



an Open Access Journal by MDPI

The Role of RNA Processing and Metabolism in Tumors

Guest Editors:

Prof. Dr. Chunjiang He

College of Biomedicine and Health, Huazhong Agricultural University, Wuhan 430070, China

Dr. Ke Chen

Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology, Wuhan, China

Dr. Xiang Zhou

 Fudan University Shanghai Cancer Center, Fudan University, Shanghai 200032, China
Institutes of Biomedical Sciences, Fudan University, Shanghai 200032, China

Deadline for manuscript submissions: closed (20 August 2022)



Message from the Guest Editors

Defects of RNA metabolism cause many human diseases, including cancer. The complete sequence of the human genome provided quite a surprise to many by revealing that more than 98% of the transcriptional output represents non-protein coding RNAs (ncRNAs). In addition to housekeeping ncRNAs (rRNAs, tRNAs, etc.), small regulatory RNAs (microRNAs, piRNAs, etc.), long noncoding RNAs (lncRNAs, >200nt), and circular RNAs are emerging as major population of eukaryotic transcripts with both reported and yet undiscovered roles in gene regulation. This special issue focuses on understanding how genetic/environmental factors affect RNA metabolism in humans and why dysregulation of this process causes cancer. Areas of interest include, but are not limited to:

1) Roles of noncoding RNAs in cancer development and therapy

2) Roles of noncoding RNAs in cancer metabolism and tumour microenvironment

3) Noncoding RNAs as novel therapeutic targets

4) Biogenesis and new action modes of lncRNAs and circular RNAs

canc relssue

5) Epigenetic regulation of noncoding RNAs

6) Nucleolar stress and cancer therapy

7) Identification of ne





an Open Access Journal by MDPI

Editor-in-Chief

Prof. Dr. Selvarangan Ponnazhagan

Department of Pathology, The University of Alabama at Birmingham, 1825 University Blvd, SHEL 814, Birmingham, AL 35294-2182, USA

Message from the Editor-in-Chief

Genes are central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fastmoving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised.

Why not consider *Genes* for your next genetics paper?

Author Benefits

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

High Visibility: indexed within Scopus, SCIE (Web of Science), PubMed, MEDLINE, PMC, Embase, PubAg, and other databases.

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

Contact Us

Genes Editorial Office MDPI, St. Alban-Anlage 66 4052 Basel, Switzerland Tel: +41 61 683 77 34 www.mdpi.com mdpi.com/journal/genes genes@mdpi.com X@Genes_MDPI