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Genetic Regulation in Ovarian Cancer

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submissions:

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Message from the Guest Editors

Dear Colleagues,

Years of research progress, underpinned by large-scale advances made in studies published by The Cancer Genome Atlas (TCGA), has identified numerous candidate gene mutations in epithelial ovarian cancer. However, the therapeutic options for ovarian cancer patients have not dramatically advanced over the last two decades.

This Special Issue presents a collection of research reports and reviews in the areas of: (1) regulation and/or expression of *BRCA1/BRCA2* gene variants; (2) genetic hereditary syndromes, such as nonpolyposis colorectal cancer (HNPCC)/Lynch syndrome; (3) DNA methylome analyses; (4) chromatin modulation in ovarian cancer; and (5) regulatory RNAs (miRNA, lncRNA, eRNA) in ovarian cancer.

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Guest Editors



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Special Issue



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Message from the Editor-in-Chief

Genes are central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fastmoving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised.

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