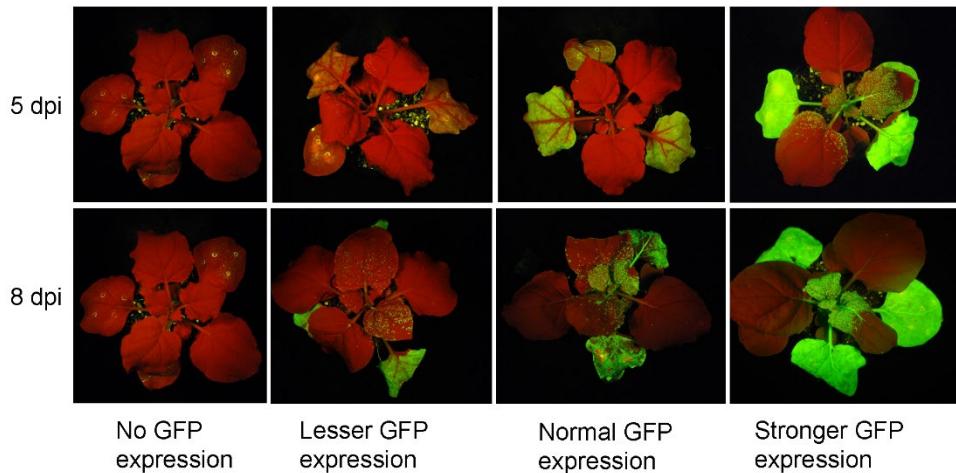
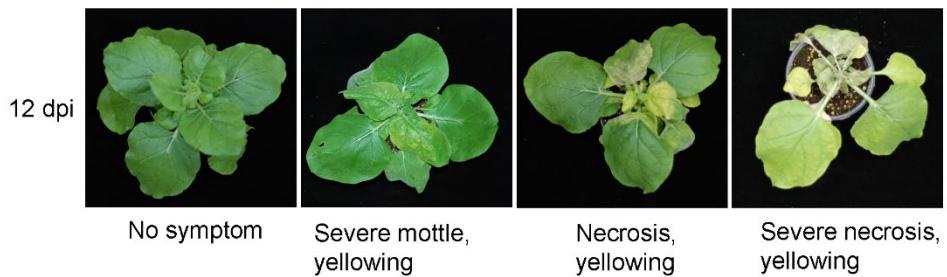
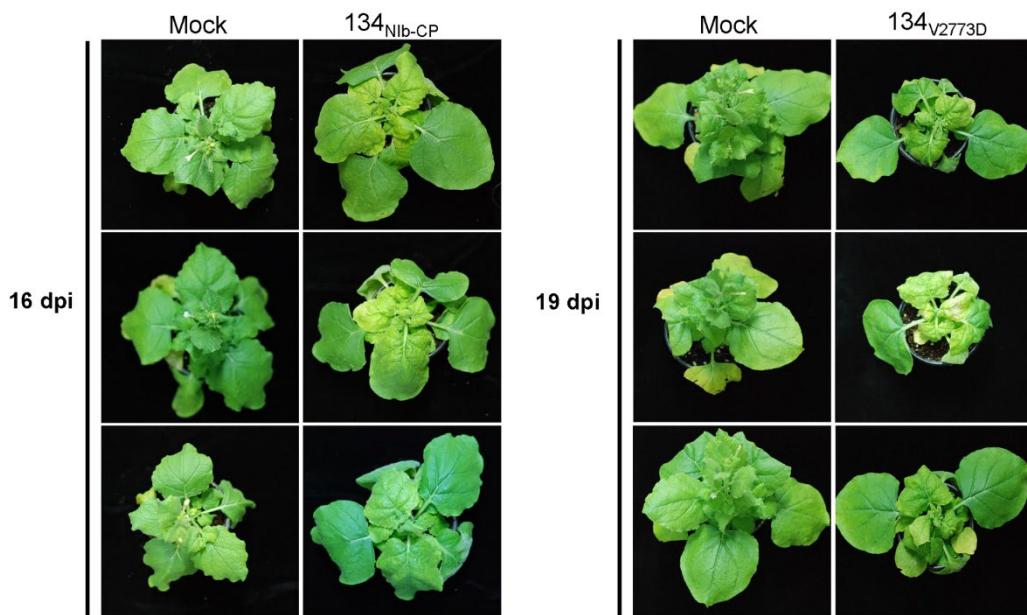


## Supplementary Material

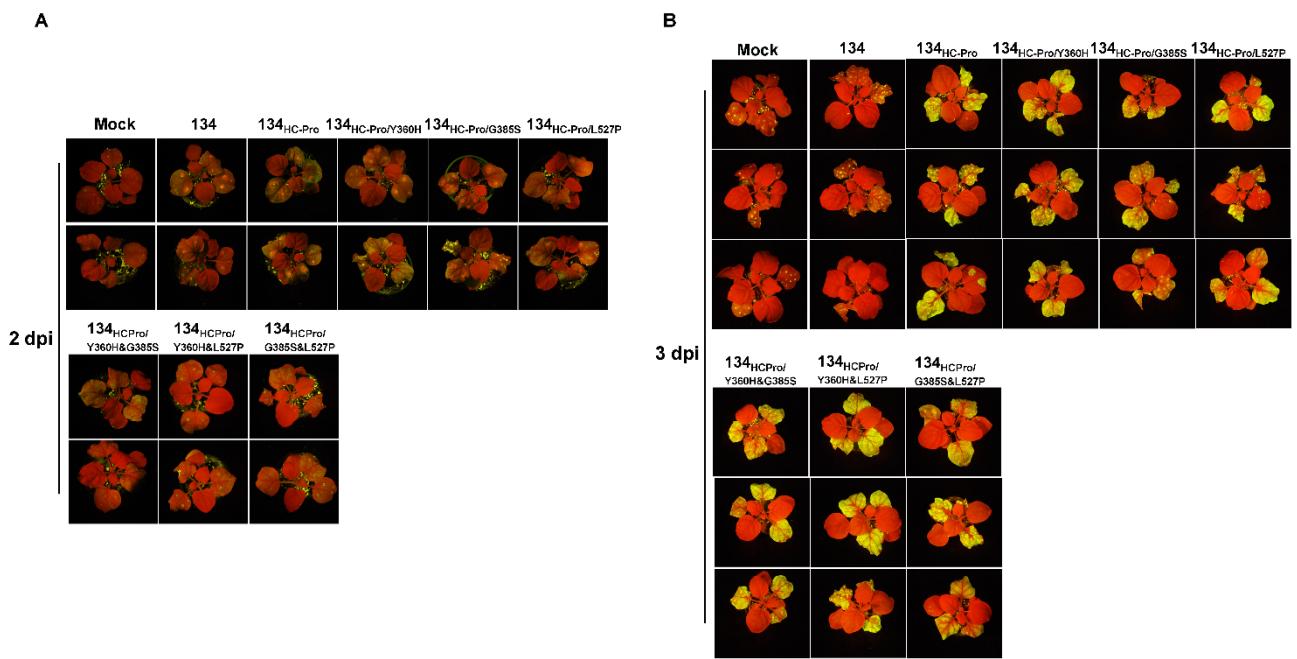
### 1.1 Supplementary Figures

**A****B**

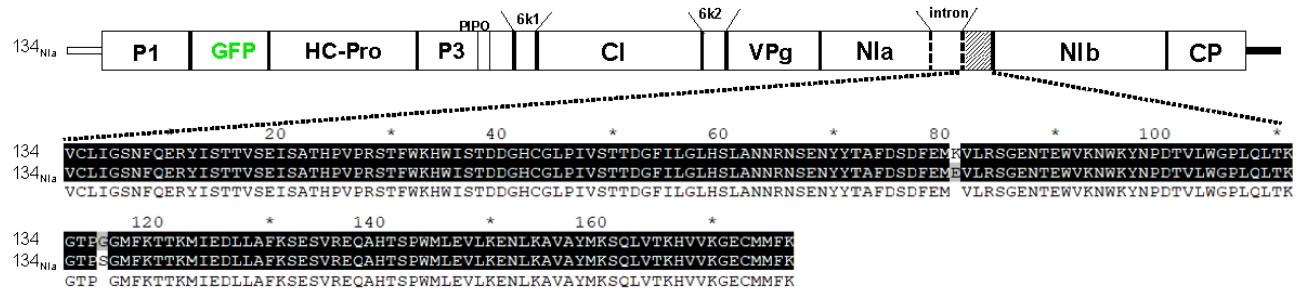
**Supplementary Figure 1.** (A) Scale card of GFP fluorescence expression. (B) Scale card of symptom expression.



**Supplementary Figure 2.** Symptom development of 134<sub>Nlb-CP</sub> and 134<sub>V2773D</sub> were monitored extra few days to confirm no necrotic symptoms. (A) Symptoms of mock and 134<sub>Nlb-CP</sub> at 16 dpi. (B) Symptoms of mock and 134<sub>V2773D</sub> at 19 dpi.



**Supplementary Figure 3.** Observation of GFP expression of 134 and 134<sub>HC-Pro</sub>-derived mutants after agroinfiltration on *N. benthamiana*. GFP fluorescence representing virus infection on plants at 2 dpi (A) and 3 dpi (B).



**Supplementary Figure 4.** Amino acid alignment sequence of the C-terminal NIa of PepMoV isolates 134 and 134<sub>NIa</sub>.

## 1.2 Supplementary Table

Primer name	Sequence 5'-3'	Restriction enzymes	Purpose
pCAMBIA 6601 Fw	GCTGGCTGGTGGCAGGATA		
P1 <i>Kpn</i> I Rv	GATGAACGTTAACGGTACCCGAATACTG TTCCATATGAAGTACAGTTGC	<i>Bsp</i> 120I <i>Acc</i> 65I	Construction of 134 <sub>P1</sub>
PepMoV 948 Fw	GTTCGCGAAAATCGGATGG		
PepMoV 2511 Rv	ACATGACACTTGGCCTGTAGATCCC	<i>Acc</i> 65I <i>Pac</i> I	Construction of 134 <sub>HC-Pro</sub>
PepMoV 2222 Fw	TGACGCAGAGCTGCCTCGTAT		
PepMoV 4727 Rv	TGCGGTGCCCTCTGTATGCG	<i>Pac</i> I <i>Sac</i> I	Construction of 134 <sub>P3-CI</sub>
PepMoV 3940 Fw	ACGGCAGGGTTTGCTGATA		
PepMoV NIa-Intron2 ST-LS1 Fw	AACAGACAGGTTGTTCTGCTTCTACCT TTGAT	<i>Sac</i> I <i>Mss</i> I	Construction of 134 <sub>CI-NIa</sub>
Intron2 ST-LS1- PepMoV NIa Rv	GAGACACACCCTAACATCACCATTTT GGTCA		Construction of 134 <sub>NIa</sub>
PepMoV 7262 Rv	TGCGTCGCAATCGACTACTCCT		
PepMoV 6988 Fw	GGGAGCAAGCACACACATCACCT	<i>Mss</i> I <i>Mlu</i> I	Construction of 134 <sub>NIB-CP</sub>
pSNU1 241 Rv	TCGCAAGACCGGCAACAGGA		
mutant4 I2374V Fw	TATCATTAGAGTTGTTGTATACATGCAGA	<i>Mss</i> I <i>Mlu</i> I	Construction of 134 <sub>I2374V</sub>
mutant4 I2374V Rv	TCTGCATGTATACAACAACTCTAATGATA		
mutant4 V2773D Fw	GAGCTTCAGGACTACCTCAGA	<i>Mss</i> I <i>Mlu</i> I	Construction of 134 <sub>V2773D</sub>
mutant4 V2773D Rv	TCTGAGGTAGTCCTGAAGCTC		
mutant4 T2789A Fw	TTGAATGTGGTGCATATGAAGTTCA	<i>Mss</i> I <i>Mlu</i> I	Construction of 134 <sub>T2789A</sub>
mutant 4 T2789A Rv	TGAACTTCATATGCACCACATTCAA		
mutant4 T2805A Fw	GATACATTGGATGCTGGAGAGGAGA	<i>Mss</i> I <i>Mlu</i> I	Construction of 134 <sub>T2805A</sub>
mutant4 T2805A Rv	TCTCCTCTCCAGCATCCAATGTATC		
PepMoV diag Fw	ATGAGCAGCTAACAGATCAGATACATTG		PepMoV detection primer
PepMoV diag Rv	CATATTCCTGACCCCAAGCA		PepMoV detection primer
PepMoV 5836 Fw	GCACAACCGTTGGCATGGGC		PepMoV quantification in RT-qPCR
PepMoV 5950 Rv	TCCATTGTGCACCTGTGAGTGG		PepMoV quantification in RT-qPCR
Nb Actin real time Fw	CCAGGTATTGCTGATAGAATGAG		Internal control for RT-qPCR

Nb Actin real time Rv	CTGAGGGAAAGCCAAGATAGAG		Internal control for RT-qPCR
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**Supplementary Table 1.** Primers and restriction enzymes used in this study.