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Bioinformatic Approaches in Cancer

Guest Editors:

Dr. Adriane F. Evangelista

Molecular Oncology Research Center, Barretos Cancer Hospital, Barretos 14784-400, Brazil

Dr. Rui Manuel Reis

 Molecular Oncology Research Center, Barretos Cancer Hospital, Barretos 14784400, SP, Brazil
Life and Health Sciences Research Institute (ICVS), School of Medicine, University of Minho, 4710057 Braga, Portugal
ICVS/3B's - PT Government Associate Laboratory, 4806909 Braga/Guimaraes, Portugal

Deadline for manuscript submissions: closed (20 May 2023)

Message from the Guest Editors

You are invited to contribute to a Special Issue of Genes that will focus on a Multi-omics Approach in Cancer. The advent of high-throughput technologies has enabled the development of methods for genome-wide characterization and detection of global transcript levels. Recently, bioinformatics tools have been developed and applied to the integration of the massive amount of data across multiple omics layers. Each type of omics data provides high-dimensional and meaningful information on the molecular mechanisms underlying the disease. An integrative approach provides a basis for the development of targeted cancer therapies.

In this Special Issue of *Genes*, we welcome reviews, new methods, and original research articles that apply different layers of multi-omics data (i.e., genome, transcriptome, epigenome, proteome, metabolome) for cancer research.

Specialsue

We look forward to receiving your contributions.



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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 is 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?

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Genes Editorial Office MDPI, Grosspeteranlage 5 4052 Basel, Switzerland Tel: +41 61 683 77 34 www.mdpi.com mdpi.com/journal/genes genes@mdpi.com X@Genes_MDPI