

## Special Issue

# Genome Informatics and Cancers

### Message from the Guest Editors

Genomic profiling has played a significant role in the rapid progress in our understanding of how cancers develop and progress, and contribute to precision in clinical care. However, the complexity and heterogeneity of cancer requires genomic profiling to be conducted along a number of orthogonal dimensions: single-cell to bulk tissue, model systems to human tissues, pediatric cancers to adult cancers, SNV to epigenetic to RNA profiling, etc. Such complex studies require an in-depth understanding of the opportunities and challenges of the complex heterogeneous data, developing associated analysis methods, and appropriate application of these methods to decipher the complexity of cancers. The journal *Cancers* invites articles in these broad areas of research for its Special Issue entitled “Genome Informatics in Cancers.” The articles may include original research, reviews, and perspectives on algorithms, statistical methods, software, and databases in all areas of cancer genomics (e.g., genomes, epigenomes, transcriptomes, regulomes, proteomes, and metabolomes).

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### Guest Editors

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### Deadline for manuscript submissions



## Cancers

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## About the Journal

### Message from the Editor-in-Chief

*Cancers* (ISSN 2072-6694) is an international, online journal addressing both clinical and basic science issues related to cancer research. The journal will continue its open access format, which will certainly evolve to ensure that the journal takes full advantage of the rapidly changing world of information and knowledge dissemination. It publishes high-quality clinical, translational, and basic science research on cancer prevention, initiation, progression, and treatment, as well as other related topics, particularly to capture the most seminal studies in the rapidly growing area of immunology, immunotherapy, and tumor microenvironment.

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### Editor-in-Chief

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