

Figure 1: Summary of fig wasps genomic TE contents. Purple and yellow background highlight pollinators and non-pollinators, respectively. (a) Phylogeny of samples in our study. (b) Correlation between TE lengths and genome sizes was significantly positive (Spearman correlation). (c) Summarizes the TE and non-TE lengths in the genomes, with significant difference in TE lengths (Mann-Whitney U test) but not in genome sizes between pollinators and non-pollinators. The asterisks indicated a significant difference between pollinators and non-pollinators ($P<0.05$).

