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## Genome Assembly and Evolutionary Analysis of Brassica Crops

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Deadline for manuscript submissions:

**closed (31 March 2023)**

### **Message from the Guest Editors**

The Brassicaceae is an important plant family with approximately 3,800 species, including commercially and economically important vegetables, fodder, oilseed and ornamental crops. Recent advancements in genomics technologies, such as long-read sequencing, chromosome confirmation capture and optical mapping, have enabled the development of highly contiguous genome assemblies. These assemblies have allowed an unprecedented view of the Brassicaceae genome organisation, providing a platform for studying genome evolution, gene expression patterns and trait adaptation. In addition, the foundational resources from the closely related species and model organism, *Arabidopsis thaliana*, can be exploited to identify and confirm candidate gene function underlying traits of interest, leading to crop improvement.

This Special Issue of *Plants* will highlight studies focused on recent genome assemblies of *Brassica* crops and their close relatives, encompassing comparative genomics, pan-genomes, transposable element diversity and evolution, epigenetics, gene function, global gene expression patterns, organelle genome assembly and evolution, evolutionary genomics, and polyploidization.



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# Special Issue



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## Editor-in-Chief

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## Message from the Editor-in-Chief

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