

# “microRNAs (miRNAs) targeting oxidant-antioxidant mechanisms in primary ciliary dyskinesia and cystic fibrosis: epigenetic insights into therapeutic biomarkers”

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Dear Editor,

Primary ciliary dyskinesia (PCD) is a rare disorder marked by abnormalities in ciliary ultrastructure and/or function which are associated with disrupted mucociliary clearance and give rise to recurrent or persistent respiratory tract infections, bronchiectasis, bronchitis as well as a broad spectrum of clinical manifestations including sinonasal disease, neonatal respiratory distress, infertility, laterality defects and heterotaxy.<sup>1</sup>

Cystic fibrosis (CF) is also classified as a rare disease and is characterized by a defective CF transmembrane conductance regulator (CFTR) gene, whereas more than 40 disease-causing genes have been implicated in PCD to date.<sup>2</sup> CFTR dysfunction is associated with altered homeostasis in various organs and biological systems, including the pancreas, salivary glands, gastrointestinal tract, reproductive organs, and the respiratory system.<sup>2,3</sup> Within the framework of pulmonary homeostasis, impairment of mucociliary clearance constitutes a crucial pathogenic mechanism in CF which is linked to a sustained hyperinflammatory milieu and progressive pulmonary injury. Additionally, a close interconnection between inflammatory responses and the oxidant-antioxidant equilibrium was evident, as demonstrated by the significant correlations observed recently in our study.<sup>4</sup> Accordingly, levels of the intracellular antioxidant enzyme glutathione peroxidase (GPx) were significantly associated with the pro-inflammatory cytokines interleukin-1 $\beta$  (IL-1 $\beta$ ), IL-6, and IL-8 in patients with PCD and CF.<sup>4</sup> From this perspective, the pro-inflammatory/oxidant milieu and cellular antioxidant mechanisms represent pivotal targets for potential therapeutic strategies in patients with PCD and CF.

CF treatment is guided by symptomatic/supportive therapeutic modalities and novel therapeutic strategies including ion channel-targeted therapies, small-molecule approaches, and gene-based therapies.<sup>2</sup> Meanwhile, PCD exhibits a comparatively lower prevalence in relation to CF, notably among Caucasian populations and its clinical management continues to be predominantly guided by CF-derived therapeutic frameworks.<sup>1,2</sup>

It is noteworthy that microRNAs (miRNAs), a class of small non-coding RNAs, have garnered increasing attention in recent years as a potential therapeutic strategy in CF, due to their roles in modulating physiological functions and mediating disease-related pathophysiological mechanisms.<sup>3</sup> Among these miRNAs, miR-155 is regarded as a pivotal pro-inflammatory miRNA in patients with CF, with its increased expression being related with elevated IL-8 expression. In this context, miR-155-based therapeutic approaches can represent a promising strategy to restrict inflammatory responses by attenuating IL-8 mediated inflammatory processes.<sup>5</sup> Given the functional crosstalk between oxidative stress and inflammation, miRNAs associated with oxidant-antioxidant homeostasis have the potential to provide a relevant conceptual framework for research in both PCD and CF. Mitochondrial dynamics are well recognized as being indispensable for the maintenance of oxidant-antioxidant balance. In this context, miRNAs targeting genes associated with mitochondrial dynamics were assessed using bioinformatic approaches based on miRNA-target prediction tools, including TargetScan, Diana-microT and miRDB in this current study. Consequently, hsa-miR-369-3p, hsa-miR-381-3p and hsa-miR-382-3p were identified as candidate



miRNAs with the potential to regulate key mitochondrial dynamics related genes, including mitofusin 1 (MFN1), MFN2, and optic atrophy 1 (OPA1).

In conclusion, a comprehensive understanding of miRNAs as epigenetic regulators of genes related to mitochondrial dynamics reflecting oxidant-antioxidant equilibrium, particularly hsa-miR-369-3p, hsa-miR-381-3p and hsa-miR-382-3p, may provide critical insights to improve therapeutic outcomes and offer substantial translational relevance in these rare diseases.

## ETHICAL DECLARATIONS

### Peer Review Process

This letter was externally peer-reviewed.

### Conflict of Interest

The author declare no conflicts of interest.

### Financial Disclosure

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### Author Contributions

The author is solely responsible for the entirety of conception, execution, analysis, and writing of the manuscript.

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