

Supporting information 6: Relative transcription for tomato defensin genes exposed to discrete biotic/abiotic stresses. Relative mRNA abundance was assessed by real-time RT-qPCR. Data were standardized as reported in the Material and Methods segment. Significant differences between treatments were determined by ANOVA and post-hoc comparisons by least significant difference ($P < 0.05$). Bars represent means (\pm SE) of three biological replications. Vertical axis shows the relative expression to the control gene Ubiquitin.

