

Special Issue

Genetics and Genomics of Silkworm

Message from the Guest Editor

The silkworm (*Bombyx mori*) has long served as a pivotal model organism in genetics, developmental biology, evolutionary biology, physiology, agricultural biotechnology, etc. Recent advancements in high-throughput sequencing, CRISPR-based genome editing, and functional genomics have revolutionized our understanding of silkworm biology. This Special Issue aims to consolidate cutting-edge research and interdisciplinary insights into the genetics and genome of the silkworm, fostering innovation in both fundamental science and applied sericulture. **Suitable topics include, but are not limited to, the following:**

Suitable topics include, but are not limited to, the following:

- The silkworm genome encompasses the genomes of various wild silkworms and different domesticated silkworm strains (*Bombyx mori*).
- The comparative genomics mining of interesting genes and functional elements.
- The identification and functional characterization of genes related to silkworm domestication, silk production, development, and adaptation.
- Epigenetic regulation and non-coding RNAs.
- Gene editing and transgenic applications.
- Genetic diversity, conservation, and breeding strategies.
- Host-microbe interactions and immunity-related genomic studies.

Guest Editor

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Message from the Editor-in-Chief

The International Journal of Molecular Sciences (*IJMS*, ISSN 1422-0067) 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, molecular biophysics, molecular medicine, and all aspects of molecular research in chemistry. *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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