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Advances in Profiling and Deciphering the Functional Role of RNA Modifications

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Deadline for manuscript
submissions:

closed (31 January 2021)

Message from the Guest Editors

In this Special Issue, we will consider outstanding reviews, research, or method manuscripts on the following topics:

- Experimental methods to profile RNA modifications
- Computational methods for the analysis of epitranscriptional data
- Insights on the functional role of epitranscriptional marks and their effectors
- Prevalence and functional role of epitranscriptional marks in diseases
- The crosstalk between epitranscriptional marks
- The crosstalk between the epitranscriptome and the epigenome



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Special Issue



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Message from the Editor-in-Chief

In the past years the growth of the epigenetic field has been outstanding, from here the need of a journal where to centralize all new information on the subject. The term epigenetics is now broadly used to indicate changes in gene functions that do not depend on changes in the sequence of DNA. *Epigenomes* covers all areas of DNA modification from single cell level to multicellular organism as well as the epigenetics on human pathologies and behavior.

Epigenomes (ISSN 2075-4655) is a fully peer-reviewed publication outlet with a rapid and economical route to open access publication. All articles are peer-reviewed and the editorial focus is on determining that the work is scientifically sound rather than trying to predict its future impact.

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