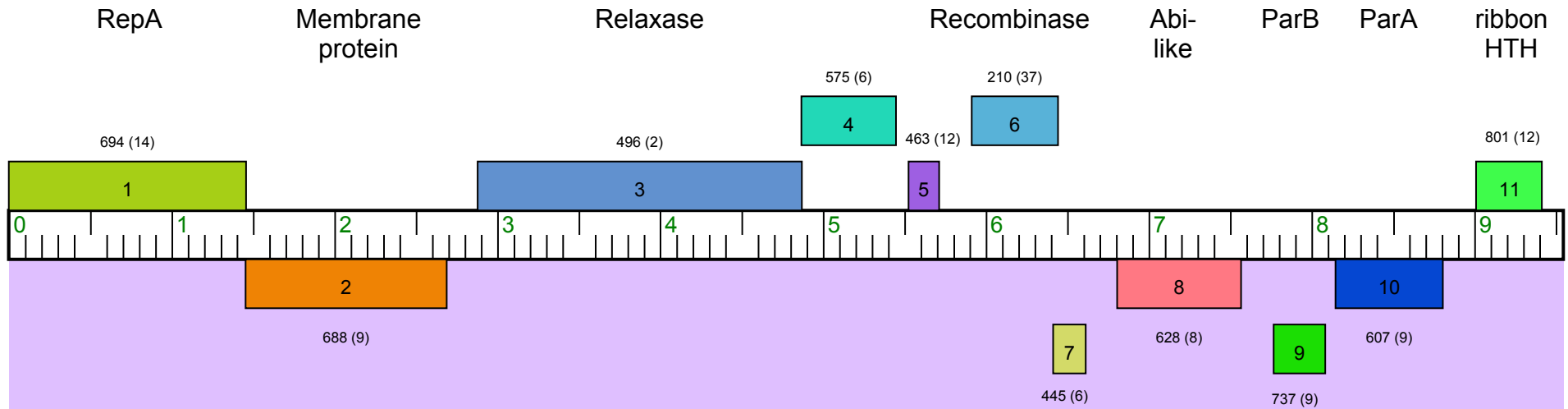


pGD08 (pA)



pGD42-2 (pA)

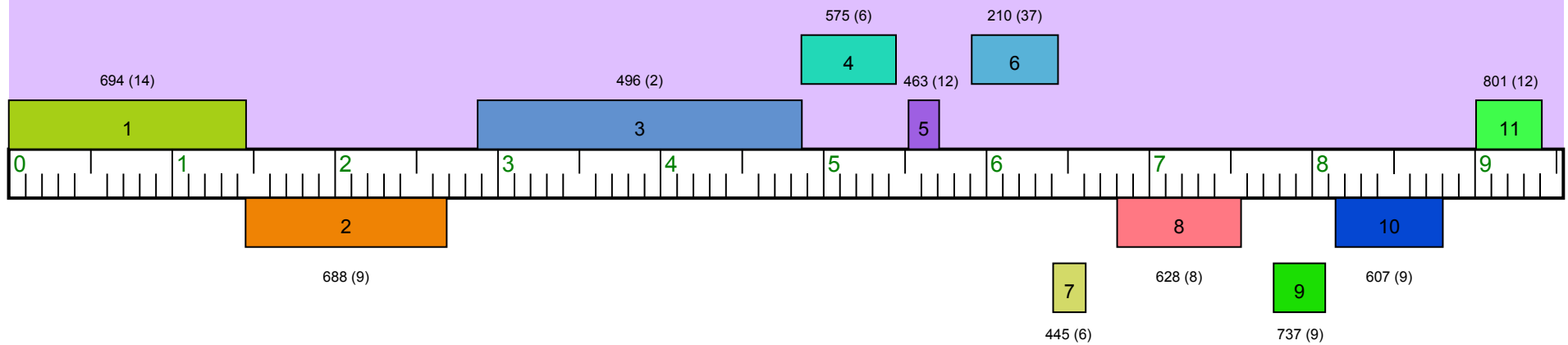


Figure S33. Genome organization and comparisons of Cluster pA. Pairwise nucleotide sequence similarity is displayed by color spectrum coloring between the genomes, with violet as most similar, and red as least similar. Genes are shown as boxes either above or below the genome indicating rightward- and leftward-transcription, respectively, each genome line; boxes are colored according to gene families, they are assigned. Maps were generated using phamerator and its database Abscessus_plasmids (version 6).