



## Insect Genome and Transcriptome Data

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Deadline for manuscript  
submissions:

**closed (31 January 2023)**

### 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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