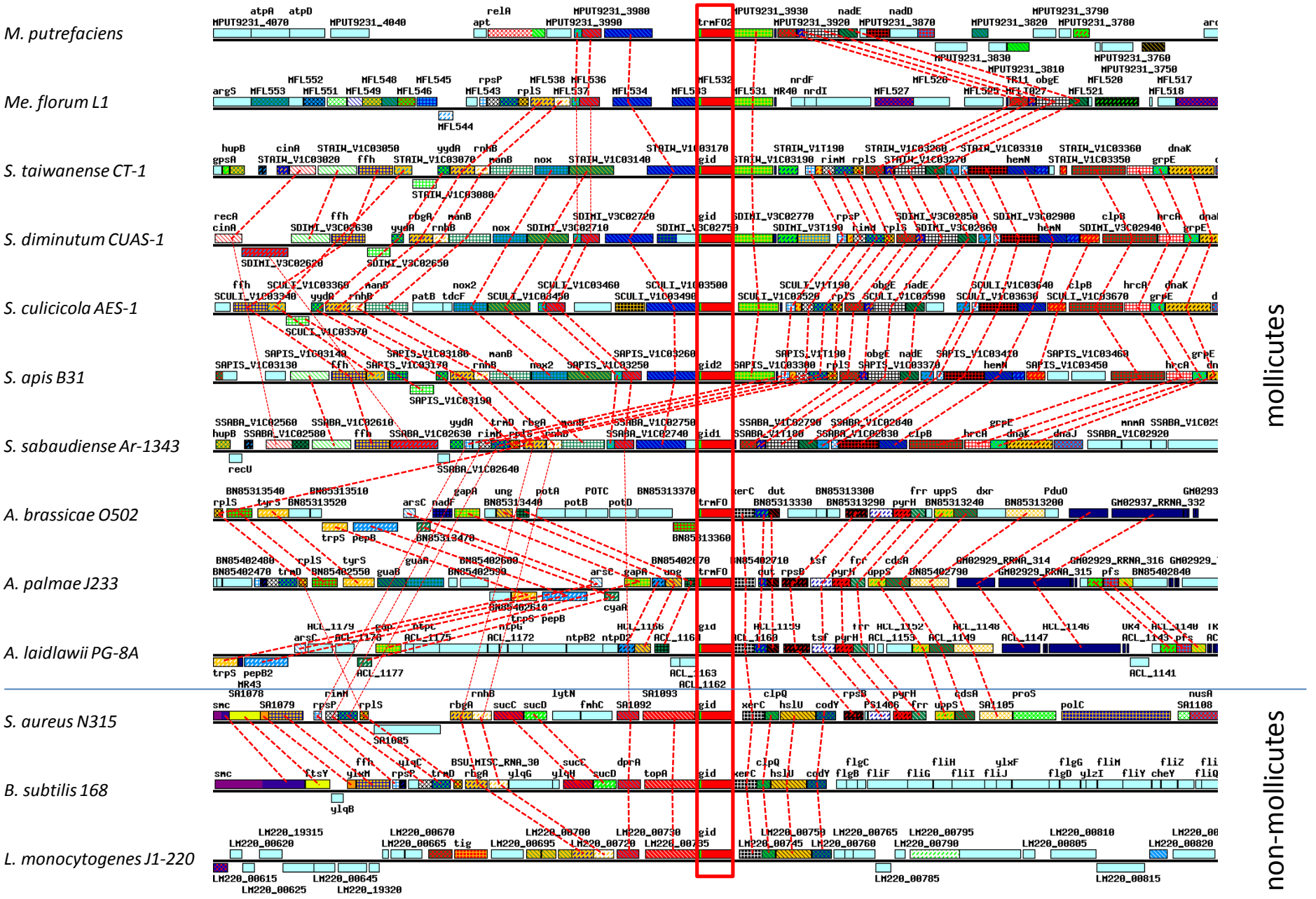

Figure S5. Genomic context of *trmFO* homologs. Analysis of synteny and genome context was performed using the MBGD database. Homologous *trmFO* genes are framed and coloured in red. Other homologs are coloured the same and connected with red dotted lines.