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Methods in MicroRNA: Biogenesis, Identification, Function and Applications

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Message from the Guest Editors

The past two decades have seen significant advances in microRNA (miRNA) research, in regards to their identification, mechanisms, and applications. Functional assessments of miRNAs have been carried out across various cellular levels, not limited to the transcriptome, epigenome, proteome, and metabolome. Information generated using state-of-the-art technologies tremendously advanced our understanding of disease mechanisms governed by miRNAs. This is a major leap forward in the era of personalized medicine. However, there is still a paucity of information that can fully explain the role of miRNAs in contributing to developmental and disease heterogeneity and complexity. To address this challenge, multi-disciplinary research efforts focusing on the combination of several validated and novel research methods need to be developed that will enable researchers to endorse the contribution of an miRNA to a certain phenotype.

In this Special Issue titled "Methods in MicroRNA: Biogenesis, Identification, Function and Applications", we welcome original research and review articles presenting miRNA-related methodologies.













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Editor-in-Chief

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Message from the Editor-in-Chief

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