

Supplementary Figure 1

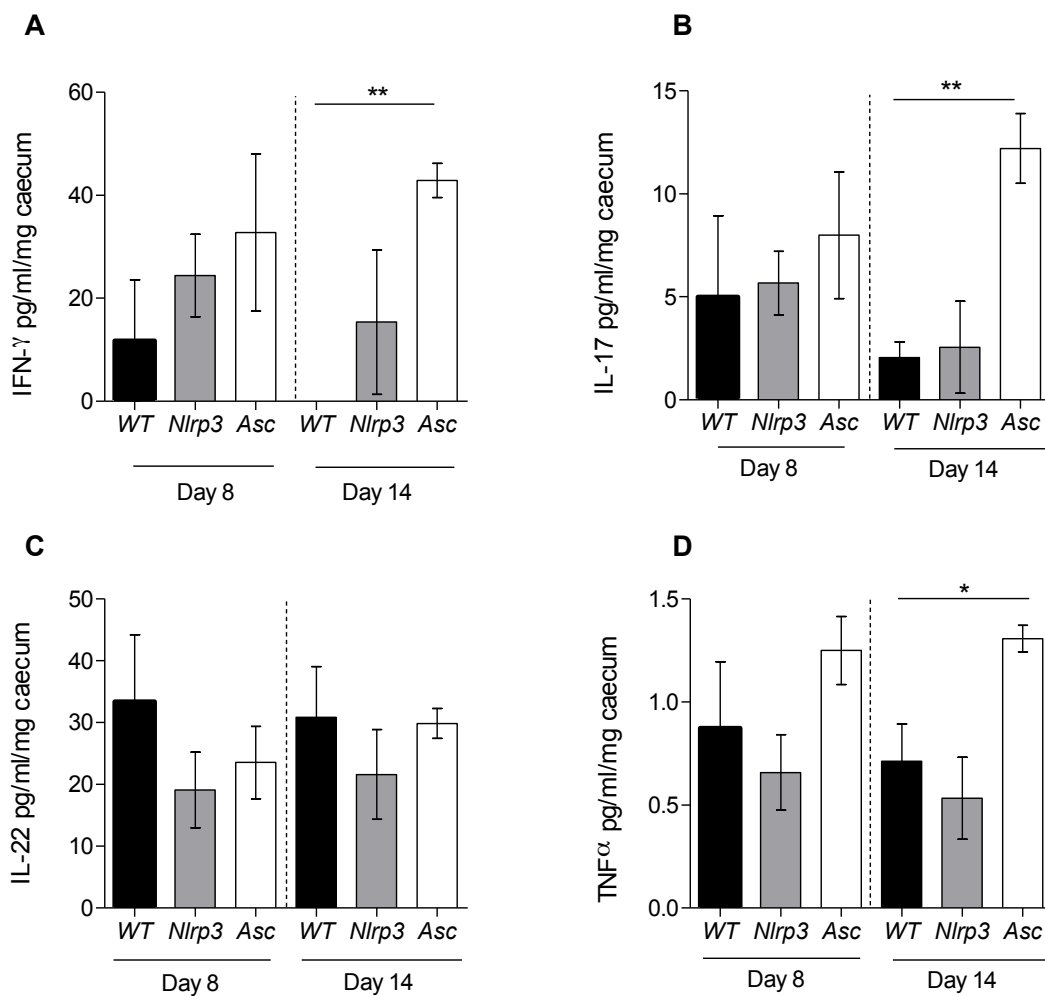


Figure S1. Induction of inflammatory cytokines upon infection with *C. rodentium*. Mice were orally infected with $\sim 10^9$ cfu of *C. rodentium* and sacrificed 8 or 14 days post-infection. Caeca were isolated and explants were cultured overnight before harvesting supernatants and analyzing for (A) IFN γ (B) IL-17 (C) IL-22 and (D) TNF α . Data represent means \pm SEM. ($n = 4$ mice per time point). Statistical analysis was performed using the Mann-Whitney test (* = $p < 0.05$). Data are representative of two independent experiments.

Supplementary Figure 2

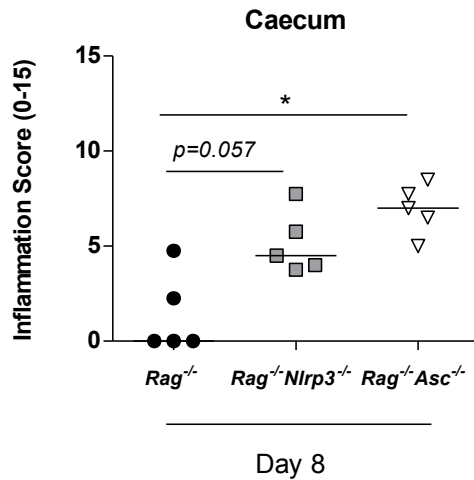


Figure S2. Nlrp3 and Asc confer protection against *C. rodentium* induced intestinal inflammation independently of the adaptive immune system. Mice were infected by oral gavage with $\sim 10^9$ cfu of *C. rodentium* and sacrificed 8 days post-infection. Inflammation scores in the caecum were assessed by scoring H & E stained histology sections and are graphically represented above. Results are from a single experiment. Horizontal bars represent the median inflammation scores. Statistical significance was determined by the Mann-Whitney test. (* = $P < 0.05$).

Supplementary Figure 3

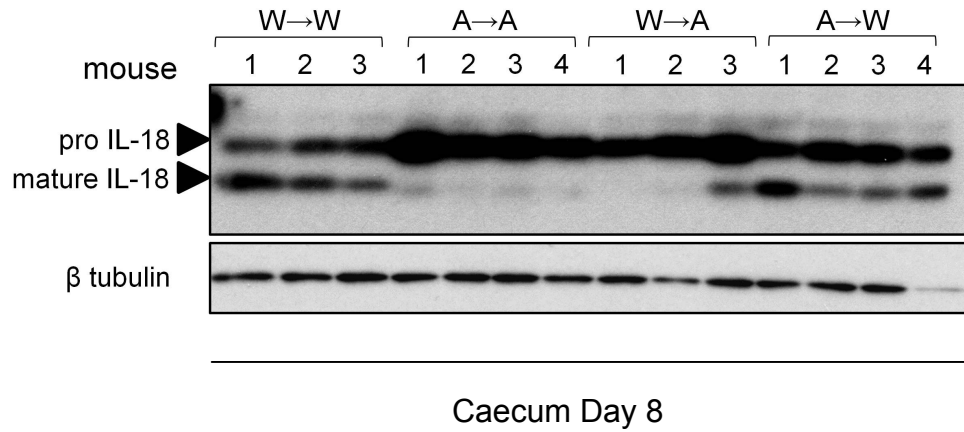


Figure S3. Non-haematopoietic cell *Asc* expression is required for IL-18 production in the intestine after *C. rodentium* infection. Irradiation bone marrow chimeras were generated as described in materials and methods, infected with $\sim 10^9$ *C. rodentium* and sacrificed 8 days later. The caeca of infected mice were then isolated and homogenized to isolate protein. Subsequently we carried out immunoblot analysis of total caecal protein extracts from WT, *Nlrp3*^{-/-} and *Asc*^{-/-} chimeras 8 days p.i., and probed with antibody against IL-18 and tubulin ($n = 3-4$ mice). Results shown are from a single experiment.

Supplementary Figure 4

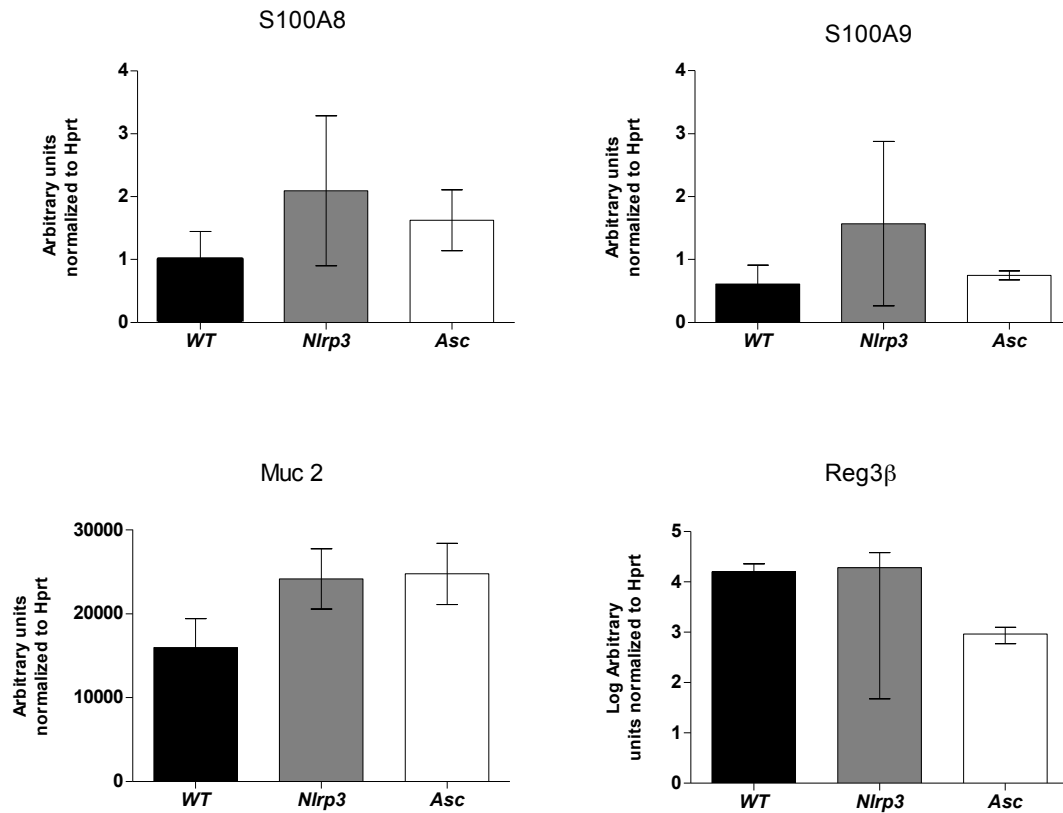


Figure S4. Comparable antimicrobial peptide (AMP) expression in inflammasome deficient mice. Expression of a variety of antimicrobial peptides and mucin were measured in the caeca of WT, *Nlrp3*^{-/-} and *Asc*^{-/-} mice at steady state by qPCR. Data are represented as mean \pm SEM. $n = 4-8$ mice and are pooled from one to two independent experiments.

Supplementary Figure 5

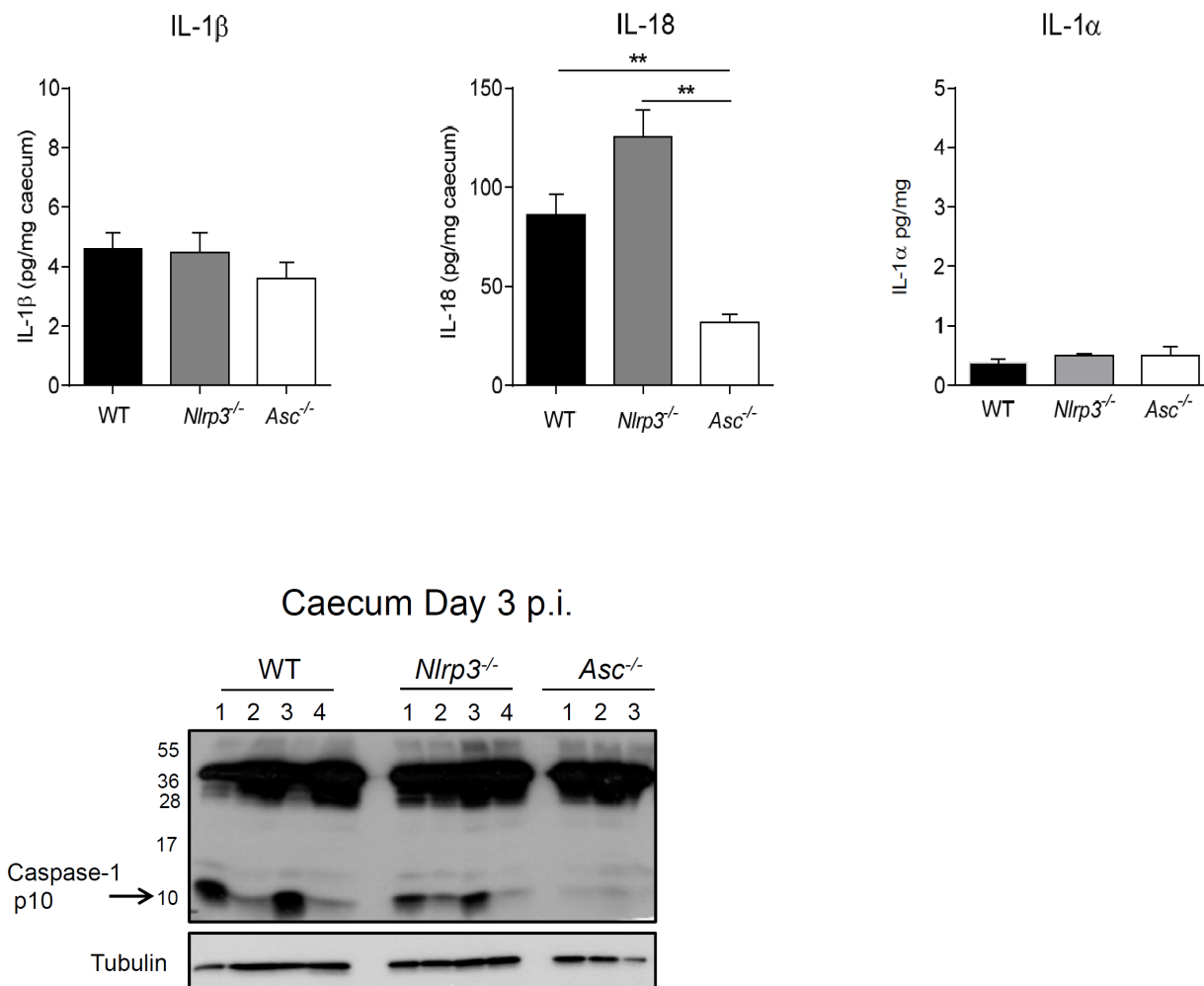


Figure S5. Early protection against *C. rodentium* is independent of caspase-1 cleaved cytokines. WT, *Nlrp3*^{-/-} and *Asc*^{-/-} mice were infected by oral gavage with $\sim 10^9$ cfu of *C. rodentium* and sacrificed 72h post infection. Caecal explants were cultured overnight before harvesting supernatant and analyzing for IL-1 β , IL-1 α and IL-18 levels by ELISA. Additionally, immunoblot analysis of caecal protein extracts from WT, *Nlrp3*^{-/-} and *Asc*^{-/-} mice was performed with antibodies to caspase-1 p10 and tubulin. Data are represented as means \pm SEM. $n = 3-4$ mice from a single experiment representative of at least two independent experiments.

Supplementary Figure 6

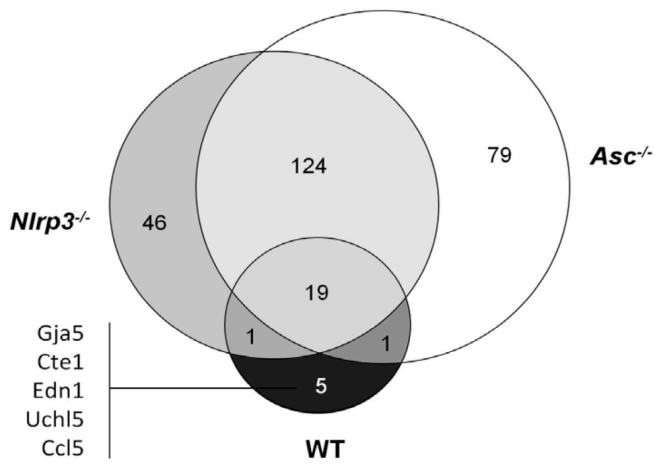


Figure S6. Genes induced upon infection in *Nlrp3*^{-/-}, *Asc*^{-/-}, and WT. WT, *Nlrp3*^{-/-}, and *Asc*^{-/-} mice were infected with $\sim 10^9$ *C. rodentium* and sacrificed 72h p.i. ($n = 3$). RNA was isolated from caeca and gene expression profiled using Illumina Mouse WG6v2 Expression BeadChip. Venn diagram of the genes induced upon infection (at least 1.5-fold 72h p.i. vs uninfected).

Supplementary Figure 7

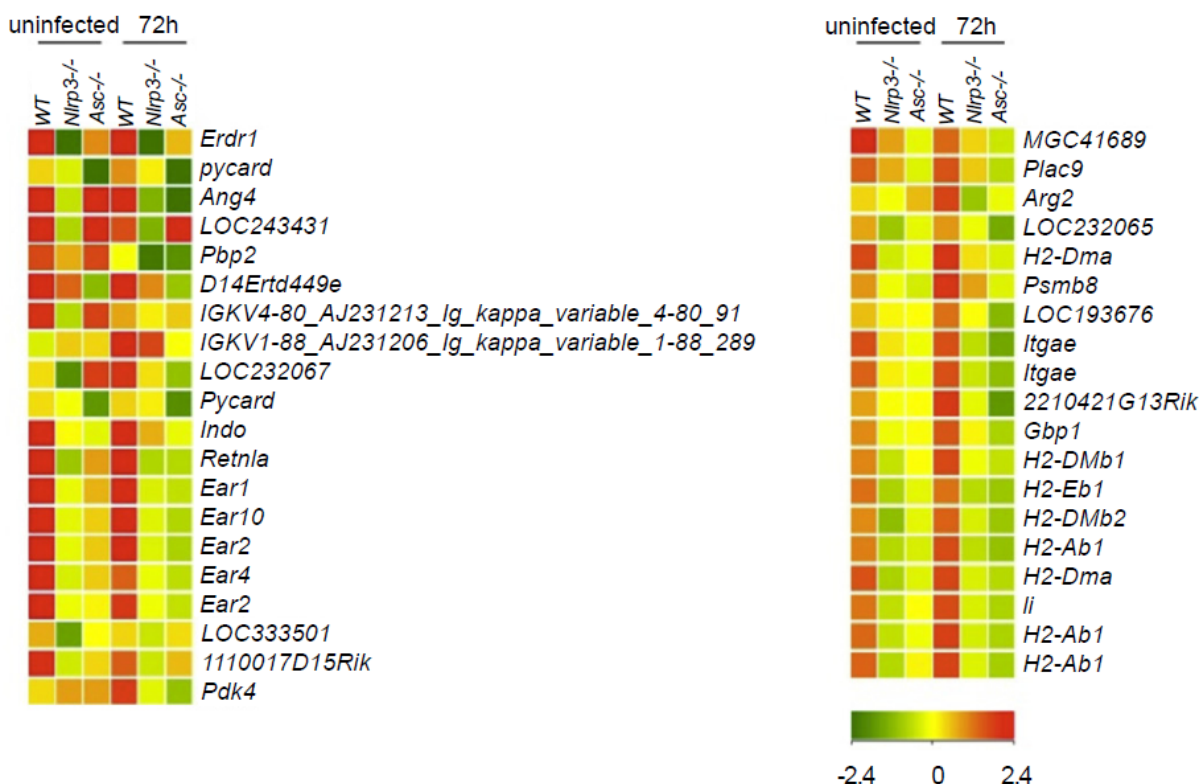


Figure S7. Differentially expressed genes between WT, *Nlrp3*^{-/-} and *Asc*^{-/-} mice. Cohorts of WT, *Nlrp3*^{-/-}, and *Asc*^{-/-} mice were either left uninfected or orally infected with $\sim 10^9$ *C. rodentium* and sacrificed at 3 days p.i. ($n = 3$ per group). Thereafter caecal tissue RNA was isolated and subjected to genome wide transcriptional profiling.

Depicted above is a heat map representing 33 unique genes significantly expressed at a lower level (at least 2-fold) in either *Nlrp3*^{-/-} or *Asc*^{-/-} relative to WT mice, during steady state (uninfected) and/or 3 days p.i.. Colour range (green to red) represent log transformed normalized intensity values.