



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

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Deadline for manuscript
submissions:

closed (20 October 2023)

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

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