



an Open Access Journal by MDPI

## Non-Coding RNA Methods

Collection Editors:

### **Dr. Piero Carninci**

Laboratory for Transcriptome  
Technology, RIKEN Center for  
Integrative Medical Sciences,  
Yokohama, Kanagawa 230-0045,  
Japan

### **Dr. Florent Hubé**

Transgenerational Epigenetics &  
Small RNA Biology, Sorbonne  
Université, CNRS, Laboratoire  
Biologie du Développement,  
Institut de Biologie Paris-Seine,  
UMR7622, 75005 Paris, France

### Message from the Collection Editors

Dear Colleagues,

Non-coding RNA (ncRNA) is a rapidly growing field in which new research methods are continuously appearing. With this Topical Collection, we aim to collect a selection of articles on newly developed techniques of special interest for researchers in the ncRNA field.

We will consider manuscripts that report on new experimental methods, as well as reviews of exceptional interest that focus on newly developed techniques for ncRNA research that include ncRNA analysis programs, ncRNA chromatography methods, ncRNA structure analysis, ncRNA function analysis, ncRNA by high-throughput screening, ncRNA database/annotation, ncRNA imaging, and methods to detect the interactome of RNA (with proteins, with chromatin, etc.).

For further reading, please visit the *Topical Collection website*.

Best regards,

Dr. Piero Carninci  
Dr. Florent Hubé  
*Collection Editors*



[mdpi.com/si/7839](https://mdpi.com/si/7839)



an Open Access Journal by MDPI

## Editor-in-Chief

### **Prof. Dr. George A. Calin**

Department of Translational  
Molecular Pathology, Center for  
RNA Interference and Non-  
Coding RNAs, University of Texas  
MD Anderson Cancer Center,  
Houston, TX 77030, USA

## Message from the Editor-in-Chief

This field finally has a dedicated journal where its broad community can communicate and exchange its latest findings in one centralized place. This field was built stone by stone from the many scientific contributions from extremely diverse horizons, studying gene silencing in plants, position effect variegation in drosophila or quelling in fungi. This field has achieved maturity, but a lot remains to be discovered! Our aim is to publish manuscripts from all horizons that will have a high impact on the development of the field. Let's have fun and wish *Non-Coding RNA* a long and rewarding life!

## Author Benefits

**Open Access:**— free for readers, with **article processing charges (APC)** paid by authors or their institutions.

**High Visibility:** indexed within **Scopus, ESCI (Web of Science), PubMed, PMC, CAPlus / SciFinder,** and **other databases.**

**Journal Rank:** JCR - Q2 (*Genetics and Heredity*) / CiteScore - Q2 (*Genetics*)

## Contact Us

---

*Non-Coding RNA* Editorial Office  
MDPI, Grosspeteranlage 5  
4052 Basel, Switzerland

Tel: +41 61 683 77 34  
[www.mdpi.com](http://www.mdpi.com)

[mdpi.com/journal/ncrna](http://mdpi.com/journal/ncrna)  
[ncRNA@mdpi.com](mailto:ncRNA@mdpi.com)  
[X@Non\\_Coding\\_RNA](https://twitter.com/Non_Coding_RNA)