



Molecular Genetics, Signaling Pathways, and Therapeutic Strategies for Liver Cancer

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Deadline for manuscript
submissions:

closed (31 October 2025)

Message from the Guest Editors

Dear Colleagues,

Liver cancer is a leading fatal human cancer. It is known to have a high degree of heterogeneity, leading to frequent treatment resistance and disease recurrence. There are a limited number of treatment options and they are not adequately effective. The hepatocarcinogenesis process of liver cancer is complex and involves the endeavor of malignant and immune components. There is an accumulation of various genetic and epigenetic alterations that pinpoint key signaling pathways or biological processes. With the emergence of different sequencing-based strategies, high-throughput screening platforms as well as computational algorithms, we can delineate the underlying molecular and cellular landscapes in liver cancer to a better extent. This Special Collection will focus on various niche areas that cover the basic as well as translational aspects of liver cancer research.

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Guest Editors





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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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