

pGD18 (pB)

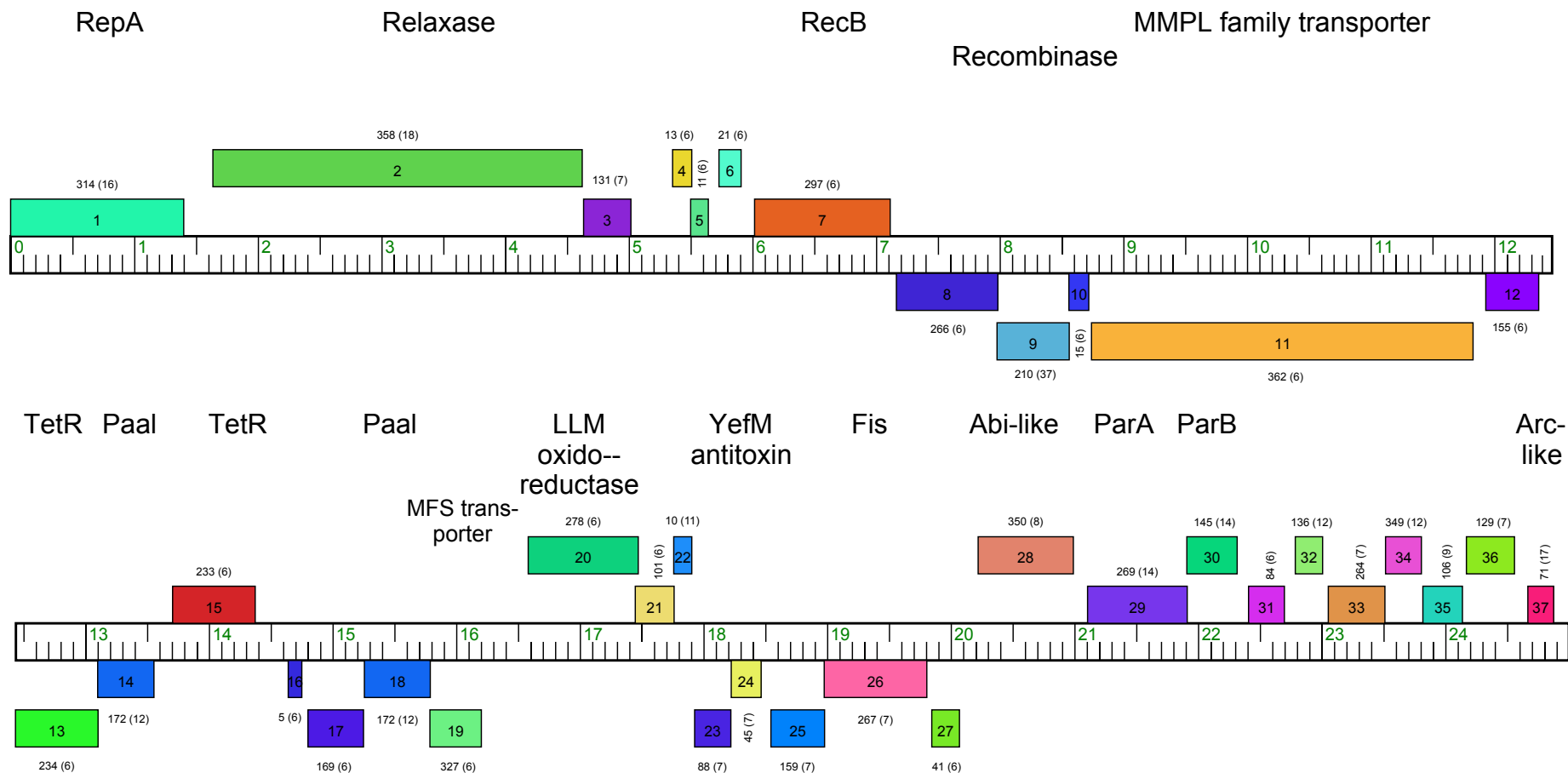


Figure S34. Genome organization of pGD18, Cluster pB. The genome of pGD18 is shown with predicted genes shown as boxes either above or below the genome indicating rightward- and leftward-transcription, respectively. Gene numbers are shown within each gene box. Family designations are shown above or below each gene with the numbers of family members in parentheses; genes are colored according to the family designations. White boxes represent 'orphans', genes with no close relatives in this dataset. Family assignments were determined using Phamerator and database Abscessus_plasmids (version 6). Predicted gene functions are indicated.