

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Gene Ontology Analysis of differentially expressed genes in survivor cells. The top 500 differentially expressed genes between 14 DPI uninfected and 14 DPI survivor cells as ranked by p value were uploaded to DAVID software for Biological Process (BP) Gene Ontology analysis. GO terms with Bonferroni adjusted $p < 0.05$ from this analysis were plotted in Fig 4F.