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# **Gene Regulation and Bioinformatics**

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# **Message from the Guest Editors**

Gene regulation is a fundamental process that governs gene expression, which in turn plays a critical role in various biological processes. Bioinformatics has emerged as a powerful tool for understanding the complex mechanisms of gene regulation and identifying novel targets for therapeutic intervention. To date, the field of gene regulation has been transformed by the advent of bioinformatics, which provides powerful tools for analyzing large amounts of genomic data.

This Special Issue aims to explore the intersection of computational and biological sciences in the context of gene regulation. We invite submissions that use bioinformatics approaches to study gene regulation at all levels, from transcriptional to post-transcriptional regulation, and in all organisms, from bacteria to humans. We are particularly interested in studies that use integrative approaches to combine genomic, epigenomic, and transcriptomic data to gain insights into gene regulation. The Special Issue will cover a broad range of topics including, but not limited to, regulatory networks, epigenomics, non-coding RNAs, machine learning algorithms, and network-based analysis.











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## Message from the Editor-in-Chief

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