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Epidemiology of Genetic Variants Associated with Inherited Cancer Syndromes including Breast or Colon Cancer

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Message from the Guest Editors

After decades of research, the major genes causing inherited breast and colon cancers have been identified. From testing one or a few genes at a time based on family history, the testing system from now on will mostly be gene panel testing or whole genome sequencing (WGS) based on incident cancer cases or early-onset cancers. Such analyses will generate previously unknown information. Determining how to interpret this information needs compilation of knowledge. This Special Issue invites the publication of results of panel testing or WGS in cancer cases and families to increase knowledge. Such knowledge is needed both for healthcare providers, to plan research projects, and for the individual care of each carrier of the genetic variants in question.

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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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