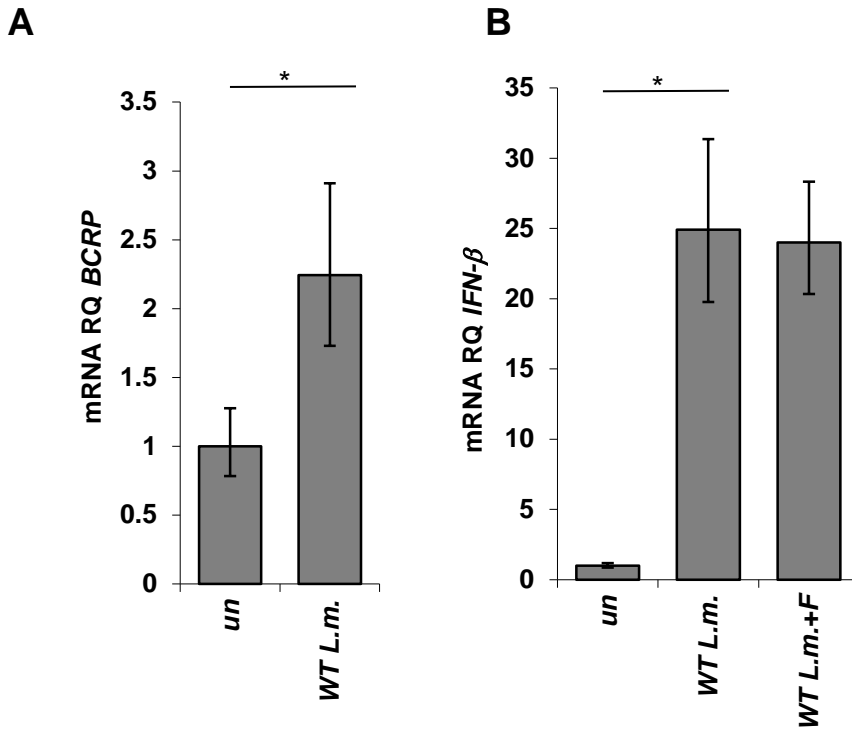**A**

B

Fig. S1. (A) RT-qPCR analysis of BCRP transcription levels in THP1 cells infected with wild-type (WT) *L. monocytogenes*. (B) RT-qPCR analysis of IFN-β transcription levels in THP1 cells infected with WT *L. monocytogenes* supplemented with 10 μM fumitremorgen C (F). Transcription levels are represented as relative quantity (RQ), relative to uninfected/untreated cells (un). Data represent at least 3 biological repeats. Error bars indicate a 95% confidence interval. * indicates p<0.01.
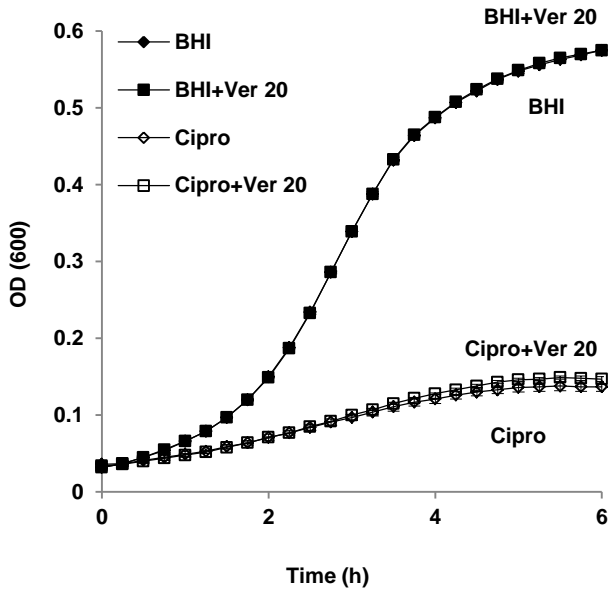
Fig. S2. Growth of WT *L. monocytogenes* in BHI medium or in BHI supplemented with 4.5 μM of ciproflaxicine (Cipro) in the presence or absence of 20 μM verapamil (Ver). Representative growth curves of 3 independent experiments are shown. Error bars represent standard deviation of a triplicate and are hidden by the symbols.
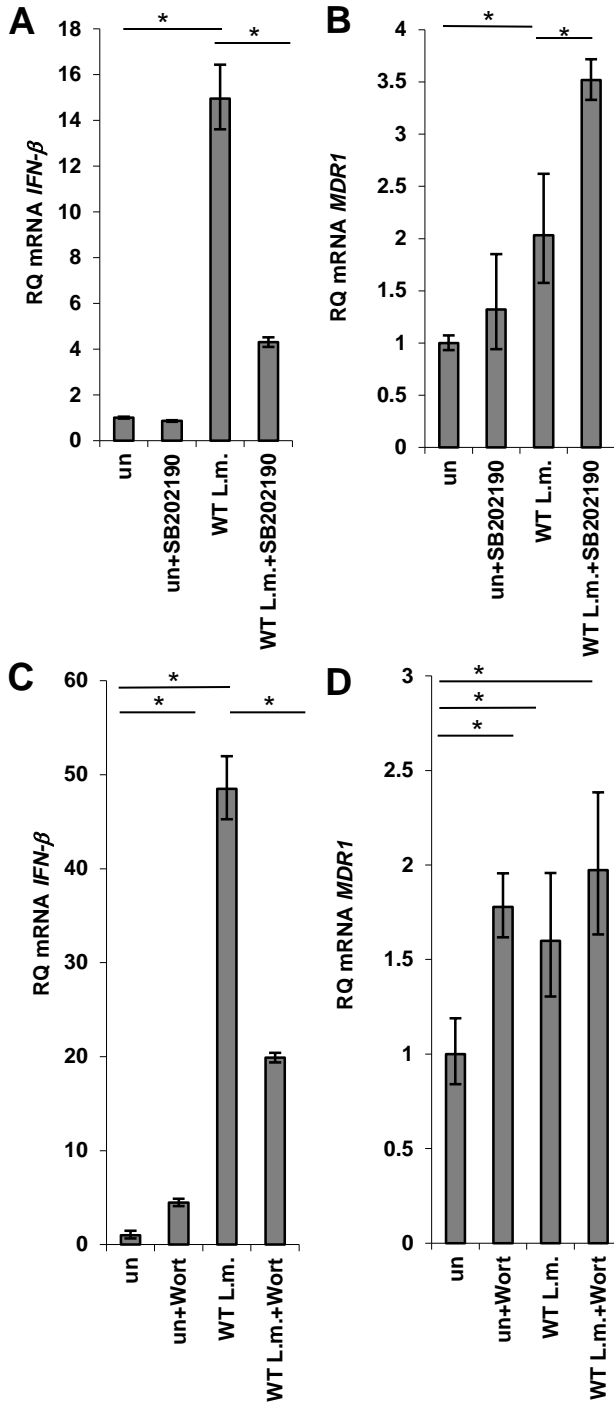
Supplementary Figure 3

Fig S3. The elicitation of IFN-β by THP1 cells in response to intracellular bacteria is mediated by p38 MAPK and PI3K. RT-qPCR analysis of IFN-β transcription levels in THP1 cells infected with WT *L. monocytogenes* and treated with 10 μM SB202190 (A, B) or 100 nM wortmannin (Wort) (C, D). Transcription levels are represented as relative quantity (RQ), relative to uninfected/untreated cells (un). Error bars indicate a 95% confidence interval. * indicates $p<0.01$. Data represents at least 3 biological repeats.