prophiGD17-2 (MabA1)
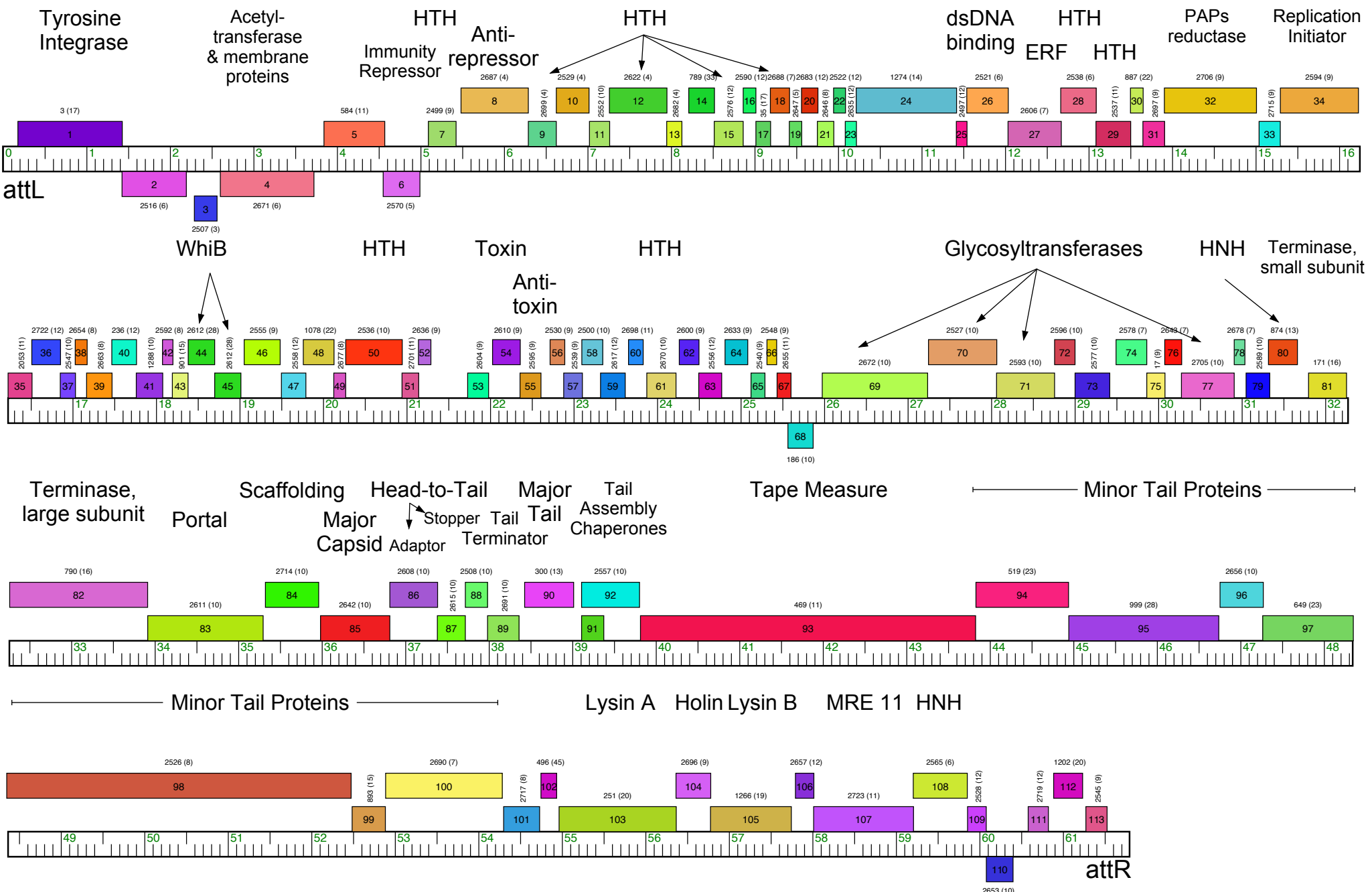

Figure S2. Genome organization of prophiGD17-2, Cluster MabA1. The genome of prophiGD17-2 is shown with predicted genes shown as boxes either above or below the genome indicating rightward- and leftward-transcription, respectively. Gene numbers are shown within each gene box. Phamily designations are shown above or below each gene with the numbers of phamily members in parentheses; genes are colored according to the phamily designations. White boxes represent 'orphams', genes with no close relatives in this dataset. Phamily assignments were determined using Phamerator and database Abscessus_prophages (version 15). Predicted gene functions are indicated.