

## Special Issue

# Insect Genome and Transcriptome Data

### Message from the Guest Editors

Dear Colleagues, This Special Issue considers manuscripts related to insect genome data (including related methods, such as metagenome or RADseq data) and transcriptome data through next-generation sequencing (NGS) for publication. To ensure data availability, raw sequence data must be registered in the international nucleotide sequence databases, namely, NCBI, DDBJ, and ENA, and submitted Supplementary Information data must be assigned a DOI using a data repository (e.g., figshare). This SI also considers manuscripts of metadata analysis. As mentioned above, an enormous amount of insect genome and transcriptome data are in public databases. Thus, novel approaches using these data can provide new knowledge contributing to molecular biology and evolutionary biology.

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### Guest Editors

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Dr. Yudai Masuoka

Dr. Akiya Jouraku

Prof. Dr. Hidemasa Bono

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### Deadline for manuscript submissions

closed (31 January 2023)



## Insects

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**Impact Factor 2.9**  
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## About the Journal

### Message from the Editor-in-Chief

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#### Editor-in-Chief

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indexed within Scopus, SCIE (Web of Science), PubMed,  
PMC, PubAg, and other databases.

##### Journal Rank:

JCR - Q1 (Entomology) / CiteScore - Q1 (Insect Science)

##### Rapid Publication:

manuscripts are peer-reviewed and a first decision is  
provided to authors approximately 18.1 days after  
submission; acceptance to publication is undertaken in 2.9  
days (median values for papers published in this journal in  
the first half of 2025).