



Figure S2. Expression profiles of *CYP83A1* (A), *FMOGS-OX2* (B), *FMOGS-OX5* (C), *CYP81F2* (D), *CYP81F4* (E) were determined 24 and 48 h after inoculation with 5×10^5 CFU/ml of WT or mock water-inoculated control, using real-time quantitative reverse transcription PCR with gene-specific primer sets. Expression in cabbage was normalized using *BoUBQ1*. Vertical bars indicate the standard error for three biological replicates. Different letters indicate a significant difference among treatments based on a Tukey's honestly significant different test ($p < 0.05$).