

Special Issue

Genomewide SNP Identification and Use for Resolving Population Structure and GWAS of Traits of Interest—3rd Edition

Message from the Guest Editors

This Special Issue will publish high-quality research papers, as well as review articles addressing recent studies involving SNP mining, resolving population structures, haplotype networking, and GWAS as emerging tools for plant breeding and ecology research. Original, high-quality contributions that have not yet been published or that are not currently under review by other journals or peer-reviewed conferences are desired.

- SNP mining and InDELs
- population structure analysis
- GWAS
- high-density QTL mapping
- haplotype networking
- genomics of wild crop relatives

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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