



Genetics and Genomics of Tomato and Solanaceae - Series II

Guest Editor:

Dr. Dilip R. Panthee

Department of Horticultural
Science, North Carolina State
University, Raleigh, NC 27608,
USA

Deadline for manuscript
submissions:

closed (1 November 2020)

Message from the Guest Editor

Dear colleagues,

With the availability of genome sequencing technology, there is an enormous amount of genomic resources available of several genotypes now. Different research groups around the world have generated a massive amount of data. The current challenge is how to extract the information from this data and use it properly in plant breeding. If this information can be utilized properly, it opens the floodgate of genetic information, making the life of plant breeders easy. They can target a gene of interest easily by using marker-assisted selection (MAS). Considering this challenge, this Special Issue will focus on the bioinformatics aspect of Solanaceae. We encourage articles focusing information on the genome sequence, molecular marker development from the sequence, and its utilization in plant breeding.

Dr. Dilip R. Panthee
Guest Editor





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Editor-in-Chief

Prof. Dr. Leslie A. Weston

Gulbali Centre for Agriculture,
Water and Environment
Research, Charles Sturt
University, Wagga Wagga, NSW
2678, Australia

Message from the Editor-in-Chief

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Agronomy Editorial Office
MDPI, Grosspeteranlage 5
4052 Basel, Switzerland

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