



Supplemental Figure 2. Distribution of gene expression changes during infection of genotype VPRIL9. (A) upper infection court; (B) lower infection court. Only genes with significant changes (TST-FDR-adjusted $p \leq 0.01$) are included. A negative fold-change indicates a reduction by that factor. The fold changes were calculated using LMMA contrast analysis with SAS Proc Mixed using GC-RMA normalized data. The mean and median of each distribution were calculated using the log-fold changes.