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Multi-omics Data Integration in Complex Diseases

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Deadline for manuscript submissions:

30 June 2024

Message from the Guest Editors

Dear Colleagues,

Each year, complex genetic diseases are responsible for the deaths of millions worldwide. Generally, genetic changes may have effects on multiple levels such as gene expression change, post-transcriptional dysregulation, protein expression change and mis-localization. Despite developments in large-scale genomics, transcriptome and proteomics technologies, it is still challenging to interpret molecular intricacies and variations at multiple molecular levels. Therefore, multi-omics-based data integration is the critical step to advance our understanding of molecular mechanisms for human disease.

Therefore, this Special Issue intends to cover manuscripts that provide valuable insight for both basic and clinical researcher communities. Particularly, we encouraged the submissions about: (1) molecular insights into the common and unique mechanism using multiple omics technologies; (2) novel computational frameworks for genetic variant classification and functional evaluation; (3) practical omics-guided diagnosis and prognosis treatment strategies.

Guest Editors













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