



Expression of Signaling Genes in Cancer and Other Pathologies

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submissions:

closed (30 June 2024)

Message from the Guest Editor

Novel sequencing techniques now allow us to produce huge amounts of transcriptomic data. The accessibility of it is also greatly facilitated through public databases such as TCGA. Today, the scientific challenge is not to measure the gene expression level, but rather to interpret the biological meaning of it. The most reasonable approach seems to be to analyze entire functional genesets, since none of the proteins stands alone, they all work in well-orchestrated and incredibly complex cross-talk of cellular pathways.

This Special Issue is meant to present new findings on how alterations in gene expression and cell signaling drive pathogenesis of cancer and other complex conditions. We invite both high-quality research and review papers concerning broadly understood regulation of cellular signaling pathways.

Looking forward to your submission.





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Editor-in-Chief

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Message from the Editor-in-Chief

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