

Number of Genes

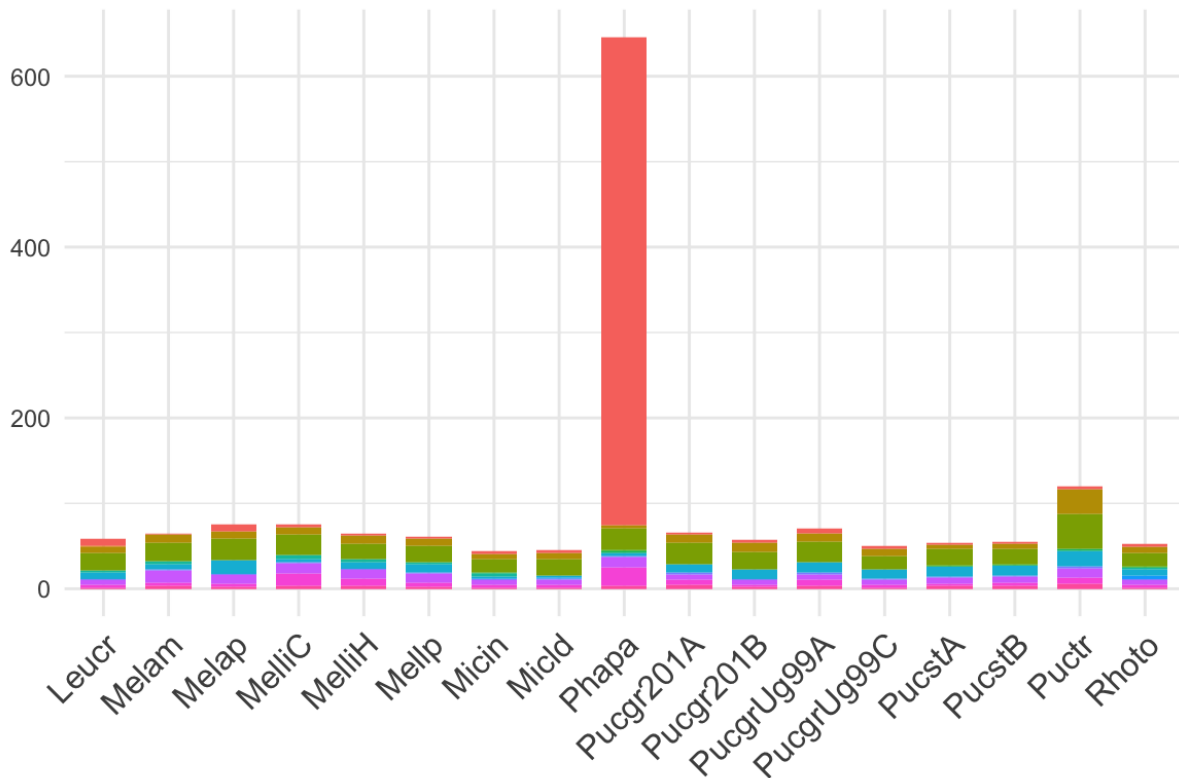


Fig. S6: TE controls involved in RNA-induced silencing complex are detected for all genomes. The number of genes involved in TE control corresponds to the RNAi pathway: Dicer-like (DCL; DCR), RNA-dependent RNA polymerase (RDP; RNT; RPAB; RPB), Argonaute (AGO; ARB) for each genome.