

Description of Additional Supplementary files

File name: Supplementary Data 1

Description: Differential expression results between Reference and Uninfected treatments.

File name: Supplementary Data 2

Description: Differential expression results between Early and Reference, and Late and Reference treatments.

File name: Supplementary Data 3

Description: k-means cluster assignments and cluster membership for all genes.

File name: Supplementary Data 4

Description: WGCNA module membership and gene level network statistics.

File name: Supplementary Data 5

Description: Module eigengene correlations with virulence, fecundity, environmental persistence, and infection treatments.

File name: Supplementary Data 6

Description: Gene Ontology enrichment analyses for differentially expressed genes and WGCNA modules.

File name: Supplementary Data 7

Description: Expression values and annotations for hub genes highlighted in the manuscript.