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Advances in Brassica Crops Genomics and Breeding

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Message from the Guest Editors

With the fast progress in sequencing technologies, a number of genomes of Brassica crops species have been sequenced and high-quality chromosome scale assemblies were obtained. Moreover, the large-scale resequencing data of germplasm resources have been made available in *B. rapa*, *B. oleracea*, and *B. napus*, which allows GWAS and domestication analysis in these important crops. These breakthroughs accelerated the investigation into the genomics of the complex Brassica genomes, the evolution of different Brassica species, functional revealing of important genes, and the molecular marker-assisted breeding of Brassica crops. The purpose of this Special Issue is to present the recent advances in genomics and breeding in Brassica crops.

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Message from the Editor-in-Chief

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