

Special Issue

Recent Advances in Horticultural Plant Genomics

Message from the Guest Editors

With the development of sequencing technologies and the reduction of sequencing costs, more and more plant genomes have been sequenced and many results have been obtained, especially with the advent and applications of second- and third-generation sequencing technologies and Hi-C technologies, which have made sequencing a reality in many complex plant genomes. Genomic research tools are widely used in horticultural plants such as fruit trees, vegetables, flowers, tea plants and Chinese herbs for molecular breeding and analysis of growth and development patterns, providing a new perspective on horticultural plant research, which would assist greater understanding of the evolutionary histories of plant species and provide genomic resources for molecular studies on the economically important traits of horticultural plants.

Guest Editors

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Message from the Editor-in-Chief

Plants is an open access journal which provides an advanced forum for research findings in areas related to plant function, its physiology, biology, taxonomy, stresses, and its interactions with other organisms. It publishes original research articles, reviews, reports, conference proceedings (peer reviewed full articles) and communications. In original research papers, it is important that full experimental details are provided. We also encourage timely reviews and commentaries on topics of interest to the plant research community.

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