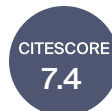


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Single-Cell Data Science (SCDS)

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Deadline for manuscript
submissions:

closed (15 September 2022)

Message from the Guest Editor

Single-cell sequencing technologies have greatly enhanced our ability to characterize heterogeneous tumor and normal cell populations to understand biological processes and identify novel therapeutic targets. Computational tools have been developed in recent years; however, analysis of single-cell sequencing data remains challenging and time-consuming, especially when integrating multiple types of data. This Special Issue of *Cancers* is devoted to papers and reviews on the latest advances and unprecedented opportunities in single-cell data science. We invite manuscripts from authors who are interested in developing novel tools, methods, and resources, as well as those who have new insights and applications to share in this area.



mdpi.com/si/56056

Special Issue



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Message from the Editor-in-Chief

Cancers is an international online journal addressing both clinical and basic science issues related to cancer research. The journal is publishing in 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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