



Figure S3. Whole genome alignment of F10 and ΦPan70. The alignment shows conserved regions between phage F10 and ΦPan70. The phage F10 (NC_007805.1) annotation is represented with arrows: light grey arrows represent contigs and dark grey arrows represent genes. The X-axis shows the base pairs (Kbp). The Y-axis shows the conservation between the two genomes. The blocks in the alignment indicate the similarity of both phages in that region and the absence of those indicate they don't share anything in that specific region.