Supplementary Figure 9. *Sphingomonas* phage vB_StuS_MMDA13 gap region diagram. The positions of putative CtrA-binding sites and half-sites in *Sphingomonas* phage vB_StuS_MMDA13 gap region are displayed as green and orange boxes, respectively. Their sequences are reported, with consensus-matching bases in bold. The location of protein coding genes is also shown. The direct repeats reported in PMID:32824138 are shown as black open boxes.