



BioTech



an Open Access Journal by MDPI

Analysis of Sequence Data from Bacterial and Human Epigenomes

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

Deadline for manuscript
submissions:

closed (31 May 2021)

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.



mdpi.com/si/61395

Special Issue