



OPEN ACCESS

EDITED AND REVIEWED BY

Jordi Pérez-Tur,
Spanish National Research Council (CSIC),
Spain

*CORRESPONDENCE

Xinxiu Xu,
✉ xinxiu.xu@vumc.org

RECEIVED 21 January 2025

ACCEPTED 12 February 2025

PUBLISHED 18 February 2025

CITATION

Xu X, Lo CW, Martin LJ, Han L and Xu D (2025) Editorial: Recent advances in causes, diagnosis, and therapeutics for congenital heart defects. *Front. Genet.* 16:1564492. doi: 10.3389/fgene.2025.1564492

COPYRIGHT

© 2025 Xu, Lo, Martin, Han and Xu. This is an open-access article distributed under the terms of the [Creative Commons Attribution License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Editorial: Recent advances in causes, diagnosis, and therapeutics for congenital heart defects

Xinxiu Xu^{1*}, Cecilia W. Lo², Lisa J. Martin^{3,4}, Lu Han^{5,6} and Dongzhu Xu⁷

¹Department of Pathology, Microbiology and Immunology, Vanderbilt University Medical Center, Nashville, TN, United States, ²Department of Pediatrics and Department of Developmental Biology, University of Pittsburgh, Pittsburgh, PA, United States, ³Division of Human Genetics, Cincinnati Children's Hospital Medical Center, Cincinnati, OH, United States, ⁴Department of Pediatrics, College of Medicine, University of Cincinnati, Cincinnati, OH, United States, ⁵Department of Pediatrics, Medical College of Wisconsin, Milwaukee, WI, United States, ⁶Division of Pediatric Cardiology, Herma Heart Institute, Children's Hospital of Wisconsin, Milwaukee, WI, United States, ⁷Department of Cardiology, Institute of Medicine, University of Tsukuba, Tsukuba, Japan

KEYWORDS

heart development, congenital heart disease, genetic causes, diagnosis, evidence-based therapies

Editorial on the Research Topic

Recent advances in causes, diagnosis, and therapeutics for congenital heart defects

Introduction

Congenital heart disease (CHD) is one of the most prevalent major birth defects, yet its causes remain largely unknown. Both genetic and environmental factors play a role. Animal and human induced pluripotent stem cell models have shown how these factors disrupt heart development (Liu et al., 2017; Xu et al., 2022), but the precise mechanisms in humans remain unclear.

Advanced genetic and genomic approaches have significantly improved CHD diagnosis and therapies, especially through prenatal genetic testing, enabling earlier and more accurate diagnosis and screening. As survival rates into adulthood improve, new research directions have emerged, including exploring the genetic basis of surgical outcomes and developing therapies to enhance the quality of life for CHD patients. The growing population of adults with CHD, who lacked access to modern genetic technologies during childhood, underscores the need for ongoing research and tailored medical care (Bhatt et al., 2015).

This Research Topic encompasses a total of 14 articles, including basic research studies, clinical case reports, and a mini review. The novel findings focus on both pediatric and adult CHD (ACHD), covering recent advances in the causes, diagnosis, and therapeutics of CHD. These studies collectively demonstrate that integrating genetic data with clinical

assessments provides crucial insights for developing precise diagnostic and therapeutic strategies.

New causes and biomarkers in heart development abnormalities

The identification of novel genetic markers and pathways is a pivotal advancement in CHD research. Several studies in this Research Topic revealed novel biomarkers and genetic factors in cardiac developmental abnormalities. [Hu et al.](#) have reported that the structural protein Sorbs2 promotes the development of the second heart field, with its deficiency resulting in atrial septal defects. Their continued research found increased cardiac macrophages in Sorbs2-deficient hearts, revealing a potential role for macrophages in responding to embryonic myocardial abnormalities. A significant breakthrough came from [Xu et al.](#), who identified Keratin 19 (Krt19) as a novel epicardial gene through cardiac single-cell mRNA sequencing analysis. This discovery provides valuable insights into epicardial contribution to embryonic and neonatal heart development, addressing a long-standing challenge in the field. Additionally, [Li et al.](#) identified novel markers associated with increased risk and pathogenesis of immune checkpoint inhibitor-associated myocarditis through serum autoantibody profiles, expanding our understanding of cardiac complications in immunotherapy.

Genetic diagnosis in human congenital heart disease

The application of advanced genetic and genomic technologies, such as microarray testing and next-generation sequencing (NGS), has revolutionized the diagnostic landscape of CHD. [Mascho et al.](#) found an association between left outflow tract obstruction and 5p deletion, with high mortality in the presence of additional copy number variants. Similarly, [Yu et al.](#) used microarray analysis to identify both *de novo* and inherited micro-CNVs at 16p13.11 in 21 Chinese patients with defective cardiac left-right patterning. Significant progress in understanding specific CHD subtypes emerged from several studies. [Li et al.](#) conducted whole exome analysis of 25 Patent foramen ovale (PFO) patients, identifying potential mutant genes like *LDLR*, *SDHC*, and *NKX2-5*, enhancing our understanding of PFO's genetic basis. [Chen et al.](#) explored the impact of a *KCNH2* missense variant on Long QT syndrome, revealing incomplete penetrance influenced by sex, potentially shedding light on the distinct penetrance behaviors and patterns of the *KCNH2* gene. In another study, [Wang et al.](#) studied complex arrhythmias in children with *RYR2* gene sequence variation, providing evidence for early detection and treatment. Additionally, [Shen et al.](#) reported a pathogenic *MEIS2* sequence variation in a patient with multiple conditions, expanding the CHD symptom spectrum associated with *MEIS2* sequence variations.

Potential therapeutics and new challenges

New research and advances in genetic diagnostics have led to evidence-based therapies. This Research Topic highlighted several promising therapeutic developments and identified critical challenges in CHD management. [Dyrka et al.](#) showed the efficacy and safety of long-term recombinant growth hormone treatment on aortic dimensions in a girl with Loeys-Dietz Syndrome. [Hiraya et al.](#) identified implications for diagnostic strategies and therapeutic approaches by performing genetic testing and human leukocyte antigen analysis in patients with hypertrophic cardiomyopathy and connective tissue diseases.

Improved management of CHD has increased the life expectancy of adults with CHD. Many adults missed out on genetic evaluations during childhood due to the lack of modern genetic technologies. [Oehlman et al.](#) highlighted the need for guidelines to enhance access to genetic services and improve medical management by studying how ACHD cardiologists offer genetic services. [Edwards et al.](#) revealed significant gaps in genetics-related care, especially for those ACHD patients with congenital and neurodevelopmental comorbidities, through a retrospective chart review.

The identification of neurological deficits in adult CHD patients, as reviewed by [Saric et al.](#), points to the necessity of integrating genomic data with clinical history for better therapeutic targeting. They also suggested that investigating the interactions among ciliary genetics, CHD, and neurodevelopment could improve therapeutic management.

Conclusion

In conclusion, this Research Topic advances our understanding of CHD by elucidating its genetic causes, improving diagnostics and therapeutics, and identifying gaps and challenges. These studies provide essential insights for greater diagnostic precision and new directions for enhancing CHD care and management. It also highlights multidisciplinary collaboration and leveraging of cutting-edge technologies will continue to drive innovations in CHD research, leading to better patient care and improved clinical outcomes.

Author contributions

XX: Writing—original draft, Writing—review and editing. CL: Writing—review and editing. LM: Writing—review and editing. LH: Writing—review and editing. DX: Writing—review and editing.

Funding

The author(s) declare that no financial support was received for the research, authorship, and/or publication of this article.

Acknowledgments

The authors thank all contributing authors, peer reviewers, and the editorial staff for their support.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

The author(s) declared that they were an editorial board member of Frontiers, at the time of submission. This had no impact on the peer review process and the final decision.

References

- Bhatt, A. B., Foster, E., Kuehl, K., Alpert, J., Brabeck, S., Crumb, S., et al. (2015). Congenital heart disease in the older adult: a scientific statement from the American Heart Association. *Circulation*. 26 (21), 1884–1931. doi:10.1161/CIR.0000000000000204
- Liu, X., Yagi, H., Saeed, S., Bais, A. S., Gabriel, G. C., Chen, Z., et al. (2017). The complex genetics of hypoplastic left heart syndrome. *Nat. Genet.* 49 (7), 1152–1159. doi:10.1038/ng.3870

Generative AI statement

The author(s) declare that no Generative AI was used in the creation of this manuscript.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

- Xu, X., Jin, K., Bais, A. S., Zhu, W., Yagi, H., Feinstein, T. N., et al. (2022). Uncompensated mitochondrial oxidative stress underlies heart failure in an iPSC-derived model of congenital heart disease. *Cell Stem Cell*. 5 (5), 840–855. doi:10.1016/j.stem.2022.03.003