Special Issue

Post-Genome Era of Brassica Vegetables

Message from the Guest Editors

The Brassica genus includes many popular vegetables, such as Chinese cabbage, pak choi, cai xin, cabbage, kale, cauliflower, broccoli, and mustard. These vegetables comprise an important part of our daily food and nutrition intake, and contribute to the agricultural economy. For their importance as vegetables and their close relationship to the model plant species Arabidopsis thaliana, the Brassica genus was among the first sets of vegetables to have their genomes sequenced. This genome information has promoted the basic research of *Brassica*, including species evolution, crop domestication, gene mapping, and molecular function dissection, as well as the application of markerbased breeding, with the help of genome and gene function information from the closely related species Arabidopsis thaliana. This Special Issue of Plants will highlight studies on genome evolution, trait dissection, gene function, and population diversity in the postgenome era of *Brassica* vegetables, which use the Brassica genome as an important research resource. We welcome various article types, including original research, reviews, or commentary articles.

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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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