

## Supplementary information

**Table S1.** Amino acid changes through each ORF from the complete genomes of MWMV-ZuM10 and WMV-MeWM7 isolates. Each aa change is based on their non-synonymous mutations, which were analysed by the sequence comparison with the corresponding Sq10 1.1 (MWMV type) and Vera (WMV) isolates.

<b>Virus/ORF</b>	<b>MWMV-ZuM10</b>	<b>WMV-MeWM7</b>
<b>P1</b>	K197E; T262I; G348S; S363L	K34E; M153V; K187N; T274A
<b>HC-Pro</b>	-	F8I; S10A; K91E; G96R; P110S; L314F; L348F; L394F
<b>P3</b>	P857S; Y872C	I2V; N30D; N206S; Y215H; I286F; G307E; K335M
<b>6K1</b>	-	S43G
<b>CI</b>	-	S249N; N291T
<b>6K2</b>	-	-
<b>Nia-VPg</b>	N2366I	E109G; S153F
<b>VPg-Pro</b>	-	M54I
<b>Nib</b>	V2423A	I42L; V434I; R482K; N505S
<b>CP</b>	-	P40Q; A41I; D21N; A52N; V58E; I243M