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Analysis, Identification and Utilization of Genetic Resources Related to Peanut

Guest Editors:

Dr. Hui Zhang

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Dr. Ming Li Wang

Deadline for manuscript submissions:

closed (25 April 2024)

Message from the Guest Editors

Dear Colleagues,

Peanut is an important oil crop belonging to Leguminosae. In this Special Issue, we intend to publish original research reports and reviews concerning the genomics and genetics of peanut. Topics might include, but are not limited to, the effects of genotype/environment/G by E on important peanut traits, using genome-wide association studies, genomic selection, or breeding lines to identify the QTLs/genes that can be potentially used in breeding through marker-assisted selection. In addition, this Special Issue also includes the detection of genes using bioinformatic analysis or omics tools. We will particularly consider manuscripts with a clear genetic component and significant impact on peanut breeding.

Dr. Hui Zhang Dr. Xinlei Yang Prof. Dr. Charles Y. Chen Dr. Ming Li Wang Guest Editors











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Editor-in-Chief

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Message from the Editor-in-Chief

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