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Mastreviruses in the African World: Harbouring Both Monocot and Dicot Species

<u>Avinash Marwal. Rakesh Kumar Verma, Megha Mishra, Rajesh Kumar & R. K. Gaur</u> Chapter | <u>First Online: 13 July 2019</u>

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Abstract

Mastreviruses are the main causal viral agent of a variety of plant diseases in African continent posing a serious threat to economically important plant species, residing/harbouring chiefly in the uncultivated ones. Mastreviruses are known to have numerous diverse carrier vectors infecting either monocotyledons or dicotyledons plants. More than 1950 Mastreviruses sequences are publically available in nucleotide database of NCBI, majority (~850) of them are reported from Africa alone. All the known Mastreviruses encompass a monopartite genomic nature (ranges from 2.5 to 2.7 kb) encoding four genes. Reports and evidences suggest a strong intra- or inter-specific recombination among the identified Mastreviruses, but the extent to which it creates diversity is still a challenging task to understand and get a clear insight. Such diversity is also supported by gene acquisition and mutations (especially point mutations

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