Supplementary Material

High-throughput sequencing facilitates the characterisation of the “forgotten” plant virus: A case of the novel henbane mosaic virus infecting tomato

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# Supplementary Figures and Tables

## Supplementary Table

Supplementary Table 1: Detailed information about HMV isolates described in the study.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Isolate name in manuscript | Isolate name | Original host plant; laboratory host plant in which the virus was maintained | Virus collection (organization) | Date (DD/MM/YYYY), place of sampling and collector (if known) | Detected virus species | NCBI GenBank accession number for HMV | NCBI SRA accession number (sRNA/rRNA depleted totRNA) |
| HMV-SI/L | HMV-SI/L | *Solanum lycopersicum*; maintained in *Solanum lycopersicum* | National Institute of Biology | 24/06/2015 Ankaran, Slovenia, collected by Patricija Pirnat | *Henbane mosaic virus, Potato virus M, Southern tomato virus* | MH779472 | SRR7734364/ SRR7734393 |
| HMV-R | HMV-R (Rothamsted) | unknown; maintained in *Nicotiana tabacum* | Institute for Sustainable Plant Protection | Originating from Rothamsted, 01/01/1963 - was collected by Dr. Osvaldo Lovisolo | *Henbane mosaic virus, Potato aucuba mosaic virus* | MH779474 | SRR7734366/ NA |
| HMV-146 | HMV-146 | *Datura inermis*; maintained in *Datura stramonium* | Institute for Sustainable Plant Protection | 19/8/1964, Torino (Botanic Garden), collected by Dr Osvaldo Lovisolo | *Henbane mosaic virus, Potato aucuba mosaic virus* | MH779473 | SRR7734365/ NA |
| HMV-PV-76 | California | *Hyoscamus niger*, Califormia; maintained in *N. tabacum* cv. Samsun | American Type Culture Collection (ATCC) | See ATCC:https://www.lgcstandards-atcc.org/products/all/PV-76 | *Henbane mosaic virus* | MH779475 | SRR7734367 /NA |
| HMV-PV-79 | Watson's Isolate A | *Hyoscamus niger*, England; maintained in *N. tabacum* cv. Samsun | American Type Culture Collection (ATCC) | See ATCC:https://www.lgcstandards-atcc.org/products/all/PV-79 | *Henbane mosaic virus, Potato virus Y* | MH779476 | SRR7734368/ NA |

## NA – not applicable

Supplementary Table 2: List of primer pairs used for PCR amplification.

|  |  |  |  |
| --- | --- | --- | --- |
| Pair number | Primer name | Primer sequence 5ˈ-3ˈ | Primer position at HMV-SI/L sequences consensus |
| 1 | P1start-FP1-R | CTTACTCTTCCTCCCCCAATTGTTCTGTTGTTCCTCC | 331-3481268-1286 |
| 2 | P1end-FHCpro-R | GCGGTTTTATCTTTCCAGTTGATTCTTTGTGGGTGTCTT | 1469-14872536-2555 |
| 3 | P3-FP3-R | TTCTCACCAAGCATTCTCCATCTTCTCATCACCATCA | 3115-31323867-3885 |
| 4 | Cl-FCl-R | AGAGTTGAGTAAGAGGGGGTAAAACACAGAAATGCCGCC | 4488-45065349-5368 |
| 5 | Clend-FNIaPro-R | ACATCAAGTAGAGAGCGGGGTTCGGAGTGTTAGTGT | 6042-60597012-7029 |
| 6 | HMV-NIb-FHMV-CP-R | GTCAAGAAGTTCAAAGGGTACACCACACCATCAATC | 8479-84969473-9490 |
| 7 | HMV-UNI-FHMV-UNI-R | TTAGCCCGATATGCTTTCCTATCTTCCACTTCAGGT | 9715-973210043-10060 |

**Supplementary Table 3:** Detailed description of annotated polyproteins for HMV isolates: HMV-SI/L, HMV-146, HMV-R, HMV-PV-76 and HMV-PV-79. Information about virus genome and its polyprotein length (nt) are given for each isolate. The corresponding number of amino acid residues, molecular weight (kilo dalton) and cleavage position for each protein are listed in separate lines. The PIPO position is indicated at the bottom of the table.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
|  | HMV-SI/L | HMV-146 | HMV-R | HMV-PV-76 | HMV-PV-79 |
| Virus length (nt)/ Virus polyprotein length (nt) | 10127/9747 | 10125/9747 | 10140/9750 | 10140/9750 | 10140/9750 |
| POLYPROTEIN | Residues No. | 3248368.04 | 3248367.65 | 3249367.19 | 3249367.26 | 3249367.27 |
| Mw (kDa) |
| P1 | Residues No. | 47654.5MQYY476-S | 476 54.3 MRYY476-S | 47753.82MRYY477-S | 47753.78MRYY477-S | 47753.78MRYY477-S |
| Mw (kDa) |
| Cleavage position |
| HC-Pro | Residues No. | 45751.54YQVG933-G | 45751.6 YQVG933-G | 45751.42YQVG934-G | 45751.4YQVG934-G | 45751.4YQVG934-G |
| Mw (kDa) |
| Cleavage position |
| P3 | Residues No. | 34840.14VKHQ1281-G | 34840.05 VKHQ1281-A | 34840.14VKHQ1282-A | 34840.15 VKHQ1282-A | 34840.15VKHQ1282-A |
| Mw (kDa) |
| Cleavage position |
| 6K1 | Residues No. | 546.09VTHQ1335-S | 546.14 VTHQ1335-S | 546.12VAHQ1336-S | 546.12 VAHQ1336-S | 546.12VAHQ1336-S |
| Mw (kDa) |
| Cleavage position |
| CI | Residues No. | 64571.87VMHQ1980-S | 64571.76 VMHQ1980-S | 64571.83VMHQ1981-S | 64571.89 VMHQ1981-S | 64571.92VMHQ1981-S |
| Mw (kDa) |
| Cleavage position |
| 6K2 | Residues No. | 535.91VAHQ2033-G | 535.86 VAHQ2033-G | 535.9VAHQ2034-G | 535.9 VAHQ2034-G | 535.9VAHQ2034-G |
| Mw (kDa) |
| Cleavage position |
| VPg | Residues No. | 16318.49PEQE2196-G | 16318.49 PEYE2196-G | 16318.57PEYE2197-G | 16318.57 PEYE2197-G | 16315.57PEYE2197-G |
| Mw (kDa) |
| Cleavage position |
| NIa-Pro | Residues No. | 26730.36VVEQ2463-A | 26730.28 VTEQ2463-A | 26730.29VTEQ2464-A | 26730.29 VTEQ2464-A | 26730.29VTEQ2464-A |
| Mw (kDa) |
| Cleavage position |
| NIb | Residues No. | 51659.06VYHQ2979-G | 51659.04VYHQ2979-A | 51658.93VYHQ2980-A | 51658.93VYHQ2980-V | 51658.93VYHQ2980-A |
| Mw (kDa) |
| Cleavage position |
| CP | Residues No. | 26930.24 | 26930.31  | 26930.33 | 26930.39 | 26930.37 |
| Mw (kDa) |

PIPO: at polyprotein sequence position: 3267 nt – 3473 nt for HMV-R, HMV-PV-76 and HMV-PV-79 isolates and 3264 nt – 3470 nt for: HMV-SI/L and HMV-146 isolates.

## Supplementary Figure

**Supplementary figure 1**: Diagnostic screening testing. A) Mechanically inoculated test plants with developed symptoms (yellow arrows indicate symptomatic plants of *Solanum lycopersicum* cv. Moneymaker, *Nicotiana rustica, Nicotiana tabacum* cv. White Burley, *Nicotiana clevelandii*). B) Observed viral particles (using TEM) in mechanically inoculated test plants.

**Supplementary figure 2:** Disease symptoms on tomato plants (*Solanum lycopersicum* cv. Moneymaker) three weeks after inoculation with: HMV-PV-76 (left), MOCK (middle) and HMV-SI/L (right). Tomato plants were mechanically inoculated as explained in section 2.2.