



Figure S4. 257 induced enhancers associated with 263 DEGs up-regulated at 4 h post infection. a Expression of enhancer eRNA and genes; dashed line shows median gene expression prior to the infection, expression in TPM was averaged across replicates. **b** Top 15 KEGG pathway maps with the lowest FDR enriched for the genes; next to the bars are the numbers of genes in the KEGG term covered by our gene list; dashed line indicates FDR = 0.05.