

Special Issue

Research Advances in the Bioinformatics of Genome Editing and Gene Function Analysis

Message from the Guest Editor

We are inviting submissions for our upcoming Special Issue (SI) focusing on the bioinformatics of genome editing and gene function analysis. Genome editing technologies have become widely used and indispensable tools for gene function analysis. Additionally, there is a growing need to process vast amounts of base sequence information obtained through Next Generation Sequencing (NGS) and large-scale image data for functional analysis. However, the details of such bioinformatics research often remain under-represented and relegated to supplementary materials. In this SI, we aim to bring these critical discussions to the forefront. We welcome a broad range of papers that shine a light on this pivotal area of research. Join us in advancing the field of genome editing and gene function analysis through your valuable contributions.

Guest Editor

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Message from the Editor-in-Chief

The *International Journal of Molecular Sciences (IJMS)* is an open access journal, which was established in 2000. The journal aims to provide a forum for scholarly research on a range of topics, including biochemistry, molecular and cell biology, and molecular biophysics. *IJMS* publishes both original research and review articles, and regularly publishes special issues to highlight advances at the cutting edge of research. We invite you to read recent articles published in *IJMS* and consider publishing your next paper with us.

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