

Supplementary Information for

ZBP1 Promotes Inflammatory Responses downstream of TLR3/TLR4 via Timely Delivery of RIPK1 to TRIF

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Figures S1 and S2



Fig. S1. (A) Relative TNF and IFN β mRNA levels, normalized to GAPDH in B6 and *Ripk1*^{-/-} mouse fetal liver macrophages stimulated with LPS for indicated time points. (B) Relative TNF and IFN β mRNA levels, normalized to GAPDH in B6 and *Trif*^{-/-} and *Tram*^{-/-} macrophages stimulated with LPS for indicated time points. For experiments with mouse fetal liver macrophages, data points indicate the mean from triplicate wells of 4 technically independent experiments. For experiments with BMDMs, data points indicate the mean from triplicate the mean from triplicate wells of 4 biologically independent experiments. Analysis of variance (ANOVA) was used for comparison between groups: ns, nonsignificant (P>0.05); *P<0.05; **P<0.01; ***P<0.001; ***P<0.001.



Fig. S2. (A) Representative 60X images of FLAG-ZBP1 staining in B6, $Zbp1^{-/-}$, and $Zbp1^{-/-}$ macrophages reconstituted with low or high levels of ZBP1 or with mutant ZBP1 constructs lacking RHIM1 (ZBP1-R1), RHIM2 (ZBP1-R2) or both Za domains. (B) Quantification of mean FLAG-ZBP1 signal intensity/cell as indicated. n=100 cells/group imaged across 4-5 fields of view. (C, D) Cell death as measured by propidium iodide incorporation in indicated cells stimulated with (C) LPS or (D) LPS/5z7. Data from cell death assays and imaging experiments are representative of 3 or more independent experiments. Analysis of variance (ANOVA) was used for comparison between groups: ns, nonsignificant (P>0.05); *P<0.05; **P<0.01; ***P<0.001; ****P<0.0001.