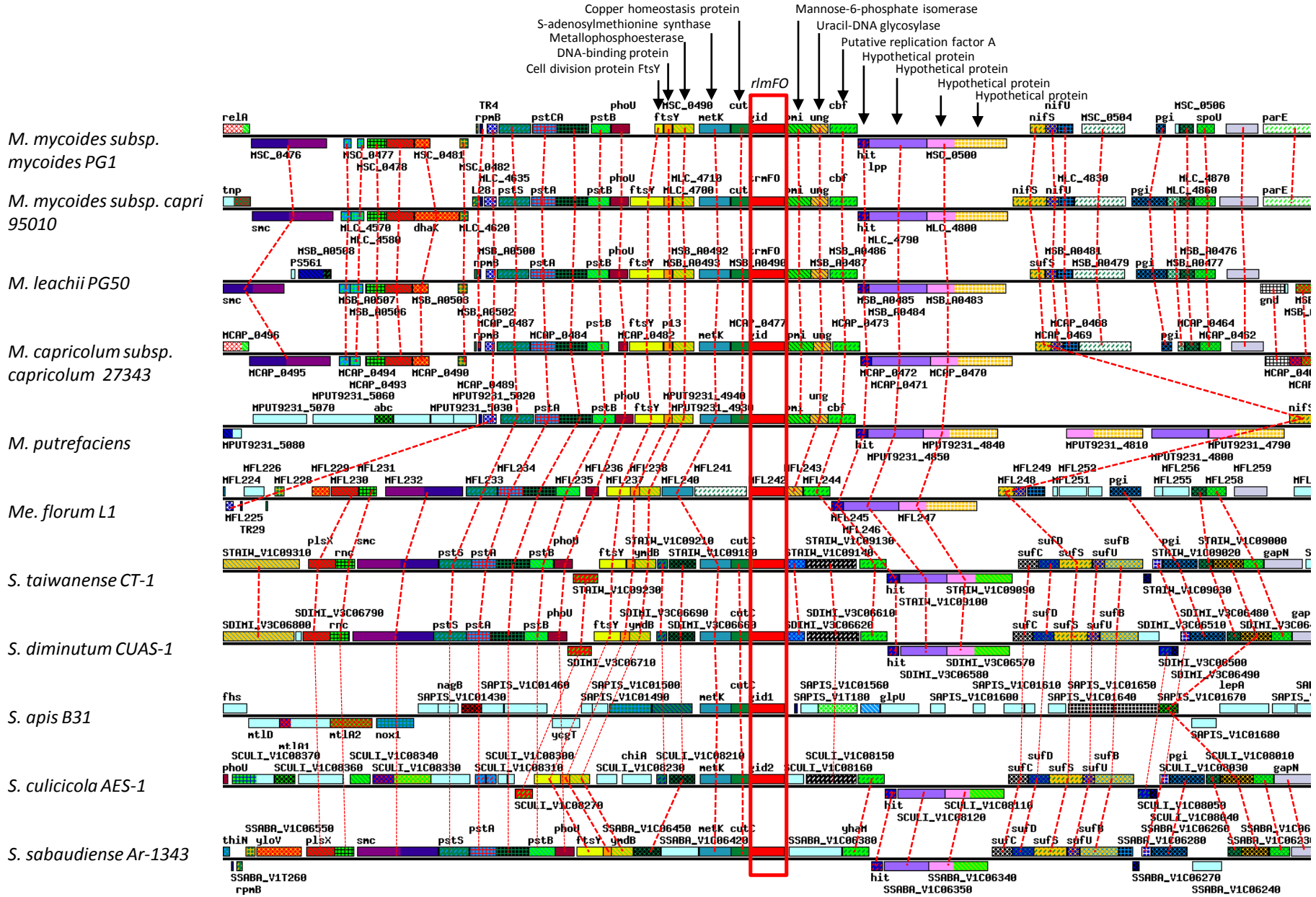

Figure S3. Genomic context of *rlmFO* homologs. Analysis of synteny and genome context was performed using the MBGD database. Homologous *rlmFO* genes are framed and coloured in red. Other homologs are coloured the same and connected with red dotted lines.