Figure S3. RT-PCR analysis of TEVΔNIb progeny at 14 dpi in 35S::NIb (lanes 1 to 3) and wild-type (lane 4) *N. benthamiana* plants inoculated with several virus combinations as indicated. Amplification products were separated by electrophoresis in an agarose gel stained with ethidium bromide. Lane 1, non-inoculated plant; lanes 2 to 4, plants inoculated with TEVΔNIb::crtB (lane 2) and co-inoculated with TEVΔNIb::LbCas12a and PVX::crFT (lane 3) or with TEVΔNIb::LbCas12a and PVX::NIb:crFT (lane 4); lane 5, DNA marker ladder with the length of some components (in bp) indicated on the right. The amplification product corresponding to the full cDNA region of LbCas12a (3,902 bp) is indicated by an arrow.