

Special Issue

Advances in Profiling and Deciphering the Functional Role of RNA Modifications

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

Guest Editors

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Deadline for manuscript submissions

closed (31 January 2021)



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About the Journal

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.

Editor-in-Chief

Prof. Dr. Ernesto Guccione

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Author Benefits

High Visibility:

indexed within Scopus, ESCI (Web of Science), PMC, PubMed, Embase, PubAg, CAPUS / SciFinder, and other databases.

Journal Rank:

JCR - Q2 (Genetics and Heredity) / CiteScore - Q2 (Biochemistry, Genetics and Molecular Biology (miscellaneous))

Rapid Publication:

manuscripts are peer-reviewed and a first decision is provided to authors approximately 25.5 days after submission; acceptance to publication is undertaken in 3.8 days (median values for papers published in this journal in the second half of 2025).