

Special Issue

The Epigenetic Roles of lncRNAs

Message from the Guest Editors

Attention has recently increased toward long non-coding RNAs (lncRNAs) because of their capacity to impact gene expression through diverse epigenetic pathways, in both cis and trans functions.

Epigenetics is a dynamic and heritable modification that happens without respect for the DNA sequence and provides a variable level of control over gene expression. Therefore, understanding the interplay between lncRNA expression patterns, localization, and associated epigenetic signatures as well as function can provide us with the information required for developing a possible combination of biomarkers and therapeutic targets.

In this Special Issue, “The Epigenetic Roles of lncRNAs”, for *Genes*, I am delighted to invite a compilation of articles exploring the epigenetic functions of lncRNAs. This Special Issue aims to investigate the many and developing functions of lncRNAs in influencing epigenetic processes, providing insight into their potential as essential controllers of gene expression and cellular function.

Guest Editors

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Deadline for manuscript submissions

closed (20 February 2025)

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About the Journal

Message from the Editor-in-Chief

Genes is central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fast-moving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised. Why not consider *Genes* for your next genetics paper?

Editor-in-Chief

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