

## Supplemental material

## Chen et al., https://doi.org/10.1084/jem.20181031

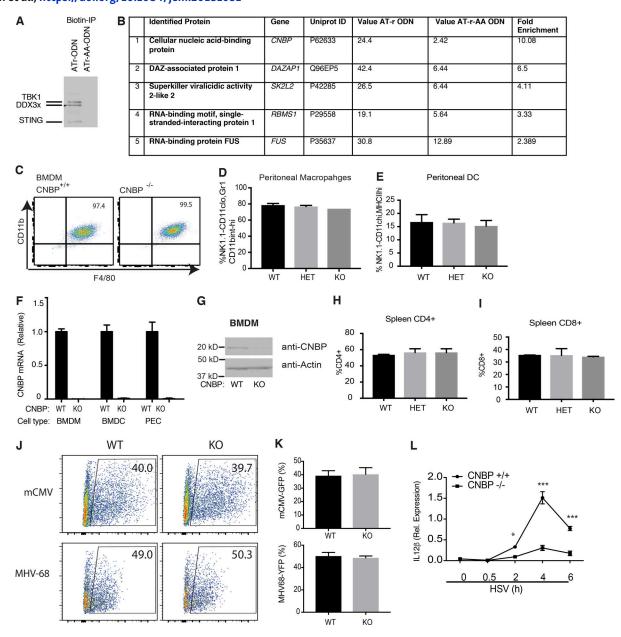


Figure S1. **Characterization of** *Cnbp***-deficient mice. (A)** Immunoblotting was performed to identify known components of the DNA-sensing pathway in ATr-ODN and a nonstimulatory ODN (ATr-AA ODN) immunoprecipitation (IP) samples from macrophage cytosolic extracts. **(B)** LC-MS analysis of proteins identified. The top five most enriched proteins identified in ATr-ODN and nonstimulatory ATr-AA-ODN complexes. **(C)** CD11b/F480 levels were measured by flow cytometry on macrophage cultures differentiated in vitro from bone marrow progenitors. **(D and E)** Percentages of PECs (D) and peritoneal DCs (E) were analyzed by their specific lineage markers from WT ( $Cnbp^{+/+}$ ), HET ( $Cnbp^{+/-}$ ), and KO ( $Cnbp^{-/-}$ ) mice. **(F and G)** CNBP expression levels in WT and KO cells were detected through RT-PCR (F) and Western blotting (G). **(H and I)** Percentages of CD4 and CD8 T cells analyzed by their specific lineage markers from WT ( $Cnbp^{+/-}$ ), HET ( $Cnbp^{+/-}$ ), and KO ( $Cnbp^{-/-}$ ) mice. **(J and K)** The replication of reporter virus mCMV-GFP or murine gammaherpesvirus 68–YFP in CNBP WT or deficient MEFs. **(L)** qRT-PCR analysis of IL12 $\beta$  mRNA in Cnbp WT or KO BMDMs mock infected or infected with HSV at different time courses. Error bars represent SD of triplicate technical replicates. All data are representative of three independent experiments with similar results. \*, P < 0.05; \*\*\*, P < 0.001.



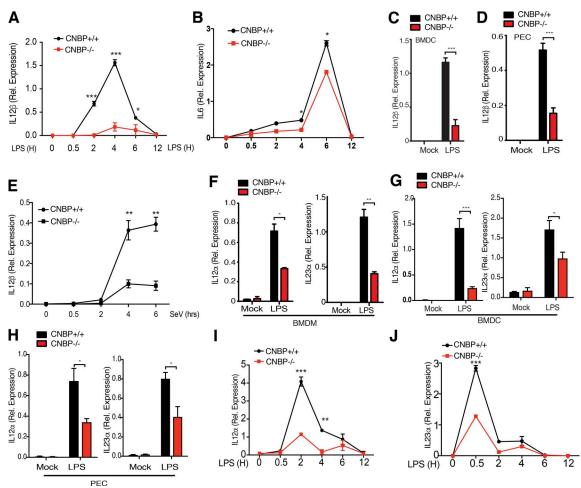


Figure S2. **Cytokine analysis in CNBP-KO cells after stimulation with various ligands at different time points. (A and B)** qRT-PCR analysis of IL-12 $\beta$  (A) or IL6 (B) mRNA in Cnbp WT or KO BMDMs left unstimulated or stimulated with LPS at different time courses. **(C and D)** qRT-PCR analysis of IL-12 $\beta$  in Cnbp WT or KO BMDC (C) or PECs (D) left unstimulated or stimulated with LPS. **(F)** qRT-PCR analysis of IL-12 $\beta$  mRNA in Cnbp WT or KO BMDMs (F), BMDCs (G), or PECs (H) left unstimulated or stimulated or stimulated or stimulated or stimulated or stimulated with LPS. **(I and J)** qRT-PCR analysis of IL-12 $\alpha$  (I) or IL-23 $\alpha$  (J) mRNA in Cnbp WT or KO BMDMs left unstimulated or stimulated with LPS at different time courses. Error bars represent SD of triplicate technical replicates. All data are representative of three independent experiments with similar results. \*, P < 0.05; \*\*, P < 0.01; \*\*\*, P < 0.001.

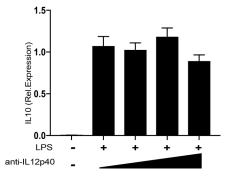


Figure S3. **IL-10 induction was not affected by IL-12.** Primary macrophages were incubated with anti–IL-12 neutralizing antibody before LPS stimulation and IL-10 mRNA expression levels were determined by qPCR. Error bars represent SD of triplicate technical replicates. All data are representative of three independent experiments with similar results.



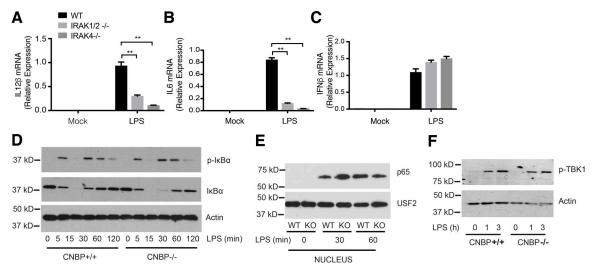


Figure S4. Analysis of the NF- $\kappa$ B signaling and cytokine inductions in IRAK1/2-/- or IRAK4-/- BMDMs. (A–C) qRT-PCR analysis of IL12 $\beta$  (A), IL6 (B), and IFN $\beta$  (C) in IRAK1/2-/- or IRAK4-/- BMDMs left unstimulated or stimulated with LPS. (D) Immunoblot analysis of phospho-I $\kappa$ B $\alpha$  or total I $\kappa$ B $\alpha$  in whole cell lysates of Cnbp WT or KO BMDMs stimulated for different times with LPS. (E) Nuclear extracts were analyzed for p65 by Western blotting in CNBP WT and KO BMDMs treated with LPS. (F) Immunoblot analysis of p-TBK1 in whole cell lysates of Cnbp WT or KO BMDMs stimulated for various times with LPS. Error bars represent SD of triplicate technical replicates. All data are representative of three independent experiments with similar results. \*\*, P < 0.01.



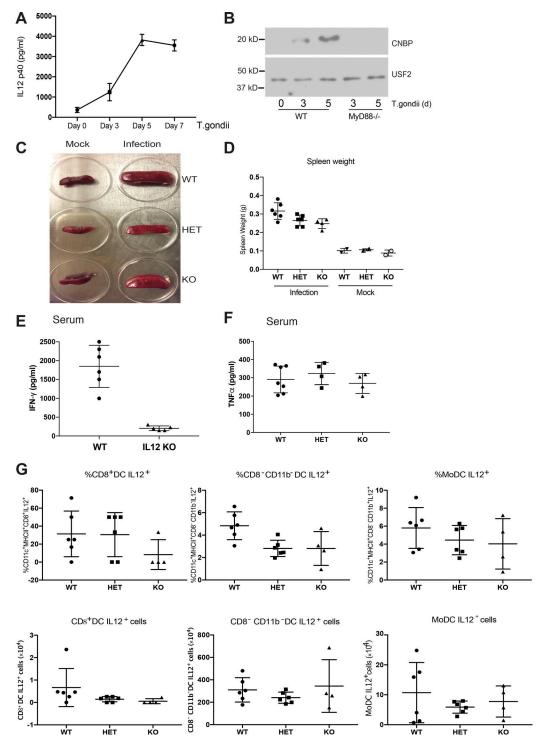


Figure S5. Analysis of CNBP-deficient mice after *T. gondii* infection. (A) ELISA quantification of IL-12p40 levels in the serum after *T. gondii* infection at different time points after infection. (B) Nuclear extracts were analyzed for Cnbp by Western blotting in spleen cells from WT and MyD88<sup>-/-</sup> mice after *T. gondii* infection at different time after infection. (C and D) Gross appearance (C) and weight (D) of the spleen in the indicated mice. (E) ELISA quantification of IL12p40 levels in the serum after *T. gondii* infection in WT and IL12<sup>-/-</sup> (KO) mice. (F) ELISA quantification of TNF-α levels in the serum after *T. gondii* infection in WT, CNBP<sup>+/-</sup> (HET), and CNBP<sup>-/-</sup> (KO) mice 7 d after *T. gondii* infection. Each symbol represents an individual mouse; small horizontal lines indicate the mean. All data are representative of at least two to three independent experiments with similar results.



Table S1. sgRNA, ChIP, and qPCR primers

| Gene           | Forward primer (5'-3')    | Reverse primer (5'-3')    |
|----------------|---------------------------|---------------------------|
| qPCR primers   |                           |                           |
| IL12b          | GGAAGCACGGCAGCAGAATA      | AACTTGAGGGAGAAGTAGGAATGG  |
| IL12a          | ACGAGAGTTGCCTGGCTACTAG    | CCTCATAGATGCTACCAAGGCAC   |
| IL23a          | CATGCTAGCCTGGAACGCACAT    | ACTGGCTGTTGTCCTTGAGTCC    |
| IL10           | CGGGAAGACAATAACTGCACCC    | CGGTTAGCAGTATGTTGTCCAGC   |
| IL6            | TACCACTTCACAAGTCGGAGGC    | CTGCAAGTGCATCATCGTTGTTC   |
| TNF-α          | GGTGCCTATGTCTCAGCCTCTT    | GCCATAGAACTGATGAGAGGGAG   |
| IFN-β          | ATAAGCAGCTCCAG            | CTGTCTGCTGGTGGAGTTCA      |
| CNBP           | ATCTGCTACCGCTGTGGTGAGT    | GCCACCTCTACCGCAGTTATAG    |
| CCL4           | TTCCTGCTGTTTCTCTTACACCT   | CTGTCTGCCTCTTTTGGTCAG     |
| CXCL9          | GGAGTTCGAGGAACCCTAGTG     | GGGATTTGTAGTGGATCGTGC     |
| IL15           | GTAGGTCTCCCTAAAACAGAGGC   | TCCAGGAGAAAGCAGTTCATTGC   |
| iNOS2          | GTTCTCAGCCCAACAATACAAGA   | GTGGACGGGTCGATGTCAC       |
| IRF7           | CAGCGAGTGCTGTTTGGAGAC     | AAGTTCGTACACCTTATGCGG     |
| IL4i1          | GGATGAGAAGACAGGCTGGATAG   | GCAGCTTCACATTATGCACCTCC   |
| Med21          | AGTGTGGTCCTCCTGCCTCTTT    | CTGTTCGTGCAATCAGTGCTGC    |
| GAPDH          | TGGCAAAGTGGAGATTGTTGCC    | AAGATGGTGATGGGCTTCCCG     |
| ChIP primers   |                           |                           |
| IL12b pro -600 | CATAAGAGACGCCCTCAAA       | TGTTTGGCTCCACGTACC        |
| IL12b pro -500 | TGGAGCCAAACAGGAGGTAAT     | GTGTTACAGGCCCAAAGAATAAA   |
| IL12b pro -400 | GGGCCTGTAACACCTACTTATTT   | CGGCAATGGCTAACCTCTC       |
| IL12b pro -300 | GCCGCCTCTATTCACCTTAG      | GACGTCGAAATCCCAGGTTA      |
| IL12b pro -200 | TCGACGTCTATATTCCCTCTGTAT  | CTAGTCTCAATTGCAACACTGAAA  |
| IL12b pro -100 | TTCAGTGTTGCAATTGAGACTA    | TGTTCCTTCTGCTGCCTTG       |
| sgRNAs         |                           |                           |
| hCNBP sgRNA1   | CACCGCCGTGTGCAGACCCGCGTG  | AAACCACGCGGTCTGCACACGGC   |
| hCNBP sgRNA2   | CACCGCGTCCGAGTCTCCGCCGCTG | AAACCAGCGGCGGAGACTCGGACGG |