naturalize in the atrial-ventricle (AV) node by binding to conductive cardiomyocytes. These resident macrophages are critically involved in cardiac function and regulate electrophysiological activity of the heart via the gap-junction protein connexin 43 (Cx43) [18]. Following the natural connection between these two studies, it is reasonable to postulate that hematopoietic mLOY may also produce mLOY macrophages that are contributing to arrhythmias commonly found in patients with cardiac failure. This would suggest two separate pathways through which hematopoietic mLOY could lead to cardiac disease in men (Fig. 1): (1) mLOY leukocytes leads to pro-fibrotic regulons that promote cardiac fibrosis during times of stress and injury leading to cardiac dysfunction, and (2) mLOY macrophages localized to the AV node and disturb electroconductive gap-junctions, and leading to arrhythmia along with cardiac fibrosis syndrome.

The adult mammal heart possesses three distinct macrophage populations (or three waves) which originate from different embryonic progenitors, including primitive yolk sac macrophages on embryonic day 7.0, fetal liver monocyte progenitors on embryonic day 9.5, and the traditionally associated hematogenous monocytemacrophage system (Fig. 1) [19–22]. In the setting of cardiac ischemia/reperfusion injuries, bone marrow monocyte production increases to produce circulating monocyte populations, which in turn localize to a myocardia macrophage niche in which these monocytes support myocardial recovery. However, the macrophages localized during embryogenesis also play a role myocardial recover. Any LOY during initial embryogenic differentiation of these cells would naturally affect the function of all macrophages in this lineage. Thus LOY loss in either embryogenic or mature hematopoietic macrophage populations would logically lead to suboptimal macrophage function and poor cardiac recovery in the adult patient [23-25]. A recent study using long term in vivo tracing of yolk sac embryonic hematopoietic stem cells (HSCs) traced these cells to the aorta-gonad-mesonephros (AGM), umbilical vessels, and other extraembryonic tissues. Importantly, these cells retain some of their hematopoietic nature even in adult mouse [26]. Samokhvalov showed yolk sac cells expressing Runx1 at embryonic day 7.5 later develop into fetal lymphoid progenitors and adult HSCs [26]. Using in vivo genetic tracing Yokomizo et al. recent showed fetal yolk sac derived HSCs expressing the liver specific transcription factor hepatic leukaemia factor (HLF) to be minimally involved in fetal hematooiesis, and instead is key in developing post-gestational intra-arterial hematopoietic clusters [27]. These results have been further confirmed using the zebrafish model [25]. Further implicating the importance of embryonic HSC in adult hematopoiesis, using Calreticulin knock-down and embryonic macrophage depletion approaches, Wattrus et al. demonstrated embryonic macrophage to be essential quality controls for establishing adult hematopoiesis [28].

Although the exact link between mLOY macrophages and cardiac conduction system is currently unknown, findings by Hulsmans et al. have identified a possible link. They identified a macrophage niche underneath atrialventricle (AV) node that puts localized macrophage directly in contact with the electroconductive cells in cardiac conduction system [18]. They found elongated cardiac macrophages expressing connexin43 to be densely intersperse with conducting cell in the distal atrioventricular node. Cardiac macrophage experimentally coupled to spontaneously beating cardiomyocytes via connexin-43-containing gap junctions were found to depolarize in a synchronized fashion with cardiomyocytes. Importantly, changing the membrane potential of these macrophages affected the resting potential of the associated cardiomyocytes and affected their repolarization, and thus their conduction ability in situ. Furthermore, conditional deletion of connexin43 in macrophages and congenital lack of macrophages delayed atrioventricular conduction in mice, resulting in these mice developing progressive atrioventricular block. These findings implicate that



Hematogenesis and developmental monocyte-macrophage system

Fig. 1 Mechanism behind the association of mLOY with cardiac fibrosis through cardiac localization of hematogenic mLOY macrophages. Although hematopoietic stem cells originally begin in the embryonic yolk sac, then the liver and dorsal aorta, by adult life, hematopoiesis lies in the bone marrow. Thus, mosaic loss of Y-chromosome in peripheral leukocytes originate in marrow myeoloid progenetors. The resultant mLOY leukocytes are then distributed throughout the body, including in to the myocardium and atrial-ventricular node, where their enrichment of TGFb1 expression and suppression of IL-1b expression promotes pro-fibrotic response to stress resulting in heart failure and arrhythmia. AVN, Atrial-ventricle node; CFB, Central fibrous body; POI, Plane of insulation

macrophage abnormality alone without intrinsic cardiac abnormality can induce aberrant cardiac conductions.

The heart is normally made of multiple heterogeneous population of cells predominantly consisting of cardiomyocytes, endothelial cells, fibroblasts, pericytes, and smooth muscle cells. Although the immune system is not classically associated with the heart, many immune cells are also found a vital component of the healthy heart [29–31]. What all these cell types, including macrophage, have in common is their response to TGF- β signaling. TGF- β signaling affects all cell types during the cardiac injury, repair and remodeling process involved in CVD [32]. TGF- β signaling is Smad-dependent. The tetrameric TGF- β receptor complex is composed of two copies each of the type-I and type-II receptors. Phosphorylation on specific sites of the type-I receptors activates their kinase domains, resulting in phosphorylation of R-Smads (regulated-SMADs) such as Smad2 and Smad3, and triggering the formation of a trimeric Smad complex with R-Smads and co-Smad (common-Smads) such as Smad4. This cytosolic phosphorylated Smad complex then transverses nuclear pore-baskets to interact with various transcription regulators that further control the transcription of specific target genes involved in cardiac fibrosis and ischemic remodeling [33, 34] (Fig. 2).

In the heart, mLOY mediated cardiac disturbance of the electroconductive gap-junction protein could occur through TGF- β signaling. Multiple studies in other cell types, such as osteocytes, have shown TGF- β 1 mediated effects on cell–cell communication via gap-junction proteins such as connexin43 and N-cadherin [35–38]. In cultured osteocytes, Fykerud et al. demonstrated Smad ubiquitination regulatory factor-2 (Smurf2) to affect gapjunctional intercellular communication via endocytosis and degradation of connexin43 and connexin45 [39].

Although no publication currently reports how Smad-TGF- β 1 signaling regulates electroconductive gap-junction proteins in the heart, since macrophages have been shown to communicate with the AV node cardiomyocytes through connexin43, and connexin43 is known to be regulated by TGF- β signaling in other cell types, regulation of mLOY in macrophages through TGF- β signaling is possible. If this hold true, abnormal Smad-TGF β signaling can affect the cardiomyocyte in two ways: (1) mLOY macrophages preferentially activate the R-Smads complex to form transcription regulatory complexes that known to trigger the pro-fibrotic axis in cardiomyocytes, leading to preferential cardiac fibrosis of surrounding cardiomyocytes; and (2) mLOY in AV node-integrated macrophages cause pathological Smad-TGF- β 1 signaling, which activates the Smurf2 pathway in surrounding cardiomyocytes, leading to endocytosis and degradation of gap-junction proteins like connexin43, eventually leading

Although peripheral mosaic loss of Y-chromosome in leukocytes/myeloid cells have been correlated with many age-related diseases in men [1, 4, 5, 9], this is the first time an animal model of peripheral mLOY has definitively showed a causative relationship [11, 12]. Through their CRISPR-cas9 hemopoietic Y-chromosome deletion mouse model of mLOY, Sano et al. have shown the mechanism behind mLOY and cardiac disease to be increased cardiac fibrosis due to pro-fibrotic leukocytes generated by mLOY. This mechanistic understanding not only provides a potential therapeutic

to loss of vital gap junction signaling, resulting in inter-

rupted conduction and cardiac arrhythmia (Fig. 2).



Fig. 2 Hypothesized mechanism of *SARA*-Smurf2 associated Connexin43/45 endocytosis and degradation via aberration of TGF- β signaling may critically cause the mosaic loss of electroconductive gap-junction protein Connexin43/45 in aging men with mLOY. An anchoring protein *SARA* (*Smad anchor for receptor activation*) binds to TGF- β receptors and activate ubiquitylation via Smurf2 (Smad ubiquitination regulatory factor-2). The Clathrin-coated early Connexin43/45 endosome then combines with cytoplasmic lysosome generated by ER-Golgi complex to form the late endosome, in which the gap-junction protein Connexin43/45 undergo degradation. This results in disassembly of gap-junctions, which could result in impaired cardiac conduction and subsequent cardiac arrhythmia. ER, endoreticulum; GRPs, Gene regulatory proteins; R-Smads, receptor-regulated Smads; C-Smads, common-partner Smads

strategy against the top cause of mortality in men, but also provides a possible treatment against other disease related to mLOY, such as cancer and age related macular degeneration.

LOY is not limited to just macrophages in ageing humans. Forsberg et al. has shown a relationship between male carcinogenesis and progressive accumulation of LOY in all cell types with increasing age [7]. Likewise Thompson et al. have suggested that macrophage LOY is simply a biomarker of general genetic instability and goes in hand with mLOY throughout the body [18]. Logically, these concomitant mutations in the affected organ result in subsequent functional consequences and contribute to age-related illness like cancer and cardiac disease. However, the recovery of these already cells after an external insult such as ischemia or reperfusion likely requires the assistance of the immune system for clearance of dead cells and recovery of injured cells. Thus, macrophages LOY leading to reduced or aberrant function likely impair immune mediated repair of injured tissues. This theory is supported by Hulsmans et al. finding that genetically modified macrophages can localized to the heart and naturalize in the atrial-ventricle (AV) node to regulate electrophysiological activity [17]. This is further supported by Sano et al. findings clearing showing macrophage LOY negatively impacts healing [11].

Abbreviations

AV	Atrial-ventricle
AVN	Atrial-ventricle node
AGM	Aorta-gonad-mesonephros
BHF-FHS	British Heart Foundation Family Heart Study
CAD	Coronary artery disease
CVD	Cardiovascular disease
CFB	Central fibrous body
C-Smads	Common-partner Smads
EMP	Erythro-myeloid progenitor
ER	Endoreticulum
IL-1β	Interleukin-1β
GRPs	Gene regulatory proteins
HLF	Hepatic leukaemia factor
HSC	Hematopoietic stem cell
mloy	Mosaic loss of Y-chromosome
ncRBC	Nucleated red blood cell
MSY	Male-specific region of Y chromosome
pEndC	Primitive endothelial cell
POI	Plane of insulation
рМас	Pre-macrophage
pMeso	Primary mesoderm
PSC	PDGFRA + stromal cells
R-Smads	Receptor-regulated Smads
SARA	Smad anchor for receptor activation
Smurf2	Ubiquitination regulatory factor-2
SCENIC	Single cell regulatory network inference and clustering
TAC	Transverse aortic constriction
TGFβ1	Transforming growth factor β1
WOSCOPS	West of Scotland Coronary Prevention Study

Acknowledgements

Not applicable.

Author contributions

MMX and XHX conceived the study. XHX, DQ and MX prepared the manuscript. ZR, DQ, MY, ZT and WM collected and analyzed data from publications. MMX, ODJ and XHX edited the manuscript. MMX and XHX finalized the manuscript. All authors read and approved the final manuscript.

Funding

This work was supported by the National Natural Science Foundation of China (#31771377/31571273/31371256), the Foreign Distinguished Scientist Program (#MS2014SXSF038), the National Department of Education Central Universities Research Fund (#GK201301001/201701005/GERP-17-45), US Maryland Stem Cell Research Fund (2009MSCRFE008300), and the Outstanding Doctoral Thesis Fund (#X2014YB02/X2015YB05). The partial collection of references for this article is supported by the Medical Scientist Training Program (MSPT) NIH to MMX.

Availability of data and materials

Not applicable.

Declarations

Ethics approval and consent to participate

All applicable international, national, and/or institutional guidelines for the care and use of animals were followed in the discussed studies.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

Author details

¹Laboratory of Cell Biology, Genetics and Developmental Biology, Shaanxi Normal University College of Life Sciences and University Hospital Medical Center, 620 West Chang'an, Chang'an District, Xi'an 710119, China. ²University of Pennsylvania School of Medicine ULAR, Philadelphia, PA 19144, USA. ³Department of Pediatrics, Morgan Stanley Children's Hospital, Columbia University, 3959 Broadway, New York, NY 10032, USA.

Received: 23 November 2022 Accepted: 19 June 2023 Published online: 24 July 2023

References

- Grassmann F, International AMD Genomics Consortium (IAMDGC), Weber BHF, Veitia RA. Insights into the loss of the Y chromosome with age in control individuals and in patients with age-related macular degeneration using genotyping microarray data. Hum Genet. 2020;139(3):401–7. https://doi.org/10.1007/s00439-019-02029-1.
- Grassmann F, Kiel C, den Hollander AI, Weeks DE, Lotery A, Cipriani V, Weber BHF, International Age-related Macular Degeneration Genomics Consortium (IAMDGC). Y chromosome mosaicism is associated with age-related macular degeneration. Eur J Hum Genet. 2019;27(1):36–41. https://doi.org/10.1038/s41431-018-0238-8.
- Forsberg LA, Gisselsson D, Dumanski JP. Mosaicism in health and diseaseclones picking up speed. Nat Rev Genet. 2017;18(2):128–42. https://doi. org/10.1038/nrg.2016.145.
- Jobling MA, Tyler-Smith C. Human Y-chromosome variation in the genome-sequencing era. Nat Rev Genet. 2017;18(8):485–97. https://doi. org/10.1038/nrg.2017.36.
- Wright DJ, Day FR, Kerrison ND, Zink F, Cardona A, Sulem P, Thompson DJ, Sigurjonsdottir S, Gudbjartsson DF, Helgason A, et al. Genetic variants associated with mosaic Y chromosome loss highlight cell cycle genes and overlap with cancer susceptibility. Nat Genet. 2017;49(5):674–9. https://doi.org/10.1038/ng.3821.
- Thompson DJ, Genovese G, Halvardson J, Ulirsch JC, Wright DJ, Terao C, Davidsson OB, Day FR, Sulem P, Jiang Y, et al. Genetic predisposition to mosaic Y chromosome loss in blood. Nature. 2019;575(7784):652–7. https://doi.org/10.1038/s41586-019-1765-3.

- Forsberg LA, Rasi C, Malmqvist N, Davies H, Pasupulati S, Pakalapati G, Sandgren J, Diaz de Ståhl T, Zaghlool A, Giedraitis V, et al. Mosaic loss of chromosome Y in peripheral blood is associated with shorter survival and higher risk of cancer. Nat Genet. 2014;46(6):624–8. https://doi.org/10. 1038/ng.2966.
- Khan SI, Andrews KL, Jennings GL, Sampson AK, Chin-Dusting JPF. Y chromosome, hypertension and cardiovascular disease: is inflammation the answer? Int J Mol Sci. 2019;20(12):2892. https://doi.org/10.3390/ijms2 0122892.
- Stubbins RE, Najjar K, Holcomb VB, Hong J, Nuñez NP. Oestrogen alters adipocyte biology and protects female mice from adipocyte inflammation and insulin resistance. Diabetes Obes Metab. 2012;14(1):58–66. https://doi.org/10.1111/j.1463-1326.2011.01488.x.
- Murtha LA, Schuliga MJ, Mabotuwana NS, Hardy SA, Waters DW, Burgess JK, Knight DA, Boyle AJ. The processes and mechanisms of cardiac and pulmonary fibrosis. Front Physiol. 2017;8:777. https://doi.org/10.3389/ fphys.2017.00777.
- Sano S, Horitani K, Ogawa H, Halvardson J, Chavkin NW, Wang Y, Sano M, Mattisson J, Hata A, Danielsson M, et al. Hematopoietic loss of Y chromosome leads to cardiac fibrosis and heart failure mortality. Science. 2022;377(6603):292–7. https://doi.org/10.1126/science.abn3100.
- Loftfield E, Zhou W, Graubard BJ, Yeager M, Chanock SJ, Freedman ND, Machiela MJ. Predictors of mosaic chromosome Y loss and associations with mortality in the UK Biobank. Sci Rep. 2018;8(1):12316. https://doi. org/10.1038/s41598-018-30759-1.
- Zeiher A, Braun T. Mosaic loss of Y chromosome during aging. Science. 2022;377(6603):266–7. https://doi.org/10.1126/science.add0839.
- Sano S, Wang Y, Evans MA, Yura Y, Sano M, Ogawa H, Horitani K, Doviak H, Walsh K. Lentiviral CRISPR/Cas9-mediated genome editing for the study of hematopoietic cells in disease models. J Vis Exp. 2019. https://doi.org/ 10.3791/59977.
- Charchar FJ, Bloomer LD, Barnes TA, Cowley MJ, Nelson CP, Wang Y, Denniff M, Debiec R, Christofidou P, Nankervis S, et al. Inheritance of coronary artery disease in men: an analysis of the role of the Y chromosome. Lancet. 2012;379(9819):915–22. https://doi.org/10.1016/S0140-6736(11) 61453-0.
- Lusis AJ. Y-chromosome genetic variation associated with atherosclerosis and inflammation. Arterioscler Thromb Vasc Biol. 2019;39(11):2201–2. https://doi.org/10.1161/ATVBAHA.119.313369.
- Eales JM, Maan AA, Xu X, Michoel T, Hallast P, Batini C, Zadik D, Prestes PR, Molina E, Denniff M, et al. Human Y chromosome exerts pleiotropic effects on susceptibility to atherosclerosis. Arterioscler Thromb Vasc Biol. 2019;39(11):2386–401. https://doi.org/10.1161/ATVBAHA.119.312405.
- Hulsmans M, Clauss S, Xiao L, Aguirre AD, King KR, Hanley A, Hucker WJ, Wülfers EM, Seemann G, Courties G, et al. Macrophages facilitate electrical conduction in the heart. Cell. 2017;169(3):510–22. https://doi.org/10. 1016/j.cell.2017.03.050.
- Epelman S, Lavine KJ, Beaudin AE, Sojka DK, Carrero JA, Calderon B, Brija T, Gautier EL, Ivanov S, Satpathy AT, et al. Embryonic and adult-derived resident cardiac macrophages are maintained through distinct mechanisms at steady state and during inflammation. Immunity. 2014;40(1):91–104. https://doi.org/10.1016/j.immuni.2013.11.019.
- Epelman S, Lavine KJ, Randolph GJ. Origin and functions of tissue macrophages. Immunity. 2014;41(1):21–35. https://doi.org/10.1016/j.immuni. 2014.06.013.
- Svedberg FR, Guilliams M. Cellular origin of human cardiac macrophage populations. Nat Med. 2018;24(8):1091–2. https://doi.org/10.1038/ s41591-018-0143-2.
- Bajpai G, Schneider C, Wong N, Bredemeyer A, Hulsmans M, Nahrendorf M, Epelman S, Kreisel D, Liu Y, Itoh A, et al. The human heart contains distinct macrophage subsets with divergent origins and functions. Nat Med. 2018;24(8):1234–45. https://doi.org/10.1038/s41591-018-0059-x.
- Peet C, Ivetic A, Bromage DI, Shah AM. Cardiac monocytes and macrophages after myocardial infarction. Cardiovasc Res. 2020;116(6):1101– 12. https://doi.org/10.1093/cvr/cvz336.
- Deniset JF, Belke D, Lee WY, Jorch SK, Deppermann C, Hassanabad AF, Turnbull JD, Teng G, Rozich I, Hudspeth K, et al. Gata6⁺ pericardial cavity macrophages relocate to the injured heart and prevent cardiac fibrosis. Immunity. 2019;51(1):131-140.e5. https://doi.org/10.1016/j.immuni.2019. 06.010.

- Chandrakanthan V, Rorimpandey P, Zanini F, Chacon D, Olivier J, Joshi S, Kang YC, Knezevic K, Huang Y, Qiao Q, et al. Mesoderm-derived PDGFRA⁺ cells regulate the emergence of hematopoietic stem cells in the dorsal aorta. Nat Cell Biol. 2022;24(8):1211–25. https://doi.org/10.1038/ s41556-022-00955-3.
- Samokhvalov IM, Samokhvalova NI, Nishikawa S. Cell tracing shows the contribution of the yolk sac to adult haematopoiesis. Nature. 2007;446(7139):1056–61. https://doi.org/10.1038/nature05725.
- Yokomizo T, Ideue T, Morino-Koga S, Tham CY, Sato T, Takeda N, Kubota Y, Kurokawa M, Komatsu N, Ogawa M, et al. Independent origins of fetal liver haematopoietic stem and progenitor cells. Nature. 2022;609(7928):779–84. https://doi.org/10.1038/s41586-022-05203-0.
- Wattrus SJ, Smith ML, Rodrigues CP, Hagedorn EJ, Kim JW, Budnik B, Zon LI. Quality assurance of hematopoietic stem cells by macrophages determines stem cell clonality. Science. 2022;377(6613):1413–9. https:// doi.org/10.1126/science.abo4837.
- Lafuse WP, Wozniak DJ, Rajaram MVS. Role of cardiac macrophages on cardiac inflammation, fibrosis and tissue repair. Cells. 2020;10(1):51. https://doi.org/10.3390/cells10010051.
- Frangogiannis NG. Cardiac fibrosis: cell biological mechanisms, molecular pathways and therapeutic opportunities. Mol Aspects Med. 2019;65:70– 99. https://doi.org/10.1016/j.mam.2018.07.001.
- Hulsmans M, Sam F, Nahrendorf M. Monocyte and macrophage contributions to cardiac remodeling. J Mol Cell Cardiol. 2016;93:149–55. https:// doi.org/10.1016/j.yjmcc.2015.11.015.
- Dobaczewski M, Chen W, Frangogiannis NG. Transforming growth factor (TGF)-β signaling in cardiac remodeling. J Mol Cell Cardiol. 2011;51(4):600–6. https://doi.org/10.1016/j.yjmcc.2010.10.033.
- Aki S, Yoshioka K, Takuwa N, Takuwa Y. TGFβ receptor endocytosis and Smad signaling require synaptojanin1, PI3K-C2α-, and INPP4B-mediated phosphoinositide conversions. Mol Biol Cell. 2020;31(5):360–72. https:// doi.org/10.1091/mbc.E19-11-0662.
- Alberts B, Johnson A, Lewis J, Raff M, Roberts K, Walter P. Molecular biology of the cell, chapter 15. 5th ed. Abingdon: Garland Science; 2008.
- Liu W, Zhang D, Li X, Zheng L, Cui C, Cui Y, Sun J, Xie J, Zhou X. TGF-β1 facilitates cell-cell communication in osteocytes via connexin43- and pannexin1-dependent gap junctions. Cell Death Discov. 2019;5:141. https://doi.org/10.1038/s41420-019-0221-3.
- Yang Y, Liu W, Wei J, Cui Y, Zhang D, Xie J. Transforming growth factor-β1induced N-cadherin drives cell-cell communication through connexin43 in osteoblast lineage. Int J Oral Sci. 2021;13(1):15. https://doi.org/10.1038/ s41368-021-00119-3.
- Liu W, Cui Y, Sun J, Cai L, Xie J, Zhou X. Transforming growth factor-β1 up-regulates connexin43 expression in osteocytes via canonical Smaddependent signaling pathway. Biosci Rep. 2018;38(6):BSR20181678. https://doi.org/10.1042/BSR20181678.
- Kotini M, Barriga EH, Leslie J, Gentzel M, Rauschenberger V, Schambony A, Mayor R. Gap junction protein Connexin-43 is a direct transcriptional regulator of N-cadherin in vivo. Nat Commun. 2018;9:3846. https://doi. org/10.1038/s41467-018-06368-x.
- Fykerud TA, Kjenseth A, Schink KO, Sirnes S, Bruun J, Omori Y, Brech A, Rivedal E, Leithe E. Smad ubiquitination regulatory factor-2 controls gap junction intercellular communication by modulating endocytosis and degradation of connexin43. J Cell Sci. 2012;125(Pt 17):3966–76. https:// doi.org/10.1242/jcs.093500.A.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.