# **Special Issue**

# Research on Mathematical and Computational Modelling of Systems Biology

## Message from the Guest Editor

Recent breakthroughs in computational systems biology have revolutionized our comprehension of complex biological systems. Central to these advances are improved algorithms that facilitate the modelling of intricate biological networks, enabling accurate simulations of cellular processes. The integration of machine learning techniques has become pivotal in analyzing large-scale genomic and proteomic datasets, thus improving the identification of critical biomarkers and therapeutic targets. The fusion of multi-omics dataencompassing genomics, transcriptomics, proteomics, and metabolomics—offers a comprehensive perspective on cellular functions and interactions, providing unprecedented insights into biological processes. Highperformance computing has been instrumental in processing extensive datasets, enhancing the development of robust predictive models. Artificial intelligence is increasingly being used in drug discovery, thus expediting the identification of promising drug candidates. This Special Issue explores these cuttingedge developments and their profound implications for the future of biomedical research.

## **Guest Editor**

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## Deadline for manuscript submissions

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The journal *Mathematics* publishes high-quality, refereed papers that treat both pure and applied mathematics. The journal highlights articles devoted to the mathematical treatment of questions arising in physics, chemistry, biology, statistics, finance, computer science, engineering and sociology, particularly those that stress analytical/algebraic aspects and novel problems and their solutions. One of the missions of the journal is to serve mathematicians and scientists through the prompt publication of significant advances in any branch of science and technology, and to provide a forum for the discussion of new scientific developments.

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