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POSTER ABSTRACT PRESENTATIONS

SESSION TITLE: POSTER SESSION 2

Abstract 602: Mir-182-5p is a Conserved Downstream Effector of Tbx5 Involved in Heart Development and Arrhythmia in Zebrafish

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Abstract

Background: TBX5 mutations cause Holt-Oram syndrome (HOS) characterized by upper limb and cardiac malformations, but can also contribute to early-onset of atrial fibrillation. Focusing on miRNAs involved in TBX5 regulatory circuits with a cardiac relevant role, we identified miR-182-5p, belonging to miR-183 cluster, found upregulated in Tbx5-depleted hearts of mouse and zebrafish embryos.

Methods: To functionally analyse the miR-182-5p role in developing heart, miR-182-5p was dysregulated in zebrafish zygotes of Tg(Myl7:EGFP) and Tg(myl7:gCaMP) transgenic lines. To stably deregulate miR-182-5p in zebrafish heart we exploited the Gal4/UAS system to restrict the miR-182 expression into cardiac context. For physiological analyses we performed the mechanogram of cardiac contraction and electrocardiogram recording. To understand miR-182-5p downstream regulation, in silico analyses, followed by ddPCR/real-time quantifications on

dissected zebrafish hearts and rescue experiments both in transient and stable miR-182-5p overexpressing zebrafish embryos were performed.

Results: Depletion of Tbx5 from cardiomyocytes increased the expression of miR-182 cluster family that is controlled by Kruppel-like factor 4 (KLF4), a transcription factor repressed by Tbx5. Both transient and stable upregulation of miR-182 in zebrafish affect heart morphology, calcium handling and the onset of arrhythmia while its cardiac-specific downregulation decreases cardiac defects in zebrafish HOS hearts. Expression analyses on selected miR-182-5p putative targets revealed that several calcium channel proteins resulted downregulated in miR-182-5p overexpressing hearts. Transgenic zebrafish line stably overexpressing miR-182-5p in the heart manifested arrhythmia overtime with or without cardiac structural defects.

Conclusion: We identified miR-182-5p as a potential suitable target to interfere in the circuit between upstream genetic abnormalities and downstream effectors leading to arrhythmia occurrence.

Footnotes

Author Disclosures: **E. Guzzolino:** None. **M. Pellegrino:** None. **N. Ahuja:** None. **D. Garrity:** None. **R. D'Aurizio:** None. **M. Groth:** None. **M. Baumgart:** None. **C. Hatcher:** None. **A. Mercatanti:** None. **M. Evangelista:** None. **C. Ippolito:** None. **E. Tognoni:** None. **R. Fukuda:** None. **V. Lionetti:** None. **L. Pitto:** None.



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