Are microRNA useful to predict prognosis in acute heart failure?

Giuseppe Limongelli^{1,2,3}, Martina Caiazza², Daniele Masarone¹

¹Cardiomyopathies and Heart Failure Unit-Monaldi Hospital, Naples, Italy; ²Department of Cardiothoracic Sciences, Università della Campania "Luigi Vanvitelli", Naples, Italy; ³Institute of Cardiovascular Sciences, University College of London, London, UK

Correspondence to: Giuseppe Limongelli, MD, PhD. Department of Cardiothoracic Sciences, Università della Campania "Luigi Vanvitelli", Naples, Italy. Email: limongelligiuseppe@libero.it.

Comment on: van Boven N, Kardys I, van Vark LC, et al. Serially measured circulating microRNAs and adverse clinical outcomes in patients with acute heart failure. Eur J Heart Fail 2018;20:89-96.

Received: 08 January 2018; Accepted: 19 January 2018; Published: 07 February 2018. doi: 10.21037/jlpm.2018.01.11 **View this article at:** http://dx.doi.org/10.21037/jlpm.2018.01.11

Introduction

Acute heart failure (AHF) is a major cause of hospital admission in patients over 65 years of age (1). Despite improvements in the pharmacologic therapy, clinical outcome remain poor with 1-year mortality rates that reach 20–30% (2). The progression of HF is linked to numerous genetic and epigenetic factors (3), including small regions of highly conserved non-coding RNAs, called microRNAs (miRNAs), that are important regulators of gene expression and modulate numerous processes implied in the pathophysiology of HF (4,5).

miRNA discovery and function

In 1993 the first miRNA, lin-4 was discovered, in Caenorhabditis elegans (C. elegans) and the interaction between lin-4 and the complementary sequence in the 3' untranslated region (3'UTR) of lin-14mRNA was shown (6). miRNAs are a class of non-coding RNAs with a length of 19–25 nucleotides that are involved in the regulation of gene expression at the post-transcriptional level (7,8). They have a tissue-dependent expression and concentration-dependent effects in pathologically affected organs (9,10).

miRNAs in heart failure

Functional studies have shown that miRNAs have a leading role in the onset and progression of HF (11), contributing to hypertrophy, fibrosis and therefore to the remodeling of the left ventricle (12). They also seems useful indicators for diagnosis and prognosis of both acute and chronic HF (13).

miRNAs and bypertrophy

Many miRNAs are implicated in the pathogenesis of cardiac hypertrophy, however among these the main one is miR-1 (14). In an experimental model of aortic coarctation Sayed *et al.* has shown that miR-1 is down regulated before the onset of cardiomyocyte hypertrophy (15). Furthermore, Elia and coworkers have shown that miR-1 interacts with insulin-like growth factor 1 (IGF-1), IGF-1 receptor and twinfilin-1, favoring cardiac hypertrophy (16).

Finally, suppression by an adeno-associated virus (AAV) of miR-1 determinate a regression of cardiac hypertrophy (17). These results suggest that miR-1 it plays a role in the development of left ventricular hypertrophy and may represent a molecular target for its regression.

miRNAs and fibrosis

miR-21 and miR-29 were identified as those most implicated in cardiac fibrosis. miR-21 determines the activation of the MAPK signal in cardiac fibroblasts through the inhibition of the extracellular inhibitor of the sprout regulated kinase 1 (Spry1) (18). On the other hand, miR-29 interacts with the genes encoding the extracellular matrix (ECM) such as fibrillin, elastin and collagen, determining their inhibition. In fact, models *in vitro* have shown that a reduction of miR-29 is associated with an ECM alma, while its overexpression causes a reduction in cardiac fibrosis. These data have also been confirmed in vivo, since the

Page 2 of 3

overexpression of miR-29b attenuates the progression of cardiac fibrosis (19).

miRNAs and diagnosis of HF

Several studies have assessed the role of circulating miRNAs in the diagnosis of heart failure. For example, in some studies, miRNAs have been able to discriminate between patients with cardiac dyspnea and those with other causes of dyspnea (20). Other studies have also described a miRNAs pattern in patients with AHF, with reduced plasma levels of some miRNAs (e.g., miR-103, miR-142-3p, miR-30b and miR-342-3p), and the increase of others (e.g., miR-499)(21).

miRNAs and prognosis in HF

Some studies have evaluated the prognostic value of circulating miRNAs in patients with acute and chronic heart failure. Qiang *et al.* have shown that elevated plasma levels of miR-508a-5p were associated with increased mortality in patients with chronic HF due to non-ischemic dilated cardiomyopathy (22). Furthermore, Ovchinnikova *et al.* have recently shown that an early reduction of some miRNAs (such as miR-18a-5p, miR-26b-5p, miR-27a-3p) were predictive of prognosis in patients with AHF (23). Therefore, miRNAs can be used as circulating biomarkers for prognosis, alone or in association with other well known markers, such as natriuretic peptides.

In a recent study (24) the plasma levels of circulating miRNAs were serially evaluated in patients hospitalized for AHF. The primary endpoint of this study was a composite endpoint of all-cause mortality and hospital readmission for HF. The study enrolled 476 patients and all the various miRNAs involved in HF were measured. miR-1306-5p was found to be positively and independently associated with all-cause mortality and hospitalization. This association was independent of NT-proBNP levels. Furthermore, repeated measurements of miR-320a, miR-378a-3p, miR-423-5p and miR-1254 were associated with the primary endpoint after adjustment for sex and age, but not after further multivariable adjustment for clinical characteristics.

The strength of this study is the serial evaluation of the miRNA, and the finding that they were associated with clinical outcome, suggesting a role for serial miRNAs evaluation in the management of patient with AHF. We postulate that patients with a lower reduction of miRNAs plasma levels after the start of HF therapy may be considered at greater risk requiring a more aggressive

management, while patients with normalization of miRNAs plasma levels may be considered at lower risk and therefore require standard care therapy.

Moreover, miR-1306-5p plasma levels was associated with the primary endpoint independently of the natriuretic peptides, suggesting an independent role for the miRNAs especially in patients with overlapping causes of natriuretic peptides increase (e.g., comorbidities, such as hepatic cirrhosis with ascites, renal insufficiency, hypothyroidism).

Conclusions

miRNAs are emerging as responsible for a wide range of physiological and pathological processes.

In the cardiovascular arena, miRNAs are primarily implicated in the progression of HF through cardiac hypertrophy and fibrosis. Further research is needed, to confirm the promise regarding these small molecules, to translate the excitement from bench to bedside, and to confirm the role of biomarkers for a tailored management of patients with HF.

Acknowledgments

Funding: None.

Footnote

Provenance and Peer Review: This article was commissioned and reviewed by the Section Editor Dr. Guo-Ming Zhang (Department of Laboratory Medicine, Shuyang People's Hospital/Shuyang Affiliated Hospital of Xuzhou Medical University, Shuyang, China).

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at http://dx.doi. org/10.21037/jlpm.2018.01.11). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

Open Access Statement: This is an Open Access article distributed in accordance with the Creative Commons Attribution-NonCommercial-NoDerivs 4.0 International License (CC BY-NC-ND 4.0), which permits the non-

Journal of Laboratory and Precision Medicine, 2018

commercial replication and distribution of the article with the strict proviso that no changes or edits are made and the original work is properly cited (including links to both the formal publication through the relevant DOI and the license). See: https://creativecommons.org/licenses/by-nc-nd/4.0/.

References

- Townsend N, Nichols M, Scarborough P, et al. Cardiovascular disease in Europe--epidemiological update 2015;36:2696-705.
- Mebazaa A, Tolppanen H, Mueller C, et al. Acute heart failure and cardiogenic shock: a multidisciplinary practical guidance. Intensive Care Med 2016;42:147-63.
- 3. Dorn GW 2nd. The genomic architecture of sporadic heart failure. Circ Res 2011;108:1270-83.
- Van Rooij E, Olson EN. MicroRNA therapeutics for cardiovascular disease: Opportunities and obstacles. Nat Rev Drug Discov 2012;11:860-72.
- Montgomery RL, Hullinger TG, Semus HM, et al. Therapeutic inhibition of miR-208a improves cardiac function and survival during heart failure. Circulation 2011;124:1537-47.
- Lee RC, Feinbaum RL, Ambros V. The C. elegans heterochronic gene lin-4 encodes small RNAs with antisense complementarity to lin-14. Cell 1993;75:843-54.
- Bentwich I, Avniel A, Karov Y, et al. Identification of hundreds of conserved and nonconserved human microRNAs. Nat Genet 2005;37:766-70.
- 8. Bartel DP. MicroRNAs: genomics, biogenesis, mechanism, and function. Cell 2004;116:281-97.
- Kozomara A, Griffiths-Jones S. miRBase: annotating high confidence microRNAs using deep sequencing data. Nucleic Acids Res 2014;42:D68-73.
- Paul P, Chakraborty A, Sarkar D, et al. Interplay between miRNAs and human diseases. J Cell Physiol 2018;233:2007-18.
- Vegter EL, van der Meer P, de Windt LJ, et al. MicroRNAs in heart failure: from biomarker to target for therapy. Eur J Heart Fail 2016;18:457-68.
- 12. Divakaran V, Mann DL. The emerging role of microRNAs in cardiac remodeling and heart failure. Circ Res

doi: 10.21037/jlpm.2018.01.11

Cite this article as: Limongelli G, Caiazza M, Masarone D. Are microRNA useful to predict prognosis in acute heart failure? J Lab Precis Med 2018;3:14.

2008;103:1072-83.

- Kalozoumi G, Yacoub M, Sanoudou D. MicroRNAs in heart failure: small molecules with major impact. Glob Cardiol Sci Pract 2014;79-102.
- 14. Sayed D, Abdellatif M. MicroRNAs in development and disease. Physiol Rev 2011;91:827-87.
- Sayed D, Hong C, Chen IY, et al. MicroRNAs play an essential role in the development of cardiac hypertrophy. Circ Res 2007;100:416-24.
- Elia L, Contu R, Quintavalle M, et. Al. Reciprocal regulation of microRNA-1 and insulin-like growth factor-1 signal transduction cascade in cardiac and skeletal muscle in physiological and pathological conditions. Circulation 2009;120:2377-85.
- Karakikes I, Chaanine AH, Kang S, et al. Therapeutic cardiac-targeted delivery of miR-1 reverses pressure overload-induced cardiac hypertrophy and attenuates pathological remodeling. J Am Heart Assoc 2013;2:e000078.
- Thum T, Gross C, Fiedler J, et al. MicroRNA-21 contributes to myocardial disease by stimulating MAP kinase signalling in fibroblasts. Nature 2008;456:980-4.
- Thum T, Catalucci D, Bauersachs J. MicroRNAs: novel regulators in cardiac development and disease. Cardiovasc Res 2008;79:562-70.
- 20. Tijsen AJ, Creemers EE, Moerland PD, et al. MiR423-5p as a circulating biomarker for heart failure. Circ Res 2010;106:1035-39.
- Yan H, Ma F, Zhang Y, et al. miRNAs as biomarkers for diagnosis of heart failure: A systematic review and metaanalysis. Medicine (Baltimore) 2017;96:e6825.
- 22. Qiang L, Hong L, Ningfu W, et al. Expression of miR-126 and miR-508-5p in endothelial progenitor cells is associated with the prognosis of chronic heart failure patients. Int J Cardiol 2013;168:2082-8.
- 23. Ovchinnikova ES, Schmitter D, Vegter EL, et al. Signature of circulating microRNAs in patients with acute heart failure. Eur J Heart Fail 2016;18:414-23.
- 24. Van Boven N, Kardys I, Van Vark LC, et al. Serially measured circulating microRNAs and adverse clinical outcomes in patients with acute heart failure. Eur J Heart Fail 2018;20:89-96.