



Supplementary Fig 1: Comparative Analysis and Phylogenetic Tree of cocktail phages. A. blast comparison of whole genome of five phages visualized with Clunker (<https://github.com/gamcil/clunker>). B. Maximum-Likelihood phylogeny of the *Karamvirus* and *Pseudotevenvirus* genera within Straboviridae based on a whole-genome alignment with bootstrap support of branches shown. The phages in the cocktail of this study are highlighted by red.