

Emerging Roles of Long Non-coding RNAs in Chromatin Organization and Gene Regulation

Bambarendage Silvestre*

Department of Medical Laboratory Sciences, AJA University of Medical Sciences, Tehran 1411718541, Iran

Introduction

Long non-coding RNAs (lncRNAs) are a diverse class of RNA molecules exceeding 200 nucleotides in length, which do not encode proteins but play critical roles in regulating gene expression. Initially considered transcriptional noise, lncRNAs have emerged as key regulators of various cellular processes, including chromatin organization and gene regulation. By interacting with DNA, RNA, and proteins, lncRNAs influence chromatin structure and gene expression patterns, thereby contributing to cellular differentiation, development, and disease. This review explores the emerging roles of lncRNAs in chromatin organization and gene regulation, highlighting their mechanisms of action and potential implications in health and disease [1].

Description

Long non-coding RNAs (lncRNAs) are pivotal in regulating chromatin organization and gene expression through various sophisticated mechanisms. These RNA molecules, which do not encode proteins, can recruit chromatin-modifying complexes to specific genomic loci, thereby influencing chromatin states and gene activity. For example, the lncRNA XIST is essential for X-chromosome inactivation, as it recruits the Polycomb repressive complex 2 (PRC2) to silence one X chromosome in female mammals. lncRNAs can also act as molecular guides, scaffolds, or decoys, directing the localization and activity of transcription factors and chromatin modifiers to regulate gene expression [2]. HOTAIR, another well-studied lncRNA, serves as a scaffold for PRC2 and LSD1, coordinating histone modifications that repress gene expression at target loci. Additionally, lncRNAs contribute to the spatial organization of the genome within the nucleus by forming nuclear substructures like paraspeckles, which are crucial for sequestering transcription factors and RNA-binding proteins to control gene expression. The lncRNA NEAT1 is integral to the formation of paraspeckles, influencing nuclear architecture and RNA processing. Furthermore, some lncRNAs can interfere with the transcription of nearby genes by their own transcriptional activity, either blocking or facilitating the transcriptional machinery's access to DNA. This multifaceted regulation by lncRNAs underscores their significant roles in cellular differentiation, development, and disease, making them a focal point for understanding gene regulation and potential therapeutic targets [3].

The roles of lncRNAs in chromatin organization and gene regulation are complex and multifaceted, reflecting their diverse mechanisms of action. By interacting with chromatin-modifying complexes, transcription factors, and other regulatory molecules, lncRNAs can fine-tune gene expression in response to developmental cues and environmental signals. The ability of lncRNAs to modulate chromatin states and nuclear architecture underscores their importance in maintaining cellular identity and function. Dysregulation of lncRNAs is implicated in various diseases, including cancers,

*Address for Correspondence: Bambarendage Silvestre, Department of Medical Laboratory Sciences, AJA University of Medical Sciences, Tehran 1411718541, Iran; E-mail: bambarendage@silvestre.ir

Copyright: © 2024 Silvestre B. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution and reproduction in any medium, provided the original author and source are credited.

Received: 29 March, 2024, Manuscript No. MBL-24-137708; **Editor Assigned:** 01 April, 2024, PreQC No. P-137708; **Reviewed:** 15 April, 2024, QC No. Q-137708; **Revised:** 20 April, 2024, Manuscript No. R-137708; **Published:** 29 April 2024, DOI: 10.37421/2168-9547.2024.13.433

neurodegenerative disorders, and cardiovascular diseases, highlighting their potential as therapeutic targets [4].

Research into lncRNAs is rapidly expanding, revealing new insights into their functions and mechanisms. Advances in high-throughput sequencing and genome editing technologies have enabled the identification and characterization of numerous lncRNAs, providing a deeper understanding of their roles in gene regulation and chromatin organization. Moreover, the discovery of lncRNA-mediated regulatory networks offers promising avenues for developing novel therapeutic strategies. Targeting lncRNAs or their interacting partners could provide innovative treatments for diseases associated with aberrant gene expression and chromatin dynamics [5].

Conclusion

Long non-coding RNAs are crucial regulators of chromatin organization and gene expression, influencing a wide array of biological processes. Their ability to interact with DNA, RNA, and proteins allows them to modulate chromatin states, gene transcription, and nuclear architecture. Understanding the mechanisms by which lncRNAs regulate these processes is essential for elucidating their roles in development and disease. Continued research into lncRNA functions and interactions will advance our knowledge of gene regulation and chromatin biology, potentially leading to novel therapeutic approaches for treating complex diseases associated with lncRNA dysregulation.

Acknowledgement

None.

Conflict of Interest

None.

References

- Mattick, John S., Paulo P. Amaral, Piero Carninci and Susan Carpenter, et al. "Long non-coding RNAs: Definitions, functions, challenges and recommendations." *Nat Rev Mol Cell Biol* 24 (2023): 430-447.
- Heidari, Reza, Mostafa Akbariomi, Yazdan Asgari and Diako Ebrahimi, et al. "A systematic review of long non-coding RNAs with a potential role in Breast Cancer." *Mutat Res Rev Mutat Res* 787 (2021): 108375.
- Plasek, Leah M., and Saba Valadkhan. "lncRNAs in T lymphocytes: RNA regulation at the heart of the immune response." *Am J Physiol Cell Physiol* 320 (2021): C415-C427.
- Vajari, Mahdi K., Susan Moradinasab, Amir-Mohammad Yousefi and Davoud Bashash. "Noncoding RNAs in diagnosis and prognosis of Graft-Versus-Host Disease (GVHD)." *J Cell Physiol* 237 (2022): 3480-3495.
- Giuliani, Bianca, Chiara Tordonato and Francesco Nicassio. "Mechanisms of long non-coding RNA in breast cancer." *Int J Mol Sci* 24 (2023): 4538.

How to cite this article: Silvestre, Bambarendage. "Emerging Roles of Long Non-coding RNAs in Chromatin Organization and Gene Regulation." *Mol Biol* 13 (2024): 433.