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## Epigenetics and Genome Evolution in Plants

Guest Editor:

**Dr. Xuan Hieu Cao**

Forest Genetics and Forest Tree  
Breeding, Georg-August  
University of Göttingen,  
Busgenweg 2, D-37077 Göttingen,  
Germany

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### Message from the Guest Editor

Accumulating evidence many of those were from plant systems suggests the contribution of epigenetic mechanisms to genome evolution. For example, DNA methylation is vital for the silencing of transposable elements, enabling the colonization of a substantial proportion of repetitive sequences in plant genomes. During evolution, these genomic elements are capable of moving and self-replicating in germlines in waves (i.e., epigenetic assimilation), potentially resulting in genetic diversity and genome plasticity. Furthermore, these epigenetically controlled dynamics may also steer the ecological and evolutionary advantage of hybridization and polyploid events by perhaps modulating the transcriptional expression of nearby genes. This special issue aims to collect a wide body of research studies dealing with epigenomics, population epigenetics, and functional genomics in model and non-model plant species that have significant relevance in genome evolution, adaptation, ecological speciation, and diversification.



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# Special Issue



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### **Prof. Dr. Dilantha Fernando**

Department of Plant Science,  
University of Manitoba, Winnipeg,  
MB R3T 2N2, Canada

## Message from the Editor-in-Chief

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*Plants* Editorial Office  
MDPI, St. Alban-Anlage 66  
4052 Basel, Switzerland

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