



viruses



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Plant Virus Metagenomics

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Message from the Guest Editor

Dear Colleagues,

With the advances in sequencing technologies and reduced costs involved, metagenomics has become an indispensable tool in various aspects of plant virus research, namely in the identification and characterization of novel viruses, virus diversity and evolution, viral pathogenesis, and virus–host interactions, as well as in RNA silencing, virus detection and diagnosis.

This Special Issue on plant virus metagenomics invites authors to submit original research or review articles that involve the application of metagenomics in any aspect of plant virus research. Researchers are encouraged to submit novel applications of advanced sequencing technologies and to develop bioinformatics tools and databases related to plant virus metagenomics.

Prof. Dr. Renyi Liu
Guest Editor



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Message from the Editor-in-Chief

Viruses (ISSN 1999-4915) is an open access journal which provides an advanced forum for studies of viruses. It publishes reviews, regular research papers, communications, conference reports and short notes. Our aim is to encourage scientists to publish their experimental and theoretical results in as much detail as possible. There is no restriction on the length of the papers. The full experimental details must be provided so that the results can be reproduced. We also encourage the publication of timely reviews and commentaries on topics of interest to the virology community and feature highlights from the virology literature in the 'News and Views' section.

Electronic files or software regarding the full details of the calculation and experimental procedure, if unable to be published in a normal way, can be deposited as supplementary material.

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