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## **Plant Genomics and Epigenomics for Trait Improvement**

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Deadline for manuscript submissions:

closed (30 November 2017)



genes







**Message from the Guest Editors** 

Dear Colleagues,

understanding of plant traits and mechanisms has been enormously improved over the past decade. Recent plant breeding programs have taken into account available genomic information, e.g., for (1) dissecting the genetic architecture of agronomic traits; (2) unlocking the hidden favorable genetic diversity from genetic resources; or (3) discovering, validating and characterizing economically important genes. Evidence has been accumulating that heritable variation of a trait is not solely determined by DNA sequence polymorphism but involves epigenetic processes that impact chromatin structure and gene expression. Especially, with the rapid development of CRISPR-Cas technology, it is believed that the promise of novel and improved crops with greater yield and tolerance to the stresses of climate change and extreme weather is around the corner. The aim of this issue is to publish original research, review articles addressing recent advances on plant genomics and epigenomics as emerging approaches for plant breeding.

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Blvd, SHEL 814, Birmingham, AL <u>G529</u>年 5月快灾间最风ffice MDPI, Grosspeteranlage 5 4052 Basel, Switzerland Genes are central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fastmoving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide 1683 77 34 manuscript handling serious peer review, and rapid publication across others which discipline of genetics. Starting in 2010, the journal is now well

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