

Cluster MabA1

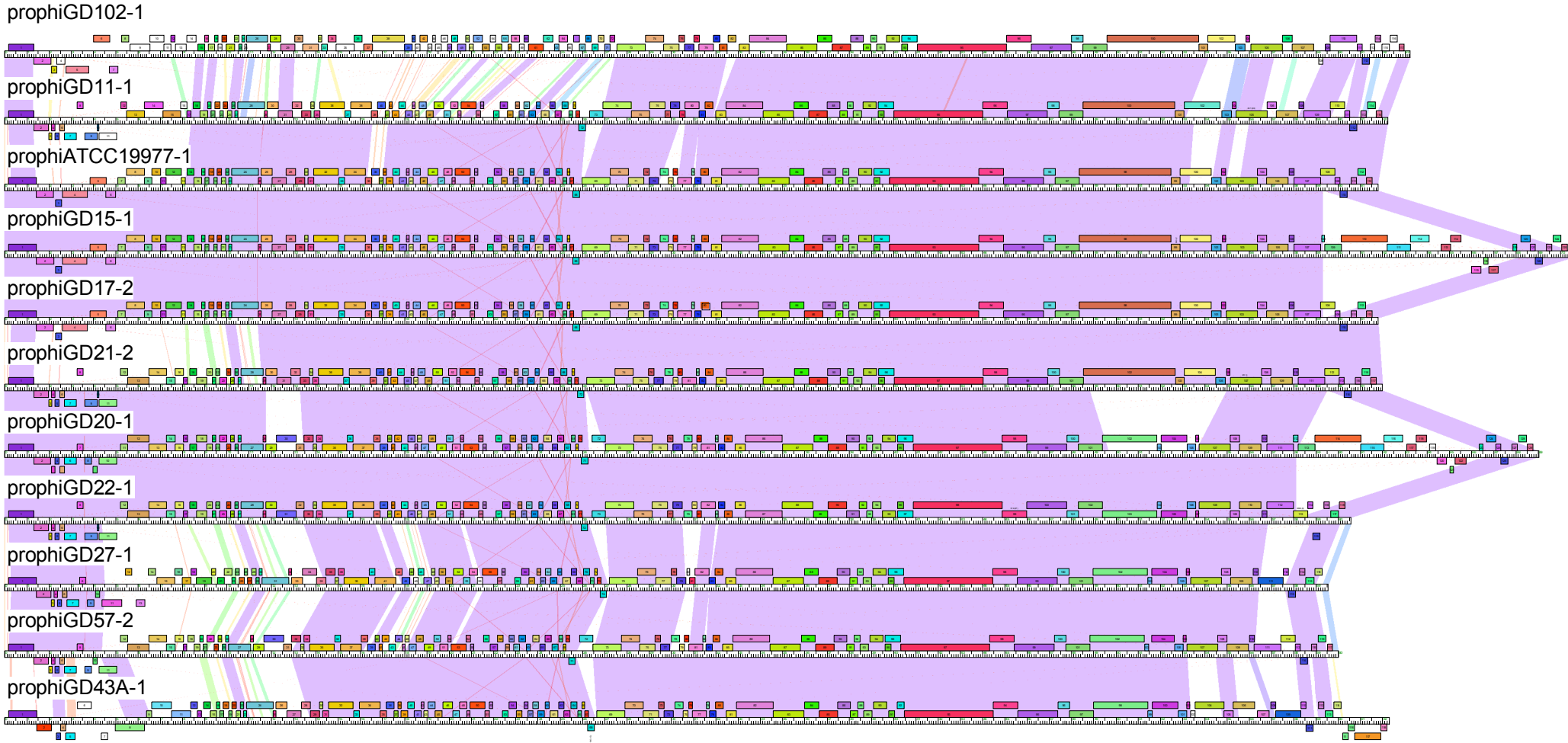


Figure S3. Genome comparisons of Cluster MabA1. Pairwise nucleotide sequence similarity is displayed by color spectrum coloring between the genomes, with violet as most similar, and red as least similar. Genes are shown as boxes either above or below the genome indicating rightward- and leftward-transcription, respectively, each genome line; boxes are colored according to gene families, they are assigned. Maps were generated using phamerator and its database Actino_prophage (version 5).