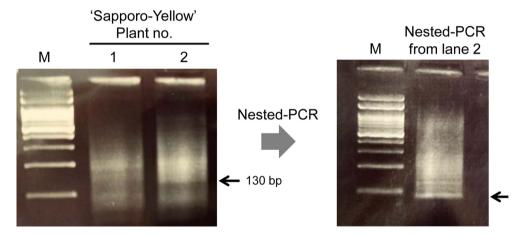
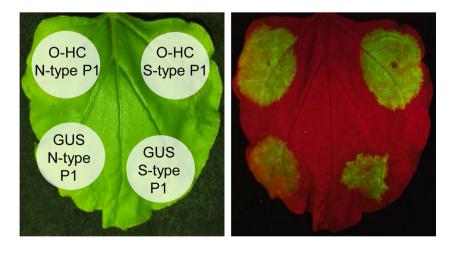


**Supplementary Fig. 3.** Detection of leek yellow stripe virus (LYSV) in onion (cv. Fresh-Red) mechanically inoculated with the sap from China garlic infected with LYSV-S. At 21 days postinoculation, the partial sequence of LYSV was reverse transcription polymerase chain reaction-amplified from the upper leaf using primer pair LYSV-5-130/LYSV-3-130. The detected bands (arrow) were excised and sequenced to confirm the S-type P1 sequence. Onion yellow dwarf virus (OYDV) was not detected in this onion at 21 days postinoculation. M, 50 bp DNA ladder.



**Supplementary Fig. 4.** Detection of leek yellow stripe virus (LYSV) in onion seedlings purchased from the market. LYSV was detected as a faint band (arrow, left gel) by the first-round reverse transcription polymerase chain reaction using primer pair LYSV5-130/LYSV3-130, but a distinct band was detected at the expected position (arrow, right gel) by the nested-polymerase chain reaction using primer pair LYSV5-nest200/LYSV3-130. The band was excised and sequenced, revealing that LYSV actually had S-type P1. M, 100 bp DNA ladder.



**Supplementary Fig. 5.** RNA silencing suppressor (RSS) activity of LYSV P1 proteins in the presence of LYSV-S-derived HC-Pro (O-HC). *Nicotiana benthamiana* leaves were coinfiltrated with *Agrobacterium* suspensions carrying GFP, O-HC and P1 at the same ratio  $(OD_{600} = 0.6)$ . GUS was used as a negative control. Image with GFP fluorescence was taken at 5 days post agroinfiltration.