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Genetic Markers and Cancer Risk

Guest Editor:

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Deadline for manuscript submissions:
closed (31 December 2020)

Message from the Guest Editor

Dear Colleagues,

All types of cancers are influenced by the genetic background of the patient and population. The influence of genetic variants and mutations ranges from rare, highly penetrant mutations in cancer syndrome genes (*TP53*, *VHL*, *BRCA1*, *BRCA2* and others) to common, low impact variants identified by genome-wide association study. Identification of genetic markers in cancer risk is an active area. For this Special Issue of *Cancers*, papers that identify risk variants, seek to understand their function, study the frequency in specific cancers, or specific populations, or that address new methods to identify variants will be considered.

Dr. Michael Dean
Guest Editor

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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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