



**Suppl. Fig. 1 TLR-dependence of genes that were found to be strongly (> 10-fold) induced by *H. pylori* in wt BMMs**  
 Genes regulated by *H. pylori* infection (3 h, MOI50) in wt and anergic BMMs were analyzed by microarray. Only genes with a >10-fold change in infected compared to uninfected wt BMMs were considered for further analysis. (A) Genes upregulated >1.5 fold more strongly in wt than MyD88/Trif<sup>-/-</sup> BMMs infected with *H. pylori* were defined as TLR-dependent. (B) Genes upregulated <1.5 fold less strongly were defined as no or low TLR dependence genes.