

Special Issue

Analysis of Sequence Data from Bacterial and Human Epigenomes

Message from the Guest Editors

This Special Issue will focus on analyzing sequence data from bacterial and human epigenomes. The scope of this Special Issue includes novel biological techniques, sequencing data, epigenomic information, and analysis tools. It also includes the analysis of bacterial data in the context of infectious diseases and human data in the context of clinical cases. We invite the research community from both academia and industry to submit their work on bacterial and human epigenomes for this Special Issue. Even though the focus of the Issue is analyzing sequence-level data, we encourage colleagues to submit relevant research work that was performed using methods and technologies apart from sequencing. This Special Issue will provide a snapshot of current methods and analyses that are being used by the scientific community for understanding bacterial and human epigenomes. We anticipate that these research findings will reveal further insights into genome structure and function.

Guest Editors

Dr. Miten Jain

UCSC Genomics Institute, University of California, Santa Cruz, CA 95064, USA

Dr. Sarah L Castro-Wallace

NASA Johnson Space Center, Houston, TX, USA

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BioTech
Editorial Office
MDPI, Grosspeteranlage 5
4052 Basel, Switzerland
Tel: +41 61 683 77 34
biotech@mdpi.com

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Editor-in-Chief

Prof. Dr. Massimo Negrini
Department of Morphology, Surgery and Experimental Medicine,
University of Ferrara, Ferrara, Italy

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manuscripts are peer-reviewed and a first decision is provided to authors approximately 22.3 days after submission; acceptance to publication is undertaken in 2.8 days (median values for papers published in this journal in the first half of 2025).