Figure S11. Myzus persicae flavivirus (MpFV) genomic (g)RNA secondary structures upstream and downstream of a predicted 5'-end of subgenomic (sg)RNA. Secondary structures of the gRNA sequences surrounding the 5'-end of sgRNA determined using Turner model 2004 at 30°C at the Webserver http://rna.tbi.univie.ac.at/cgi-bin/RNAWebSuite/RNAfold.cgi are presented as images exported from the Webserver. Positions of 5'- and 3'-termini of the folded gRNA sequences are indicated and the sgRNA 5'-terminal nucleotide is circled. The two most stable hairpins located downstream of the sgRNA 5'-end are indicated with pink ovals. Color code indicates base pair probabilities ranging from 0 (blue) to 1 (red).

