



First report of *Canna yellow mottle virus* on *Canna* from India

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Canna (*Canna indica*) is a popular landscape plant known for its attractive foliage and flowers and is grown in tropical and subtropical countries including India. Various cultivars of *canna* are being maintained as germplasm at the Botanic Garden, NBRI, Lucknow. During a survey in September 2012, necrotic striping, severe veinal chlorosis, yellow mosaic and veinal streaking symptoms were observed on the leaves of different *canna* cultivars (Fig. 1a-d), with the disease incidence between 17 and 35% at NBRI. Subsequently, similar symptoms were found on *canna* in other gardens in Lucknow. Symptoms observed were similar to those described previously for *canna* yellow mottle disease and associated with *Canna yellow mottle virus* (CaYMV) (Pappu *et al.*, 2008; Marino *et al.*, 2008).

For virus detection, total genomic DNA was isolated from leaf samples of 13 plants with symptoms and one without, and subjected to PCR using CaYMV specific primers (CaYMV-3 and CaYMV-4; Momol *et al.*, 2004). PCR resulted in an amplicon of the expected size (~550 bp) in all the 13 samples with symptoms, but not in the sample from the symptomless plant, indicating the presence of CaYMV in the diseased plants. Subsequently the 13 amplicons obtained from 10 different cultivars of *Canna indica* and *C. flaccida* were cloned, sequenced and deposited in GenBank (Table 1). These accessions, in BLASTn analysis, were shown to share 93-99% nucleotide sequence identity with the available CaYMV isolate sequences from around the world. Further, phylogenetic analysis of sequences under study by MEGA5.1 (Tamura *et al.*, 2011) using a neighbour-joining method clearly clustered them together with CaYMV (Fig. 2). Based on sequence comparison and phylogenetic analysis, the virus isolates in this study were identified as CaYMV. To the best of our knowledge, this is the first report

of natural occurrence of CaYMV on *canna* from India. CaYMV may be considered as posing a phytosanitary risk for further spread in *canna* since it is being propagated through suckers or rhizomes.

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Figure 1

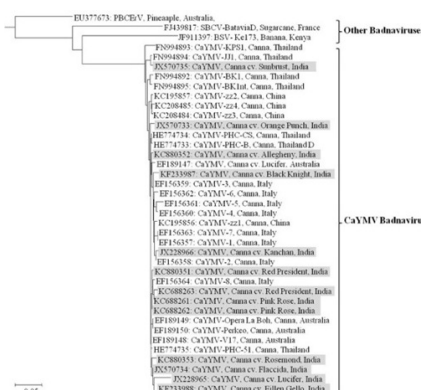


Figure 2

Table 1. Details of *Canna yellow mottle virus* sequences isolated from different cultivars of *Canna* spp., submitted to GenBank database.

| Isolate number | Accession number | Species/cultivars of <i>canna</i> | Sequence length (bp) |
|----------------|------------------|-------------------------------------|----------------------|
| AK-1 | JX228965 | <i>Canna indica</i> cv. Lucifer | 565 |
| AK-2 | JX228966 | <i>C. indica</i> cv. Kanchan | 565 |
| AK-3 | JX570733 | <i>C. indica</i> cv. Orange Punch | 565 |
| AK-4 | JX570734 | <i>C. flaccida</i> | 565 |
| AK-5 | JX570735 | <i>C. indica</i> cv. Sunburst | 565 |
| AK-6 | KC688261 | <i>C. indica</i> cv. Pink Rose | 441 |
| AK-7 | KC688262 | <i>C. indica</i> cv. Pink Rose | 529 |
| AK-8 | KC688263 | <i>C. indica</i> cv. Red President | 472 |
| AK-9 | KC880351 | <i>C. indica</i> cv. Red President | 565 |
| AK-10 | KC880352 | <i>C. indica</i> cv. Allegheny | 565 |
| AK-11 | KC880353 | <i>C. indica</i> cv. Rosemond Coles | 565 |
| AK-12 | KF233987 | <i>C. indica</i> cv. Black Knight | 525 |
| AK-13 | KF233988 | <i>C. indica</i> cv. Eileen Gallo | 486 |

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