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# **Carcinogenesis as an Evolutionary Process**

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Deadline for manuscript submissions: closed (10 November 2022)

### Message from the Guest Editor

In the first phase of carcinogenesis, a cell may acquire a mutation that permits it to proliferate abnormally. In the next phase, other mutations allow for the expansion of cell number and eventually metastasize in distant organs. Analyses of the mutation landscapes and evolutionary trajectories of various tumor tissues have identified a constrained set of driver genes as the key genes whose mutation is most likely to initiate carcinogenesis. And their mutations are subject to positive selection during tumor evolution. Several types of approaches exploit this principle for the identification of genes that drive carcinogenesis.

As a result of combined efforts, providing a catalogue of driver mutations that could give rise to cancer has been achieved. However, a rather large fraction of cancer driver genes remain to be identified.

Another outstanding problem concerns the existence of tumor essential genes.

This Special Issue aims to focus on the toolkit of cancer driver genes and tumor essential genes. The significance of this question is that novel driver genes and tumor essential genes may prove to be valuable targets for cancer therapy.

**Special**sue



mdpi.com/si/119025





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### Message from the Editor-in-Chief

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