

Special Issue

Targeting Epigenetic Network in Cancer

Message from the Guest Editors

As many sporadic cancer cases have been shown to be the result of a variety of modifiable lifestyle factors, including poor dietary habits, sedentary lifestyle, obesity, cigarette smoking, and exogenous hormones, a dysregulated epigenetic code seems to play an important role in sporadic cancer development. Three main epigenetic processes, including DNA methylation, histone modifications, and non-coding RNA mechanisms, are concerted in the epigenetic network to determine the transcriptional activity of certain genes. This Special Issue plans to give an overview of the most recent advances in the field of medical epigenetics in cancer and its application as a potential source of cancer biomarkers and anti-cancer therapies. Potential topics include, but are not limited to: DNA methylation in cancer; Histone modifications in cancer; Non-coding RNA mechanisms in cancer; RNA methylation in cancer; Epigenetic cancer biomarkers; Epigenetic anti-cancer therapy; Epigenetic chemoprevention.

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Deadline for manuscript submissions

20 February 2025



International Journal of Molecular Sciences

an Open Access Journal
by MDPI

Impact Factor 4.9
CiteScore 8.1
Indexed in PubMed



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Molecular Sciences*
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Message from the Editor-in-Chief

The International Journal of Molecular Sciences (*IJMS*, ISSN 1422-0067) is an open access journal, which was established in 2000. The journal aims to provide a forum for scholarly research on a range of topics, including biochemistry, molecular and cell biology, molecular biophysics, molecular medicine, and all aspects of molecular research in chemistry. *IJMS* publishes both original research and review articles, and regularly publishes special issues to highlight advances at the cutting edge of research. We invite you to read recent articles published in *IJMS* and consider publishing your next paper with us.

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