

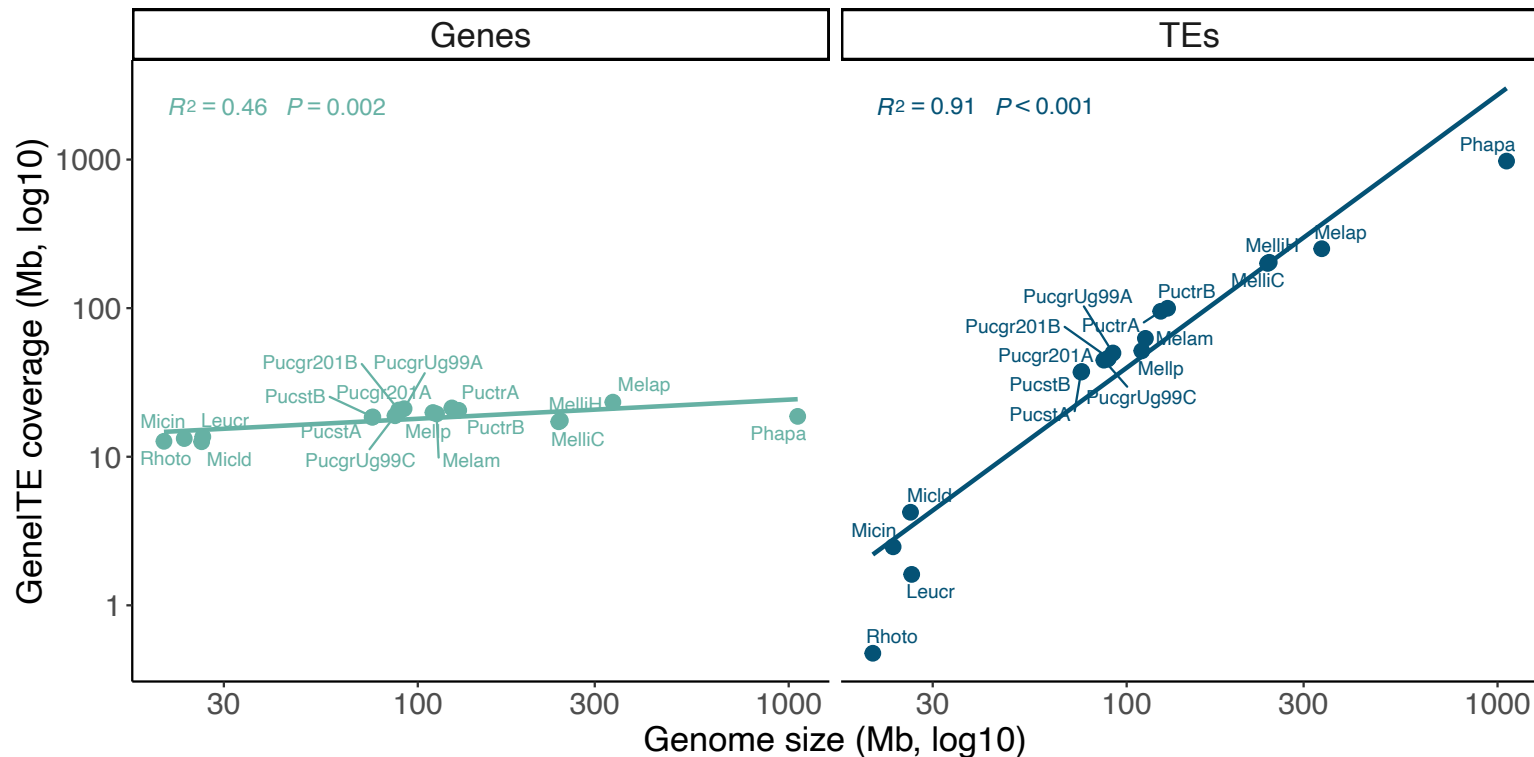


Fig. S1: Association between genome size and gene/TE content. Genome sizes (Mb) were plotted against genes or TE coverage for each Pucciniomycotina species. The lines represent the linear regression.