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Insights in Genetic Medicine

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Message from the Guest Editor

Dear Colleagues,

Cancer is one of the leading causes of illness and death worldwide. Different clinical, epidemiological, molecular studies have shown advances in the basic knowledge of the biology of cancer, its diagnosis, and treatment. In this sense, the advances in knowledge of the variants in different metabolic pathways that participate in the processes and susceptibility to the development of cancer have been of great importance. Thus, it has been observed that the frequency of these variants depends to a large extent on the study population, so it is interesting to know data specific to each region. In Mexico, breast, colon, and lung cancer are among the main causes of death, which is why our group is interested in the great diversity of genes that participate in the evolution of cancer, as well as in its treatment. Among the genes that have attracted attention are, for example, the KRAS, ESR1, ESR2, ERBB2, TNF alpha, VEGF, eNOS, Tp53, RAD1, MTHFR, CBS, and TYMS genes.

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