



S6 | Hierarchical clustering of 5042 genes across 785 experiments, representing 77 host-pathogen interactions. The position of the common host response, a group of genes upregulated by pathogens in many experiments and expanded in FIG. 1, is indicated by a red bar. The blue bar marks the position of the group of genes discussed in the text that are only induced in DCs. See Figure 1 (article) for details.