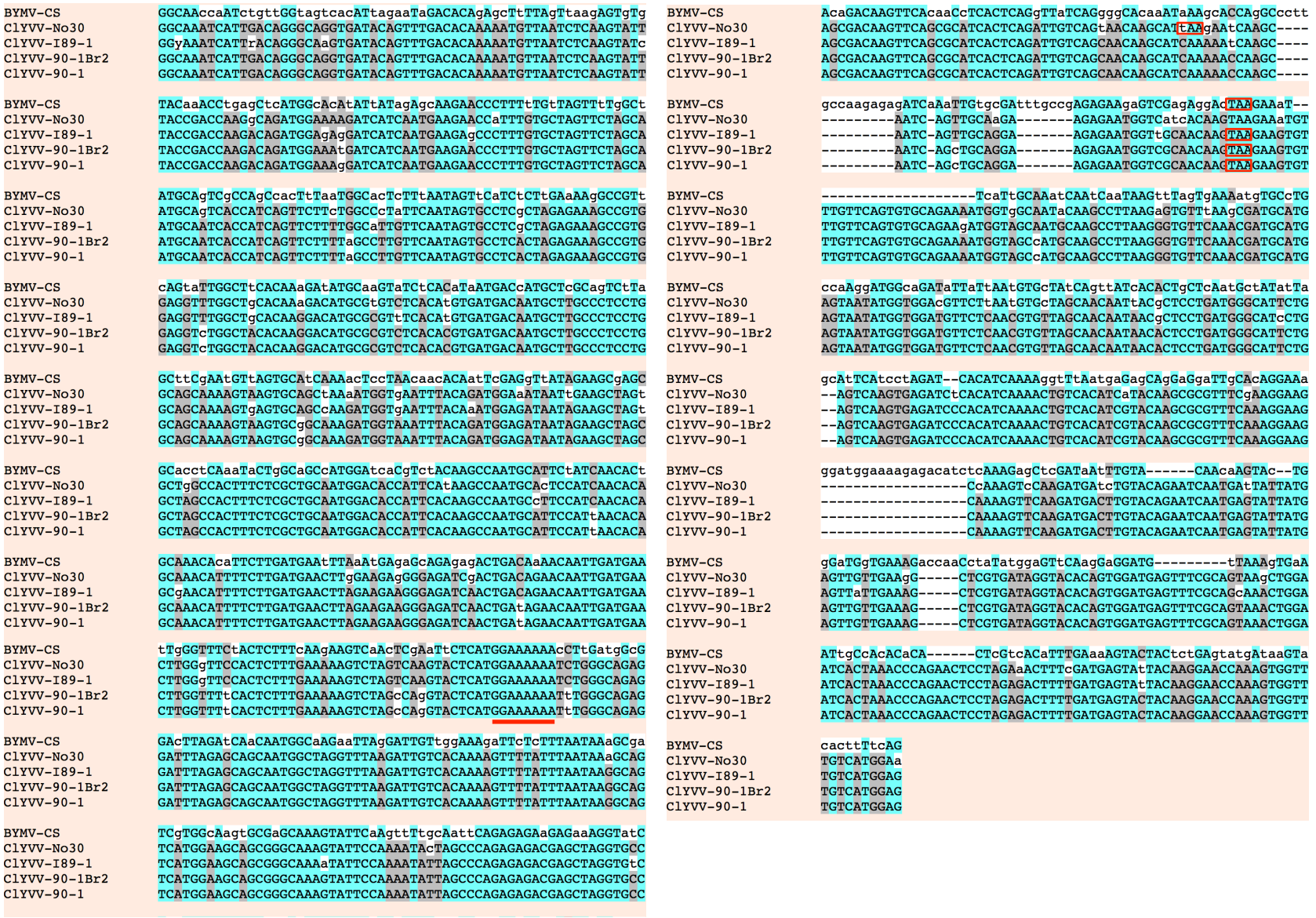

FIG S1 Multiple alignment of nucleotide sequences of ClYVV and BYMV *P3* genes. Alignment was performed using the program MUSCLE (3.8) (http://www.ebi.ac.uk/Tools/msa/muscle/)(28). Accession numbers: BYMV-CS (AB373203), ClYVV-No.30 (AB011819), ClYVV-I89-1 (LC096082) and ClYVV-90-1 Br2 (AB732962). Red underline indicates the G_2A_6 motif. Red boxes indicate the stop codon of the PIPO reading frame in each sequence.