

Supporting information:

Virus variants with differences in the P1 protein coexist in a Plum pox virus population and display particular host-dependent pathogenicity features

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Figure S1. Multiple amino acid alignment of the P1 proteins of 97 isolates belonging to seven Plum pox virus strains recovered from public databases. The amino acid conservation at positions W29 and V139E is highlighted.

Figure S2. Alignment of small RNAs to the reference PS-MCL genome. The green bars represent the unique small RNAs (length between 20 and 25 bases) aligned to the two strands of the PS-MCL genome. Only perfect matching was allowed.

