



Systems Medicine and Bioinformatics

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Message from the Guest Editors

In precision medicine, determining the right treatment for the right patients based on a precise diagnosis is the ultimate goal. It guides us to provide considerable progress to our knowledge of disease etiology and pathogenesis. To achieve this goal, molecular features and clinical phenotypes are used to perform the correct stratification of patients and classify the corresponding features. Recent advances in sequencing and multi-omics technologies transform biomedical research and change healthcare and medicine by providing big data as molecular and clinical features. In parallel, advanced statistical, computational, and mathematical tools to analyze, integrate, interpret big heterogeneous datasets, and develop multilevel models are demanded.

We encourage the submission of original and review articles to this Special Issue of the Journal of Personalized Medicine that cover the "Systems medicine and bioinformatics". Articles focused on empirical and computational studies, implementation of new models along with the validations, focusing on the barriers to expanding systems medicine into personalized and precision medicine will be welcome.





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Message from the Editor-in-Chief

Journal of Personalized Medicine (JPM; ISSN 2075-4426) is an international, open access journal aimed at bringing all aspects of personalized medicine to one platform. *JPM* publishes cutting edge, innovative preclinical and translational scientific research and technologies related to personalized medicine (e.g., precision medicine, pharmacogenomics/proteomics, systems biology, ‘omics association analysis). *JPM* is covered in Scopus, the Science Citation Index Expanded (SCIE), PubMed, PMC, Embase, and other databases.

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