

Figure S4. Violin plots of distances between TEs and all pairs of predicted genes or pairs of different classes of predicted genes (Effectors predicted by EffectorP or predicted by SignalP (SP), and secondary metabolites (SM) predicted by AntiSMASH). Black point represent the mean distance of each class. Significativity of the Wilcoxon rank tests are indicated by NS (Not significant, p-value > 0.05), * (slightly significant, 0.05 > p-value > 0.01) or *** (highly significant, p-value <<< 0.001).

