



## Identification and Molecular Characterization of Plant Virus

Guest Editor:

**Dr. Igor Koloniuk**

Department of Plant Virology,  
Institute of Plant Molecular  
Biology, Biology Centre of the  
Czech Academy of Sciences, v.v.i.,  
370 05 České Budějovice, Czech  
Republic

Deadline for manuscript  
submissions:

**closed (30 January 2022)**

### Message from the Guest Editor

Dear Colleagues,

Ongoing technological advances in high-throughput sequencing and its wide adoption have led us to discoveries of numerous novel viral species. While a number of findings are related to established viral families, some are pieces of the puzzle that do not fall into place. That and the metagenomic nature of many studies raise a question of host affiliations and correctness of host assignment.

Brief reports and research articles describing novel plant viruses and mixed infections, including infections of multiple strains, will be considered. Publications on the metagenomic analysis of plant samples, including viruses of plants and the associated organisms, are especially welcome in the Special Issue.

Dr. Igor Koloniuk  
*Guest Editor*





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## Editor-in-Chief

### 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  
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