Special Issue

Genetics and Genomics of Sweet Potato

Message from the Guest Editors

Providing ample food for the ever-growing population is a major challenge of our time, especially in rapidly changing climate conditions. Sweet potato (Ipomoea batatas [L.] Lam., Convolvulaceae), among the most widely cultivated staple crops worldwide, is a valuable source of human food, animal feed and industrial raw material. Sweet potato exceeds most other staple foods in vitamins A and C, \(\mathbb{\mathbb{K}}\)-carotene, anthocyanins, calcium and dietary fiber. Consequently, sweet potato could be employed as an excellent source of natural healthpromoting compounds. In recent years, sweet potato has been in the spotlight of agricultural biotechnology and has been considered as a biological model for storage root formation. Although studies on genetics and genomics have contributed to progress on sweet potato research during the past decade, there is still a gap in knowledge compared with other crops. This Special Issue aims to integrate recent research in sweet potato biology by expanding our knowledge in various fields, such as genetics, molecular biology, functional genomics, biotic and abiotic stress responses, and omics studies.

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Message from the Editor-in-Chief

Genes is central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fast-moving field. There is a need for good quality, open access journals in this area, and the Genes team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised. Why not consider Genes for your next genetics paper?

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