Cystic Fibrosis and Personalized Medicine

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Deadline for manuscript
submissions:
closed (25 November 2021)

Message from the Guest Editors

Cystic fibrosis is the most prevalent monogenetic disease worldwide. Over 2000 different defects in the cystic fibrosis transmembrane conductance regulator (CFTR) gene have been described, not only leading to a highly diverse phenotype of the disease, but also to the development of a plethora of gene-defect specific therapies. Many new techniques and models have been developed to unravel the genotype-phenotype relationship and to develop and apply new molecules which can modify CFTR-protein function. In addition to clinical and radiological evaluation, functional measurements such as sweat testing, intestinal current measurements, and patient-derived cell models from airway and gut tissue have become available. Today, these tools are more and more helpful in driving treatment of cystic fibrosis into theratyping and real personalized medicine. For this Special Issue, we invite review articles and original studies addressing the latest developments in patient-specific drug development and evaluation of effects in patients with cystic fibrosis.

mdpi.com/si/58483

Special Issue

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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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