## New Disease Reports

## Association of *Cotton leaf curl Multan virus* and its associated betasatellite with leaf curl disease of *Hibiscus rosa-sinensis* in India

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Hibiscus (*Hibiscus rosa-sinensis*) is a perennial ornamental plant grown for its attractive flowers. In September 2011, symptoms of severe leaf curling, vein thickening and enation were observed on 31% (13/42) hibiscus plants (Fig. 1) grown in a garden at Lucknow, India. The symptoms and whitefly infestation suggested a begomovirus infection.

Leaf samples from 13 diseased and four healthy hibiscus plants were collected. Total DNA was isolated and subjected to PCR using degenerate primers begomovirus (CP-F: 5'-GCATCTGCAGGCCCACATYGTCTTYCCNGT-3' and CP-R: 5'-AATACTGCAGGGCTTYCTRTACATRGG-3'). All samples of symptomatic leaves yielded a ~1.2 kb PCR product indicative of a begomovirus. Full-length genomes were amplified by the rolling circle amplification method (~2.7 kb) from two samples, and cloned and sequenced (GenBank Accession Nos. JN880418 and JN807763). Nucleotide sequence analysis revealed 99% identity between the two isolates, and 90-93% identity and close phylogenetic relationships with isolates of Cotton leaf curl Multan virus (CLCuMV) (Fig. 2). Hence, two new variants of CLCuMV were identified. Association with a betasatellite tested by PCR with specific was primers (Beta 01: 5'-AGCCTTAGCTACGCCGGAGC-3' Beta 02: and 5'-GCTGCGTAGCGTAGAGGTTT-3'). Again, all 13 samples of symptomatic leaves produced the expected product of ~1.3 kb. One

21

0

20

23

0

amplicon was cloned and sequenced (KT447040). The sequence had 98% nucleotide identity and close phylogenetic relationships with *Cotton leaf curl Multan betasatellite* (CLCuMB) isolates (Fig. 3), and was concluded to be an isolate of CLCuMB. The disease incidence was confirmed during a survey of the same garden in September 2015. Twenty three of the 42 hibiscus plants investigated by dot blot analysis using a CLCuMV-specific probe (derived from clone JN880418) were found to be infected (Fig. 4).

The natural occurrence of CLCuMV and CLCuMB has been reported on hibiscus from China (Mao *et al.*, 2008), whereas *Cotton leaf curl Burewala virus* with CLCuMB on hibiscus has been reported from Pakistan (Akhtar *et al.*, 2014). However, this is the first report of CLCuMV and CLCuMB on hibiscus from India.

## References

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Figure 4

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