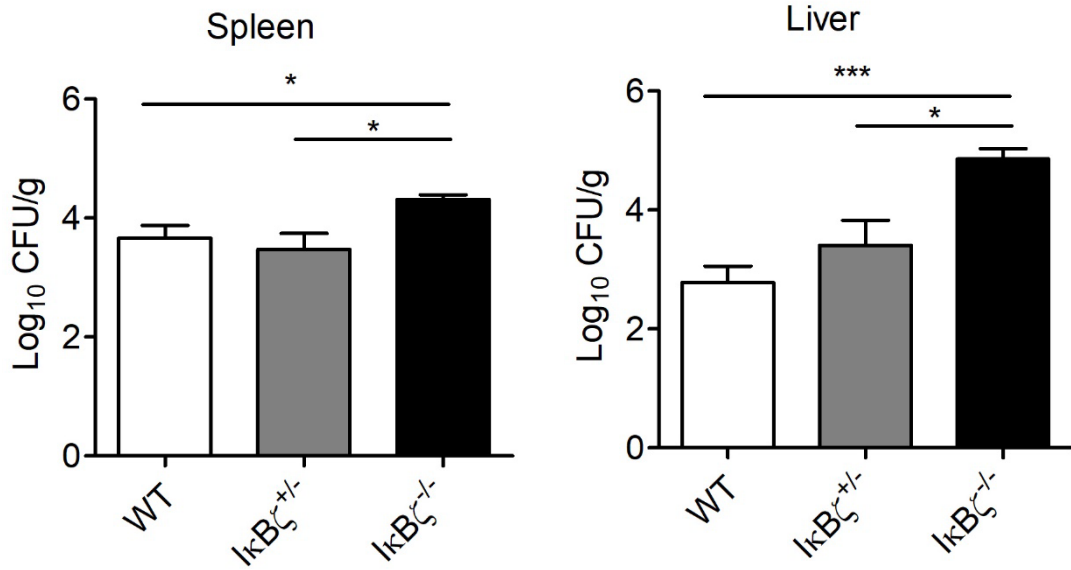


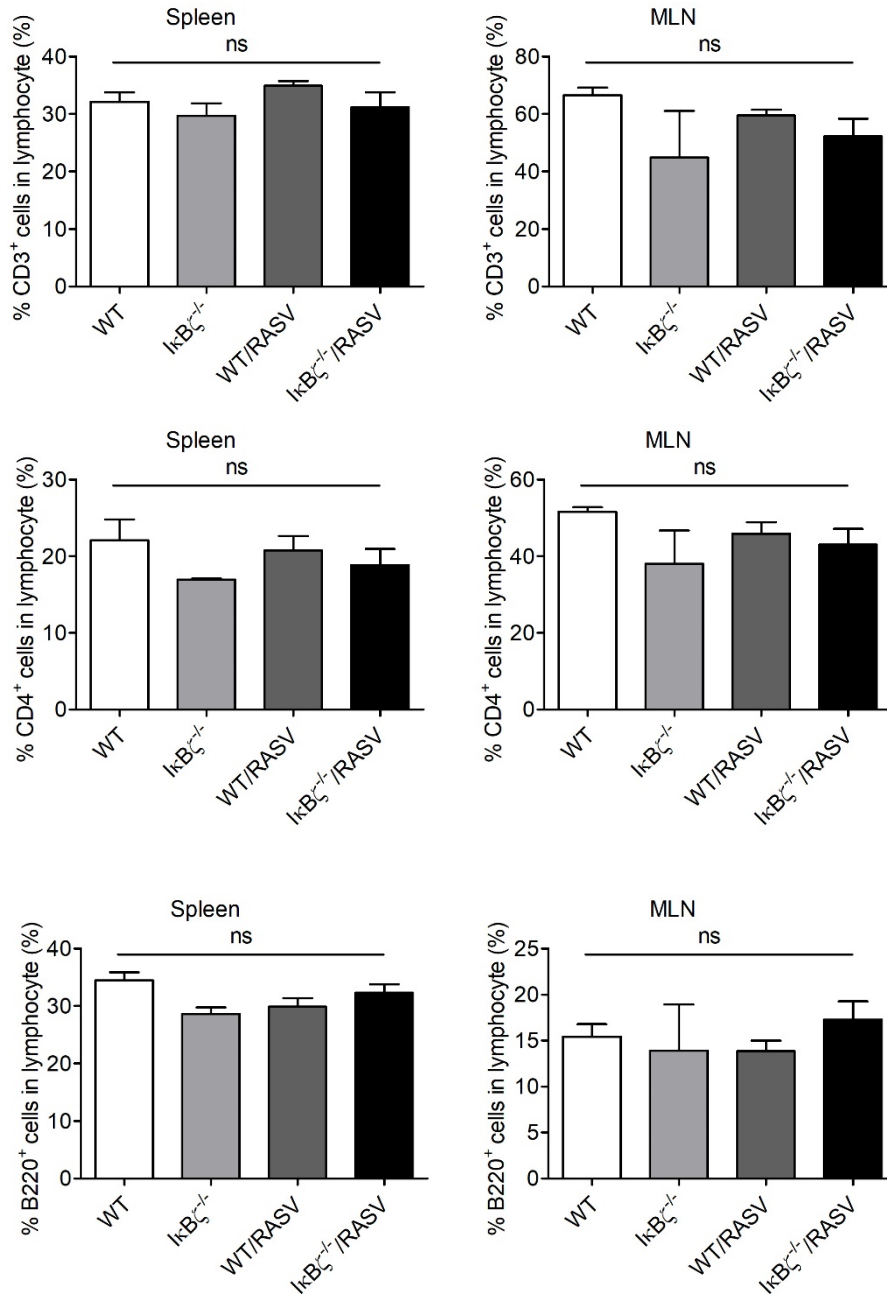
**I κ B ζ facilitates protective immunity against *Salmonella* infection via Th1
differentiation and IgG production**

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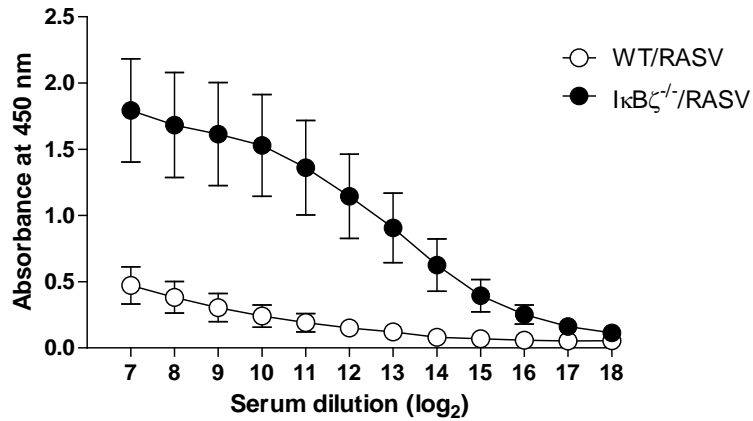
Supplementary Figure 1

Colony forming units of *Salmonella* UK-1 strain were counted with tissue homogenates obtained from liver and spleen at 9 days after oral infection of 10^7 CFU/mouse. Tissue homogenates was applied onto XLD agar plate and incubated about 24 hours. * $P < 0.05$ and *** $P < 0.001$ based on unpaired t-test.



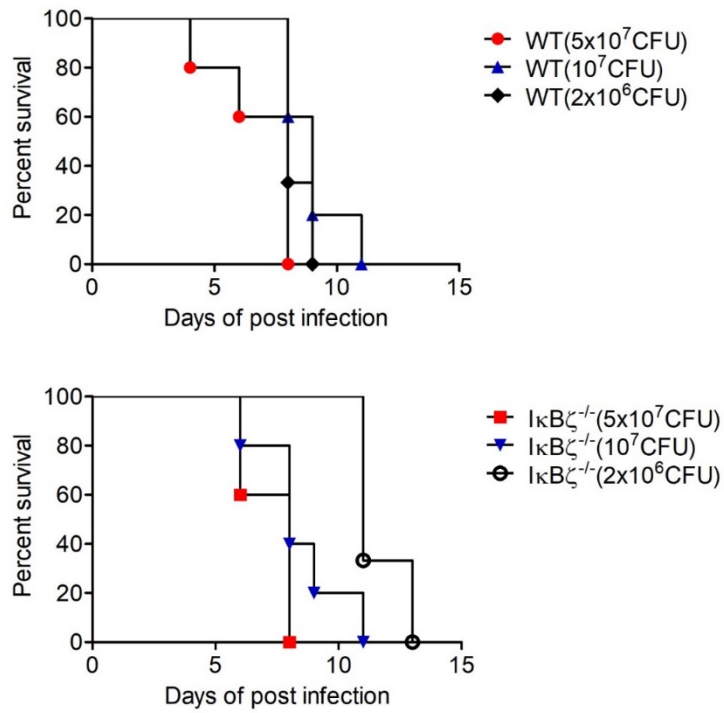
Supplementary Figure 2

CD3⁺, CD4⁺ and B220⁺ cells from spleen and lymph nodes were analyzed through flow cytometry. Spleen and mesenteric lymph nodes (MLN) were obtained after two-times administration of RASV with 14 days interval. ns, not significant; based on ANOVA with Bonferroni's multiple comparison test.



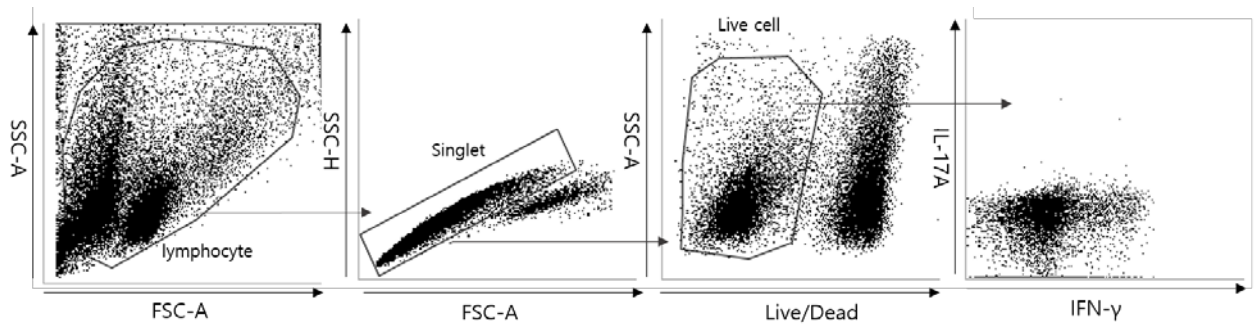
Supplementary Figure 3

Levels of *Salmonella*-specific IgG were evaluated by ELISA using whole cells of RASV as an antigen. Sera were obtained from groups of mice after two-times immunizations with RASV. For quantification of RASV-specific antibody in mouse serum, 10^7 CFU/well of RASV-whole cell bacteria were coated onto 96 well plate overnight. Further step was proceeded as described in method. Sera were next diluted by two-fold starting from 1:200 dilution.



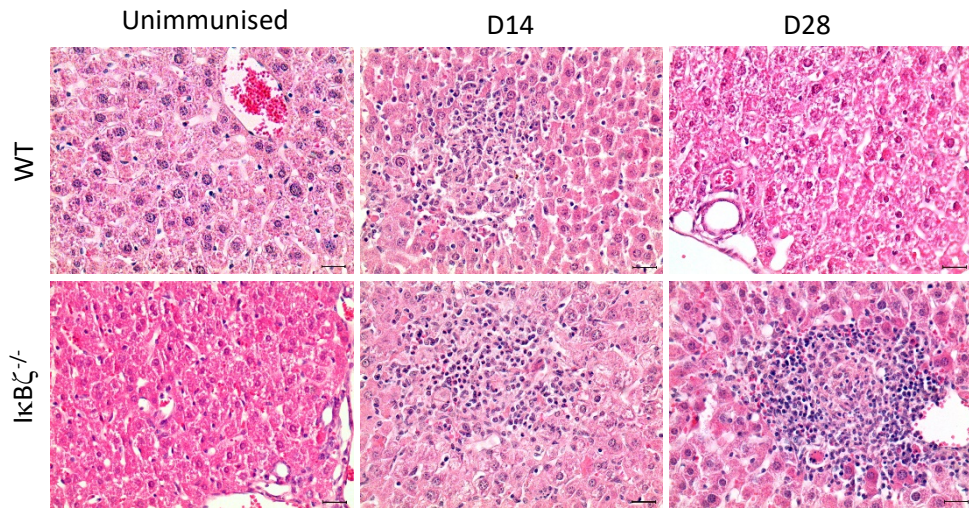
Supplementary Figure 4

Wild-type (WT) and $I\kappa B\zeta^{-/-}$ mice were orally challenged with 2×10^6 , 10^7 , and 5×10^7 CFU of a lethal wild-type *Salmonella* strain (UK-1) per mouse.



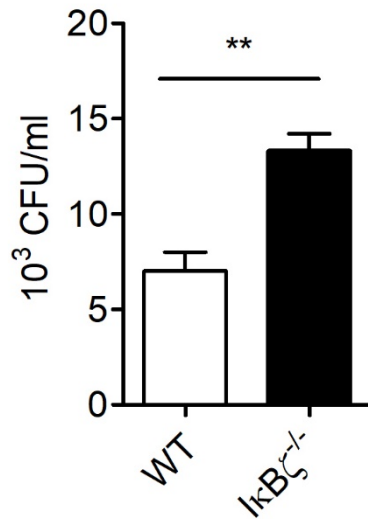
Supplementary Figure 5

Gating strategy to characterize interferon- γ and IL-17A producing cells differentiated by 10 ng/ml of IL-12.



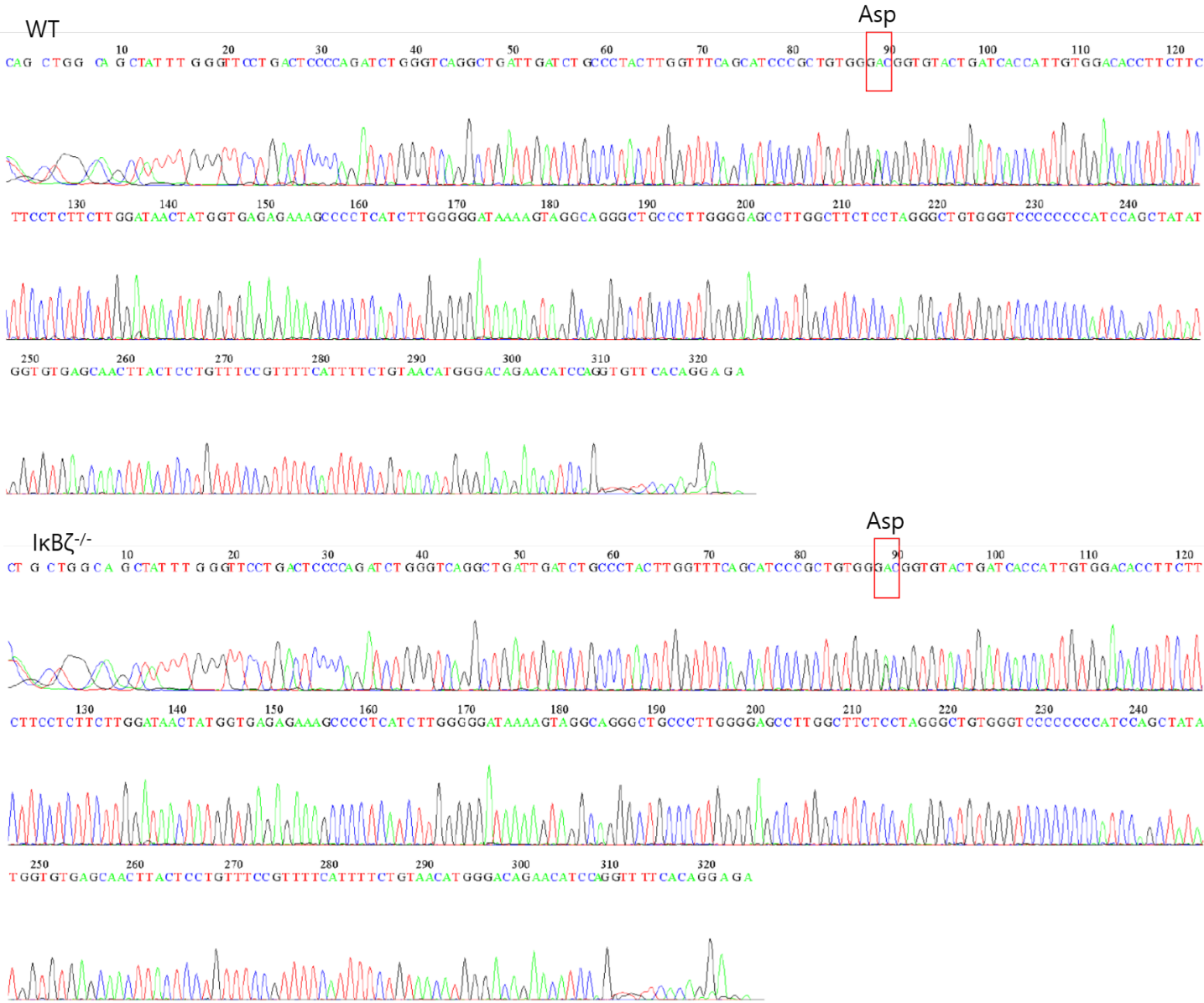
Supplementary Figure 6

High-magnified pictures of Figure 2C. Scale bar is 25 μ m.



Supplementary Figure 7

Bone marrow derived macrophages (1×10^5 cells/ well) from WT and *IκBζ*^{-/-} mice were co-cultured with 10 MOI of live UK-1 for 4 hours, and then lysed with PBS containing 1% Triton X-100. *Salmonella* CFU was checked by plating serial dilutions onto XLD agar plates.



Supplementary Figure 8

The genetic status of *Nramp1* was analyzed by genomic DNA sequencing. Tail DNA isolated from WT and *IkB ζ ^{-/-}* mice were analyzed using the following primers; Forward 5'-TTC AAC ACA ACC CAC ACT CC-3', Reverse 5'-CCT GTG ACA CCT GGA TGT TCT-3'.