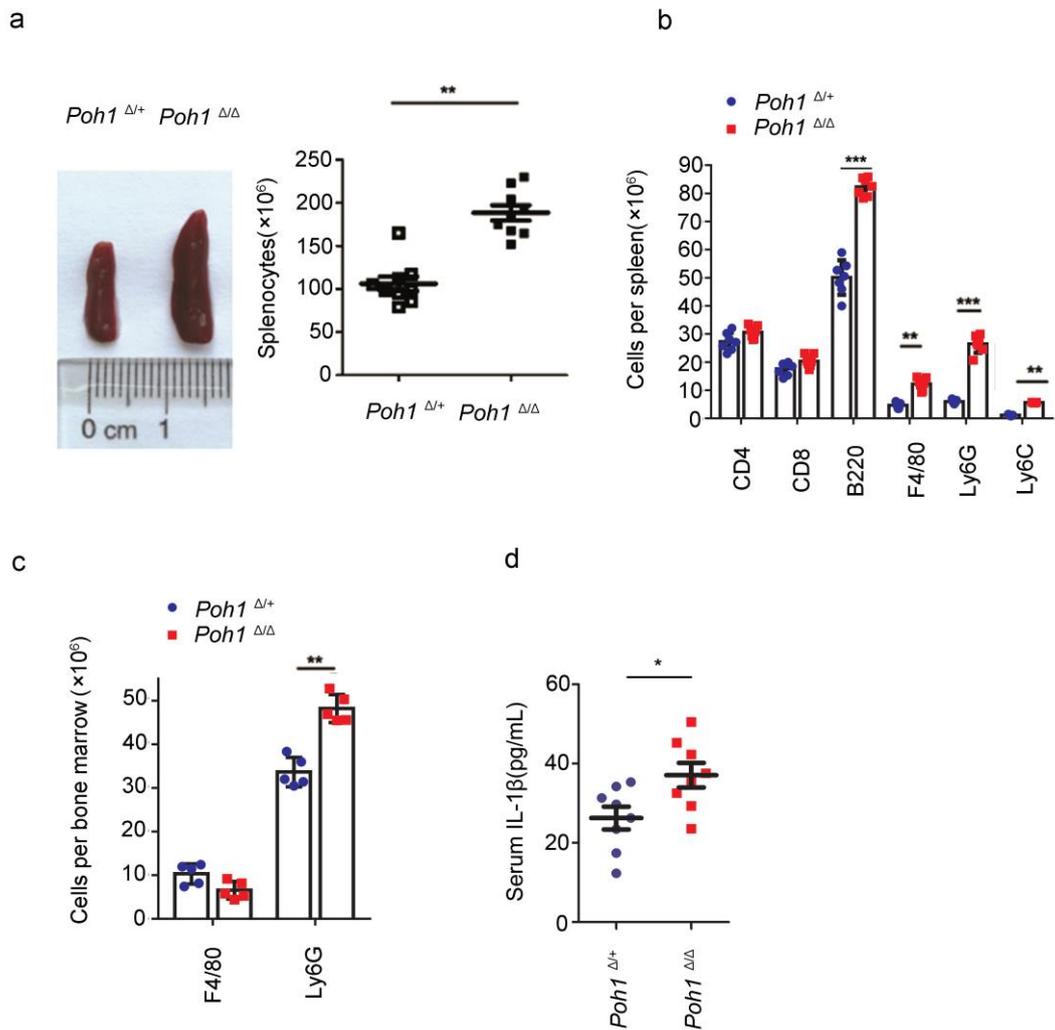


Supplementary Information

POH1 deubiquitinates pro-interleukin-1 β and restricts inflammasome activity

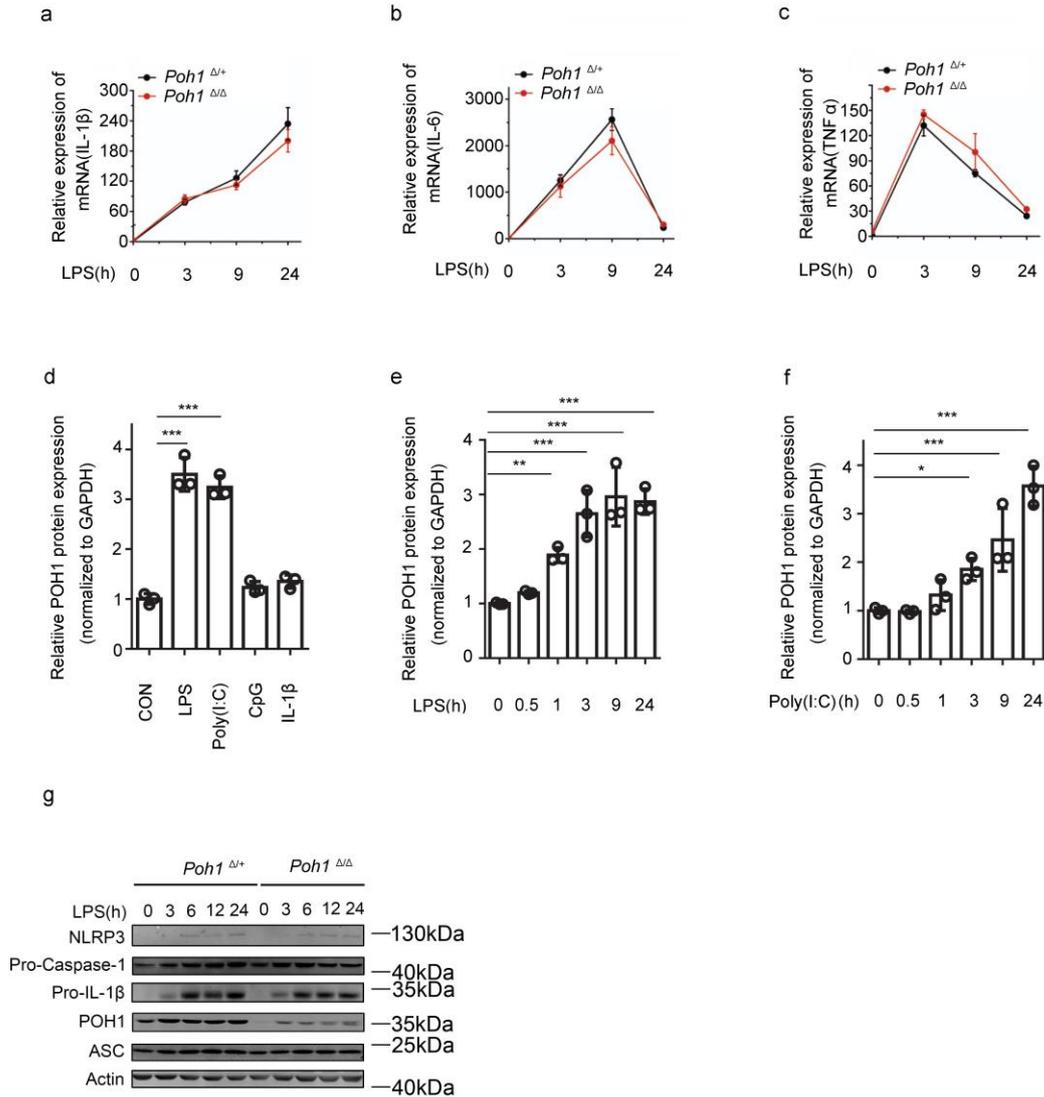
Zhang et. al.



S1

Supplementary Figure 1. **a** Morphology (left panel) and cell numbers (right panel) of spleens in *Poh1*^{Δ/+} and *Poh1*^{Δ/Δ} mice (n=9 per group). **b, c** Single cell suspensions were prepared from the spleen and bone marrow, (**b**) absolute numbers of CD4⁺ or CD8⁺ T cells, B220⁺ B cells, CD11b⁺ F4/80⁺ macrophages, CD11b⁺ Ly6G⁺ neutrophils and CD11b⁺ Ly6C⁺ monocytes in the spleens of *Poh1*^{Δ/+} and *Poh1*^{Δ/Δ} mice were measured (n=7 per group); or (**c**) absolute numbers of CD11b⁺ F4/80⁺ macrophages and CD11b⁺ Ly6G⁺ neutrophils in the bone marrow (n=5 per group) were measured. **d** ELISA quantification of the IL-1 β levels in the serum of *Poh1*^{Δ/+} and *Poh1*^{Δ/Δ} mice

($n=8$ mice per group). Data are pooled from (a) three or (b-d) two independent experiments (mean \pm s.d. in a-c; mean \pm s.e.m. in d), * $p<0.05$, ** $p<0.01$, *** $p<0.001$ (two tailed Student's t-test).

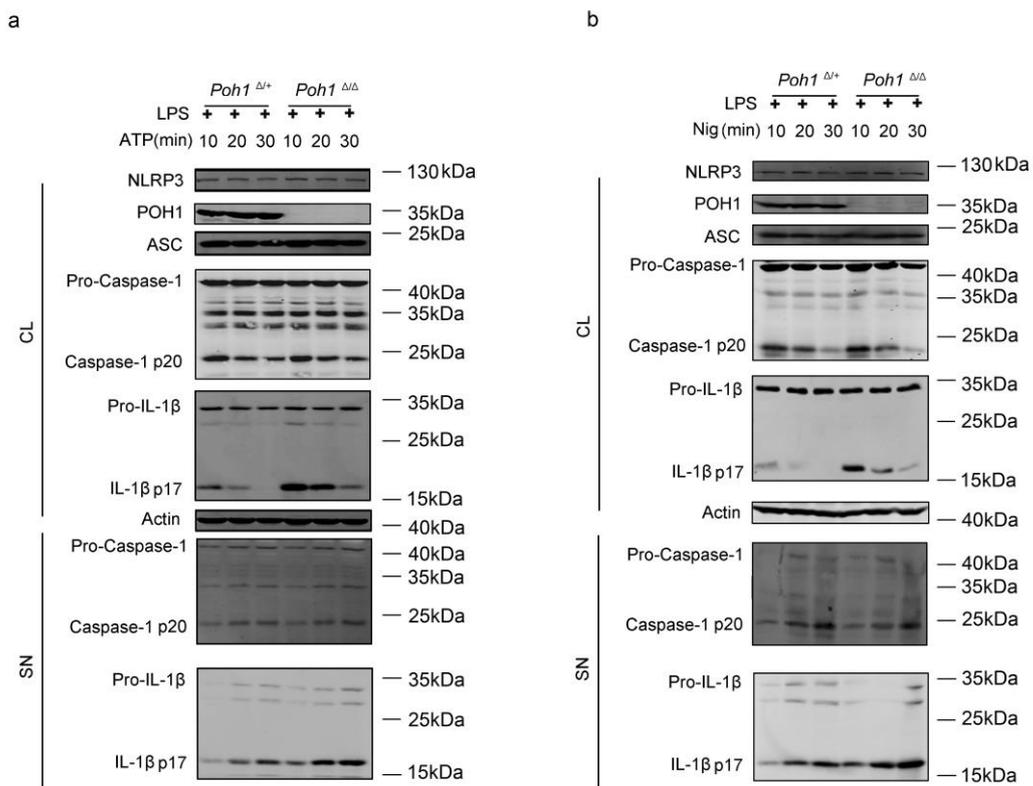


S2

Supplementary Figure 2. a-c BMDMs from *Poh1* ^{$\Delta/+$} and *Poh1* ^{Δ/Δ} mice were treated with LPS as indicated, then the mRNA levels of (a) IL-1 β , (b) IL-6 and (c) TNF α were measured by RT-PCR.

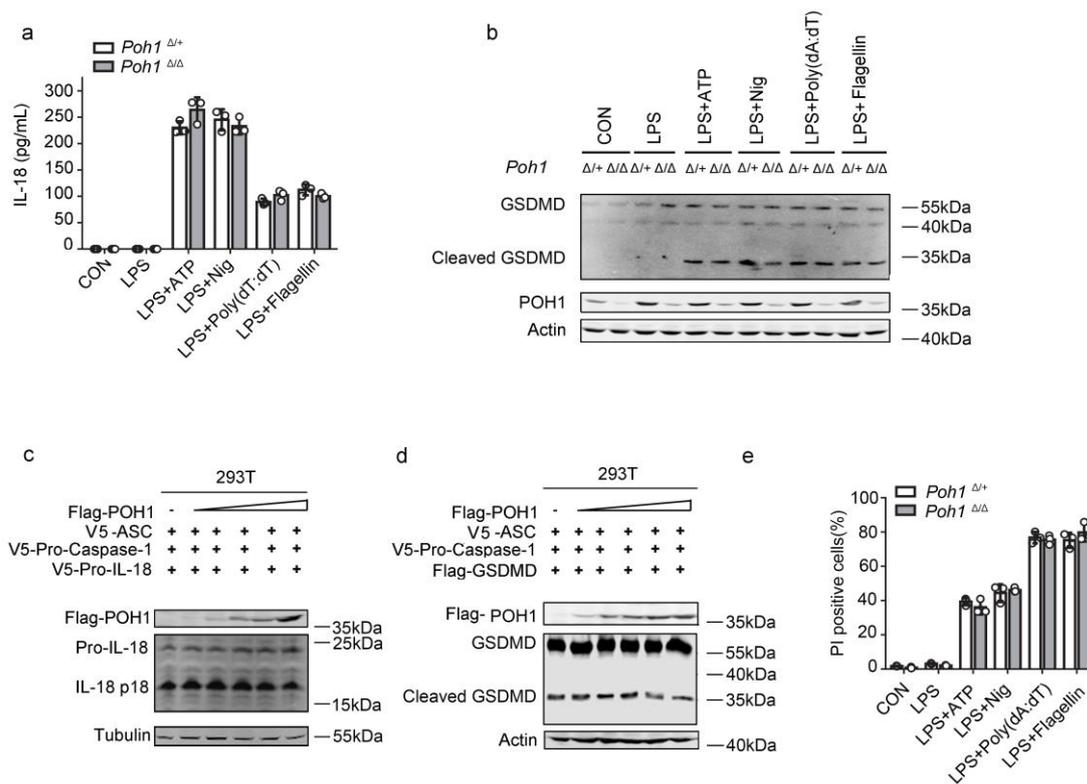
Similar results were obtained from three independent experiments. The results represent the

mean \pm s.d. of three independent sets of experiments. **d-f** Quantification of western blot signal intensity for **(d)** Fig. 2a, **(e)** Fig. 2b and **(f)** Fig. 2c from three independent experiments, data presented as the mean \pm s.d. , * p <0.05, ** p <0.01, *** p <0.001 (one-way ANOVA with Dunnett's post-hoc test). **g** BMDMs from *Pohl* ^{Δ /+} and *Pohl* ^{Δ / Δ} mice were treated with LPS as indicated, then the cell lysates were IB with the indicated antibodies. Similar results were obtained from three independent experiments.



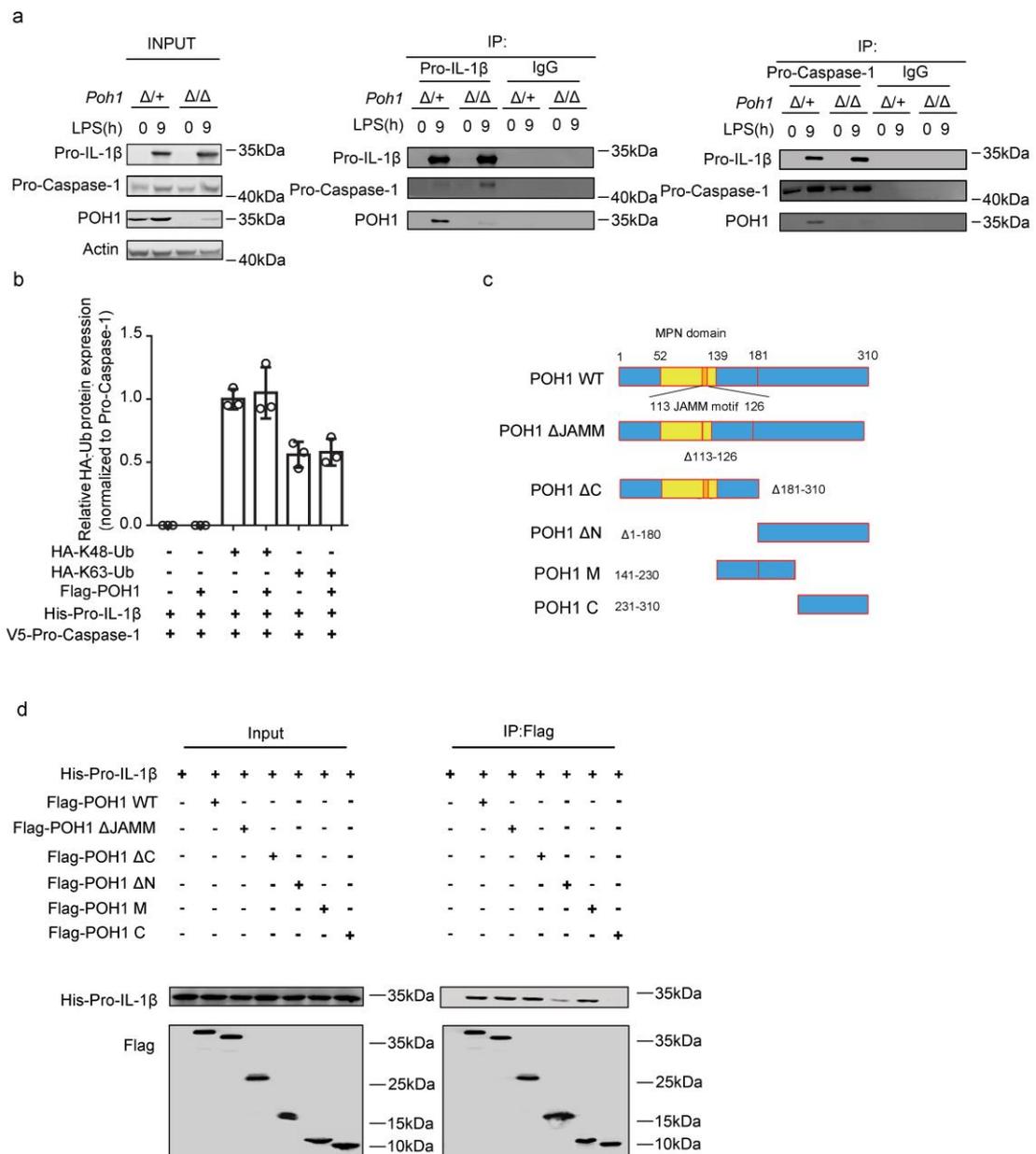
S3

Supplementary Figure 3. a, b BMDMs from *Pohl* ^{Δ /+} and *Pohl* ^{Δ / Δ} mice were primed with LPS, then treated with **(a)** ATP and **(b)** nigericin as indicated. The cell lysates and supernatants were IB with the indicated antibodies. Similar results were obtained from three independent experiments.



S4

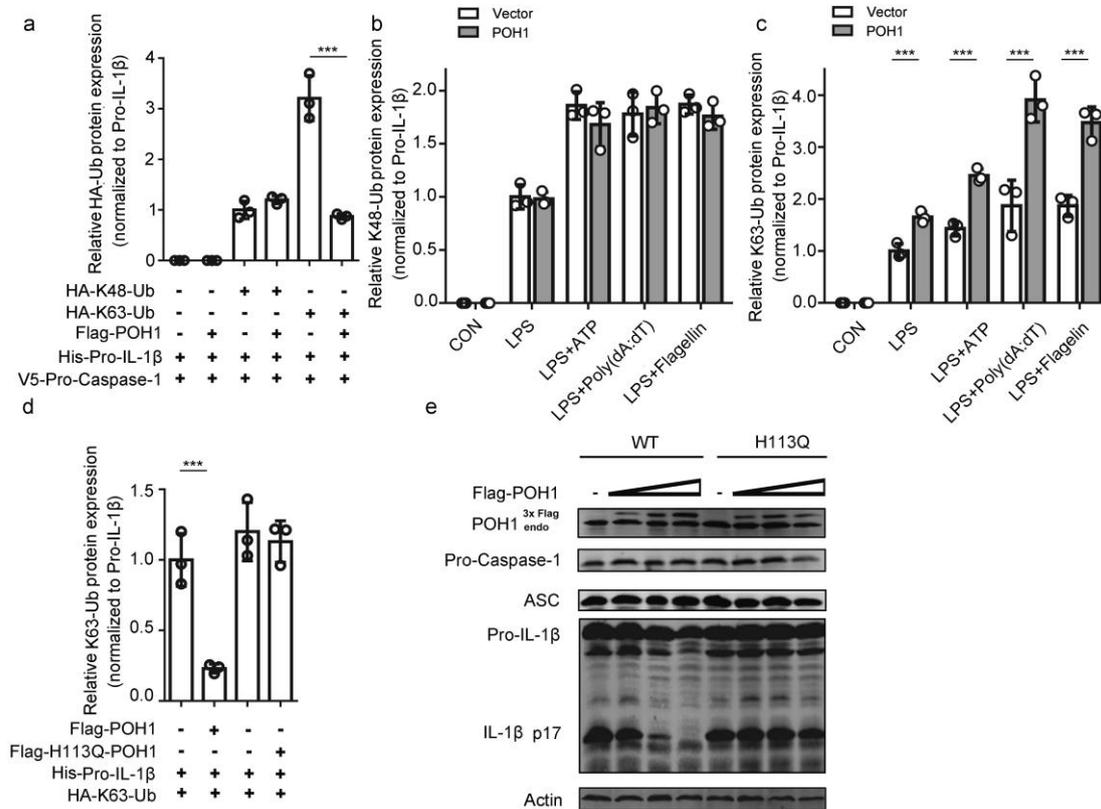
Supplementary Figure 4. a, b *Poh1*^{Δ/+} and *Poh1*^{Δ/Δ} BMDMs were primed with LPS for 12-14 hrs and then stimulated with inflammasome agonists as indicated. **(a)** The IL-18 levels in supernatants were measured by ELISA. **(b)** The cell lysates were IB with the indicated antibodies. **c, d** HEK293T cells were transfected with Flag-POH1, V5-caspase-1, V5-ASC and along with **(c)** V5-pro-IL18 or **(d)** Flag-GSDMD, then the cell lysates were subjected to IB with the indicated antibodies. **e** BMDMs were treated as in **a**, then cell pyroptosis was measured by PI staining. Similar results were obtained from three independent experiments. The results represent the mean±s.d. of three independent sets of experiments.



S5

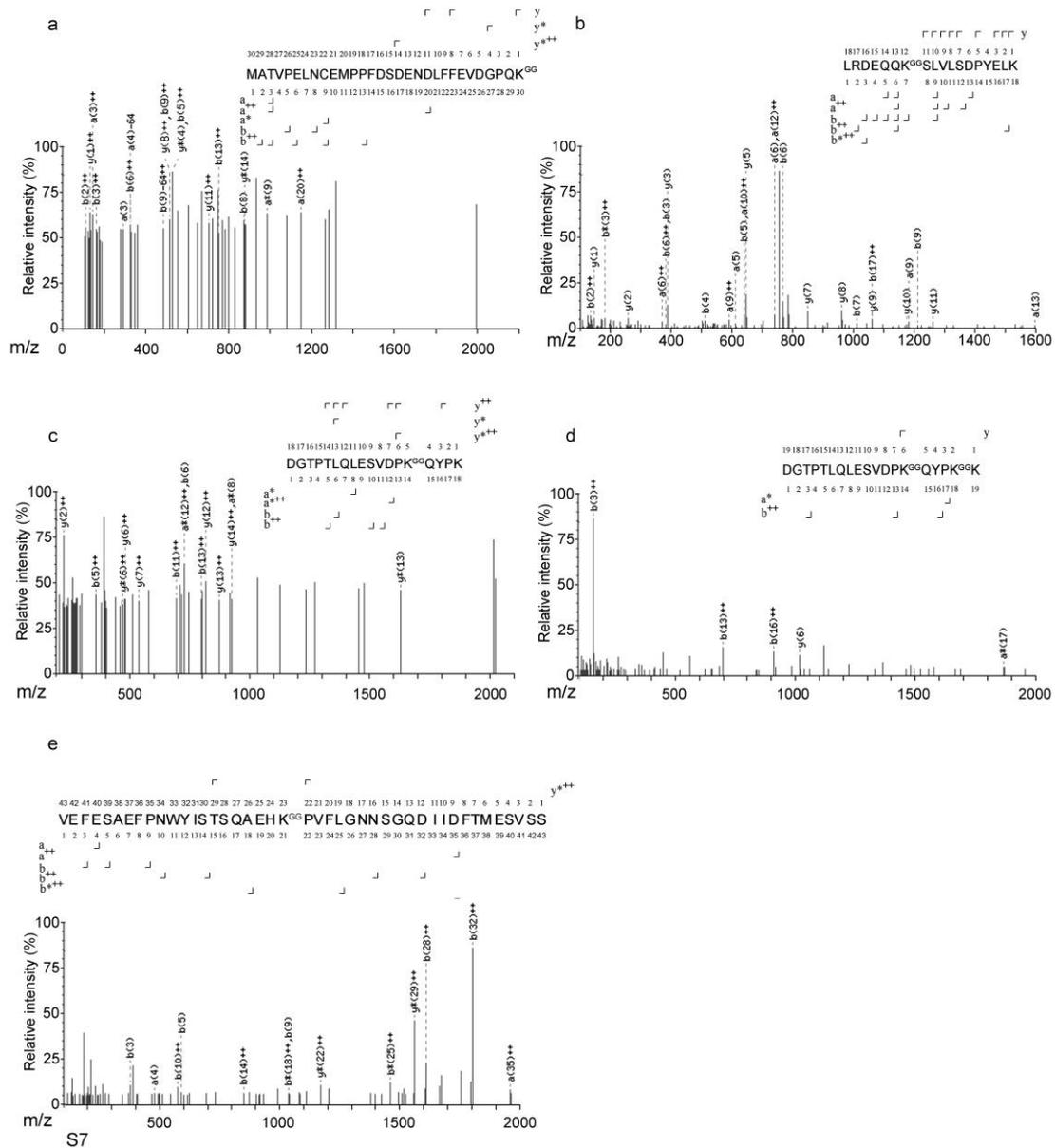
Supplementary Figure 5. a BMDMs from *Poh1* $\Delta/+$ and *Poh1* Δ/Δ mice were stimulated with LPS for 9 hrs, then cells were subjected to IP with an anti-pro-IL-1 β or an anti-pro-caspase-1 antibody and IB with the indicated antibodies. **b** Quantification of ubiquitinated pro-caspase-1 for Fig. 5d from three independent experiments, data presented as the mean \pm s.d. **c** Schematic diagram of mutants of POH1. **d** HEK293T cells transfected with a plasmid encoding His-pro-IL-1 β , Flag-POH1 (WT) or its mutants were subjected to IP with an anti-Flag antibody and IB with the indicated antibodies.

Similar results were obtained from three independent experiments.



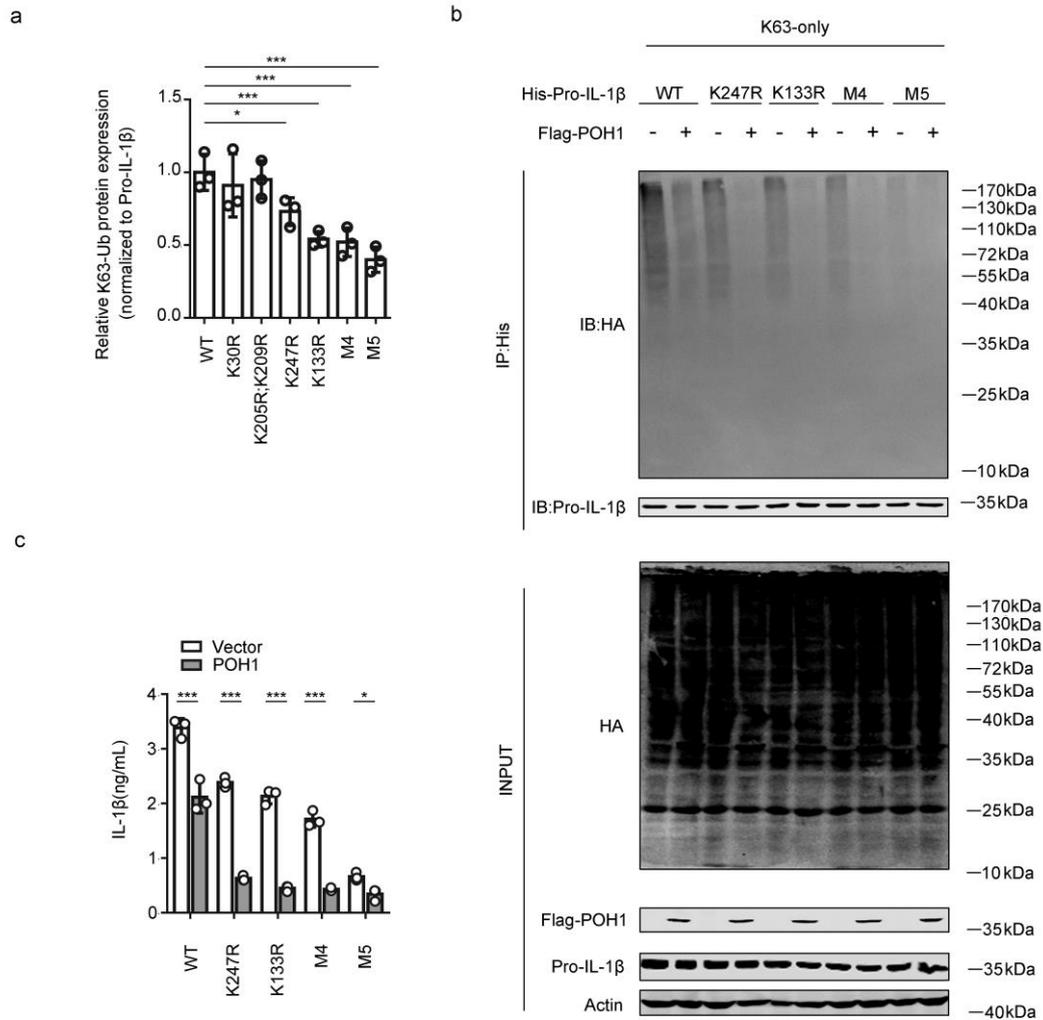
S6

Supplementary Figure 6. a-d Quantification of ubiquitinated pro-IL-1β for **(a)** Fig. 5d, **(b-c)** Fig. 5e and **(d)** Fig. 5f from three independent experiments, data presented as the mean \pm s.d., *** $p < 0.001$. (two tailed Student's t-test). **e** HEK293T cells were transfected with Flag-caspase-1, Vsv-ASC, His-pro-IL-1β, Flag-POH1 (WT) or Flag-H113Q-POH1, then cell lysates were subjected to IB with the indicated antibodies. Similar results were obtained from three independent experiments.



Supplementary Figure 7. a-e HEK293T cells transfected with a plasmid encoding His-pro-IL-1 β , Flag-ASC, V5-pro-caspase-1 and HA-K63-only Ub were subjected to IP with an anti-His antibody and prepared for MS analysis. Sequence ions are labelled (C-terminal ion, up the sequence of indicated peptide; N-terminal ions, below the sequence of indicated peptide). Data-base search identifies 5 peptides of pro-IL-1 β , containing GG (diglycine) remnant on (a) K30 (theoretical monoisotopic m/z = 3616.5320), (b) K133 (theoretical monoisotopic m/z = 2274.1855), (c) K205 (theoretical monoisotopic m/z = 2057.0317), (d) K205/K209 (theoretical monoisotopic m/z =

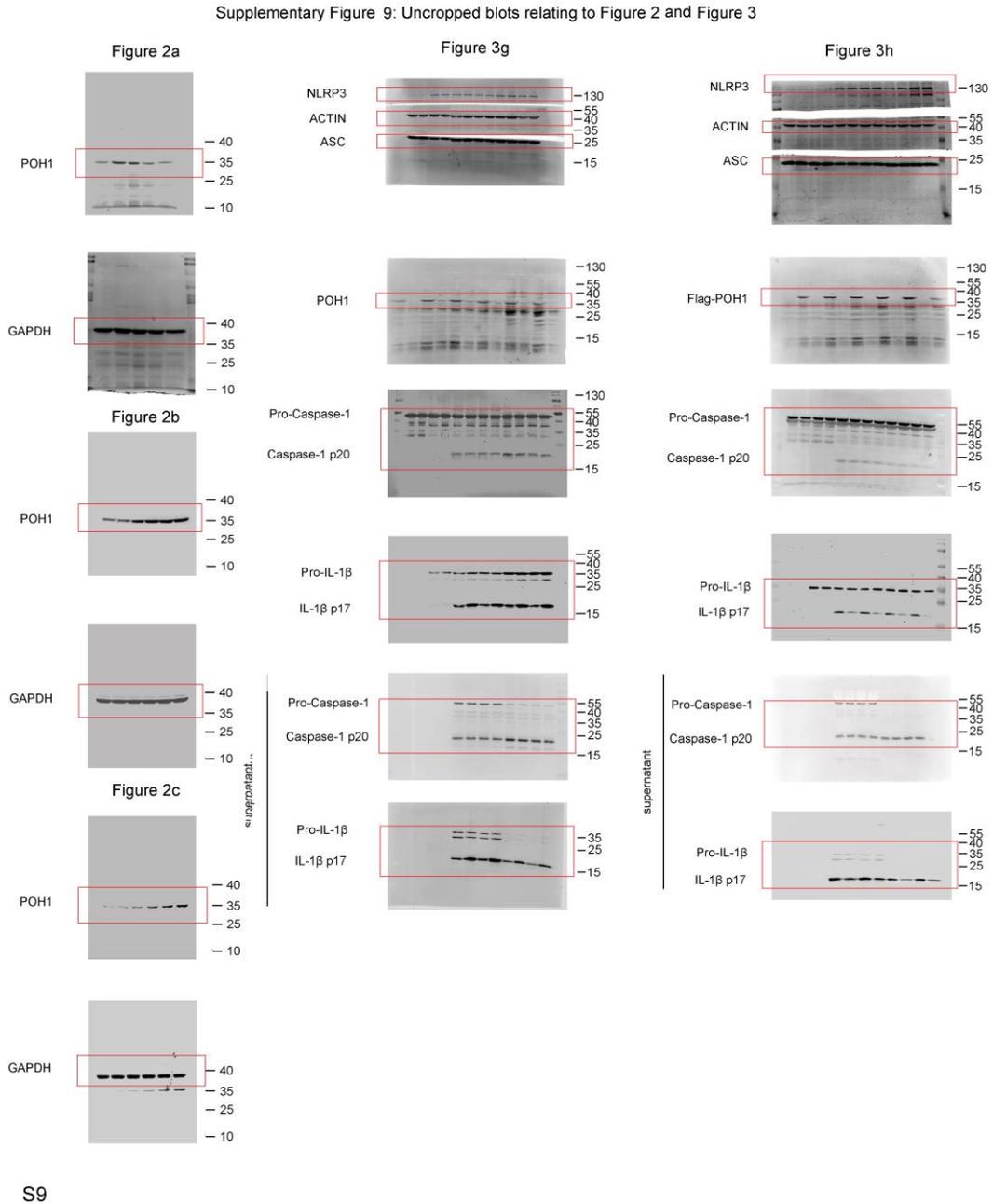
2413.2125) and (e) K247 (theoretical monoisotopic $m/z = 4836.2177$) residues.



S8

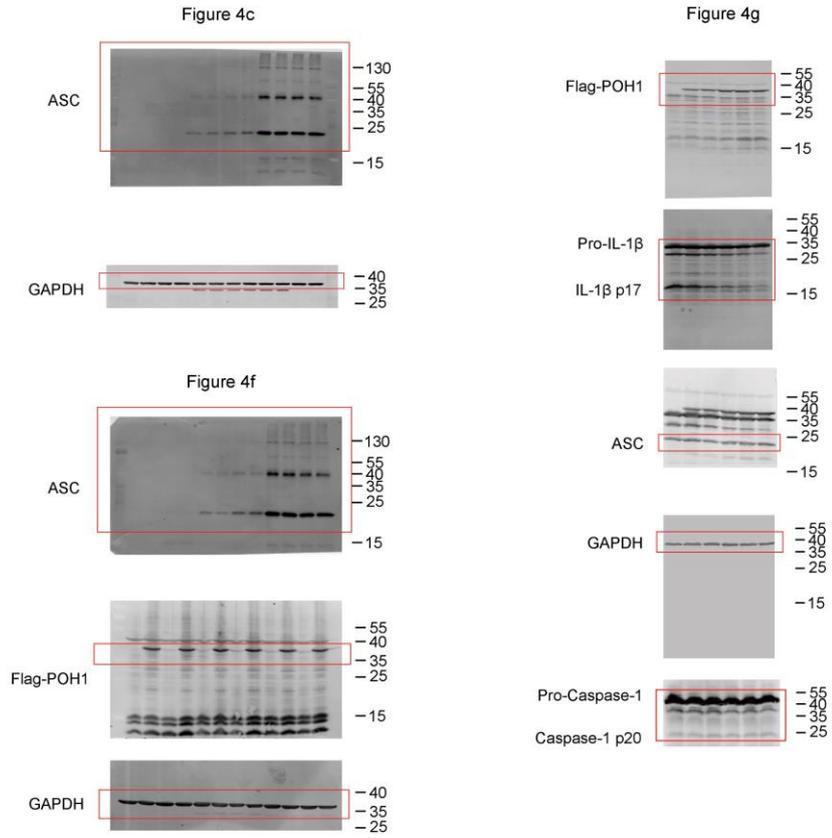
Supplementary Figure 8. a Quantification of ubiquitinated pro-IL-1 β for Fig. 6d from three independent experiments, data represent the mean \pm s.d., * $p < 0.05$, *** $p < 0.001$ (one-way ANOVA with Dunnett's post-hoc test). **b, c** HEK293T cells were transfected with Flag-POH1, V5-caspase-1, V5-ASC, His-pro-IL-1 β (WT) or its mutants, along (b) with or (c) without HA-tagged K63-only

Ub, then (b) cells were subjected to IP with an anti-His antibody and IB with the indicated antibodies; (c) The IL-1 β levels in supernatants were measured by ELISA. Similar results were obtained from three independent experiments. The results represent the mean \pm s.d. of three independent sets of experiments. * $p < 0.05$, *** $p < 0.001$ (two tailed Student's t-test).



Supplementary Figure 9. Uncropped blots relating to Figure 2 and Figure 3.

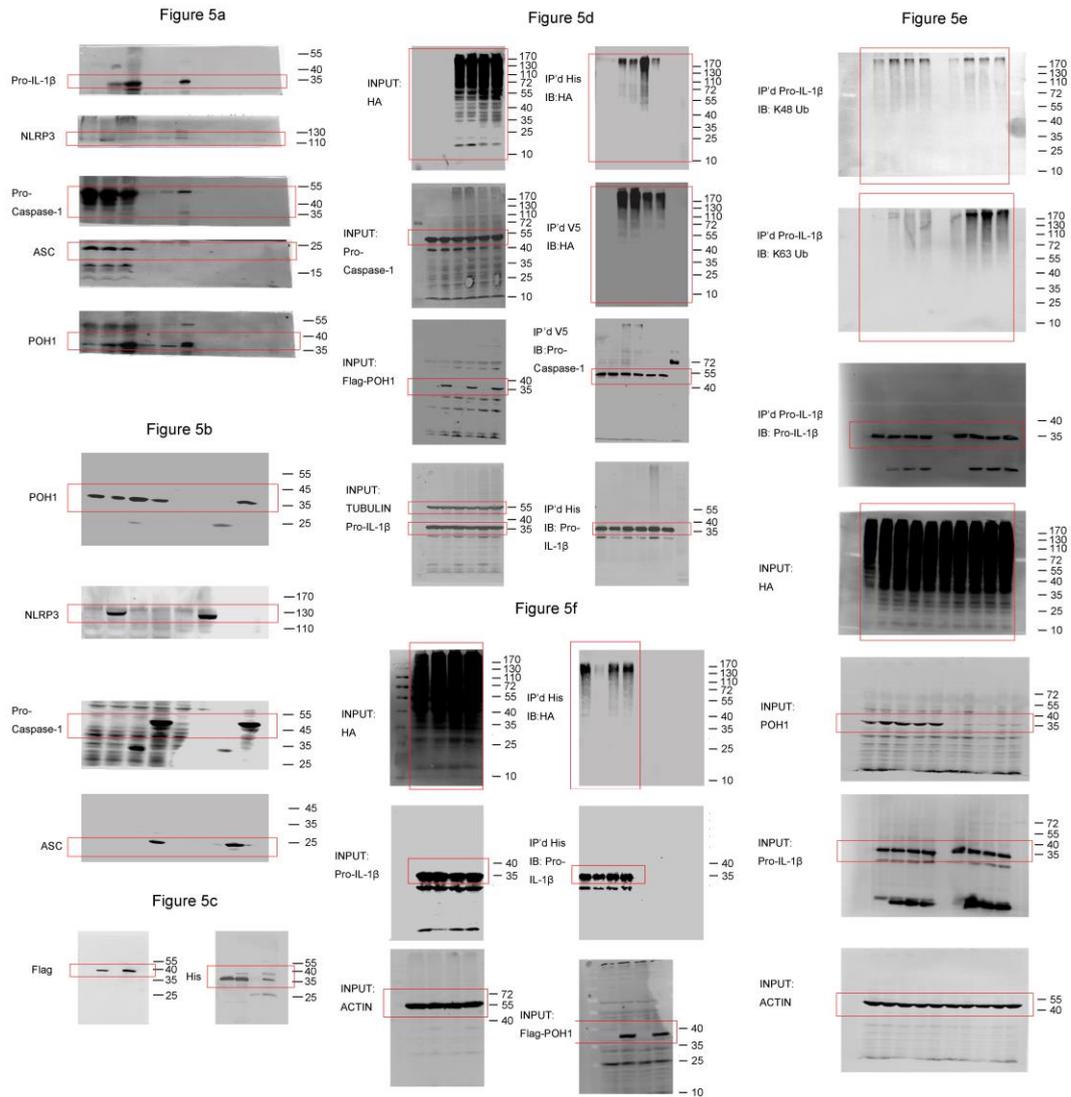
Supplementary Figure 10 : Uncropped blots relating to Figure 4



S10

Supplementary Figure 10. Uncropped blots relating to Figure 4.

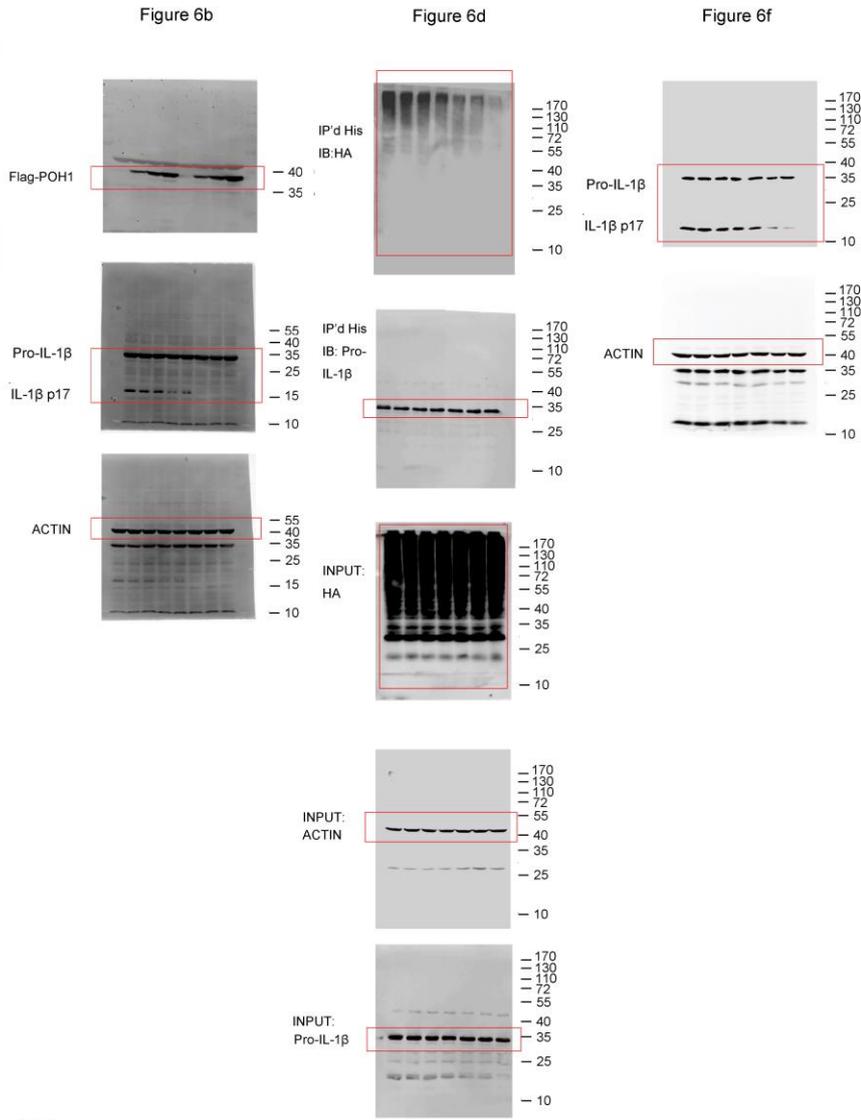
Supplementary Figure 1: Uncropped blots relating to Figure 5



S11

Supplementary Figure 11. Uncropped blots relating to Figure 5.

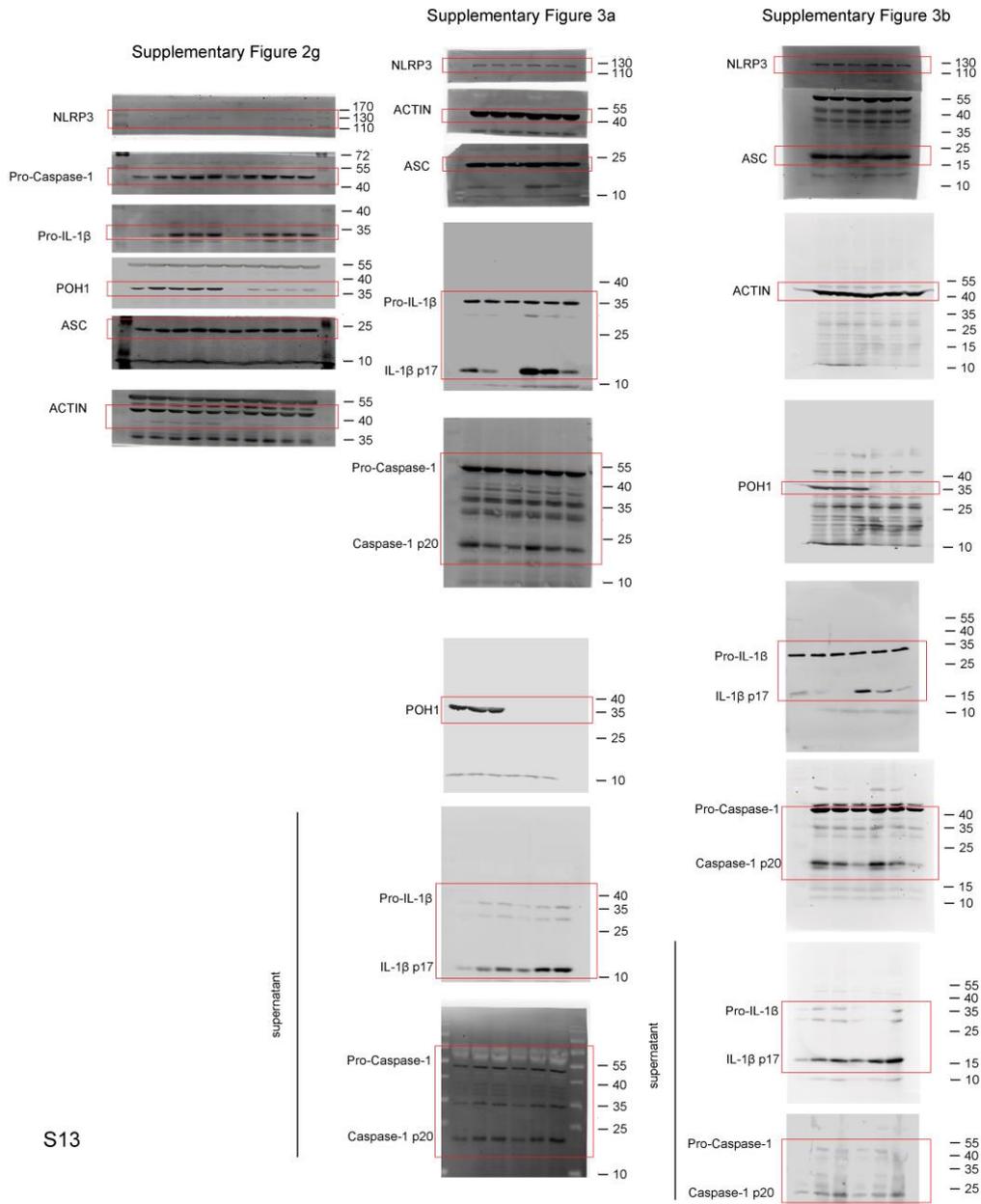
Supplementary Figure12: Uncropped blots relating to Figure 6



S12

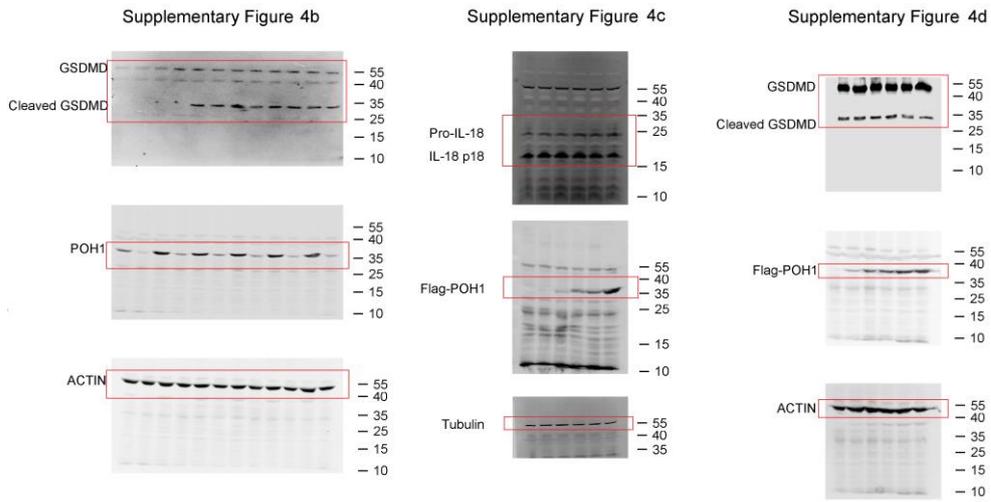
Supplementary Figure 12. Uncropped blots relating to Figure 6.

Supplementary Figure 13: Uncropped blots relating to Supplementary Figure 2 and Supplementary Figure 3



Supplementary Figure 13. Uncropped blots relating to Supplementary Figure 2 and Supplementary Figure 3.

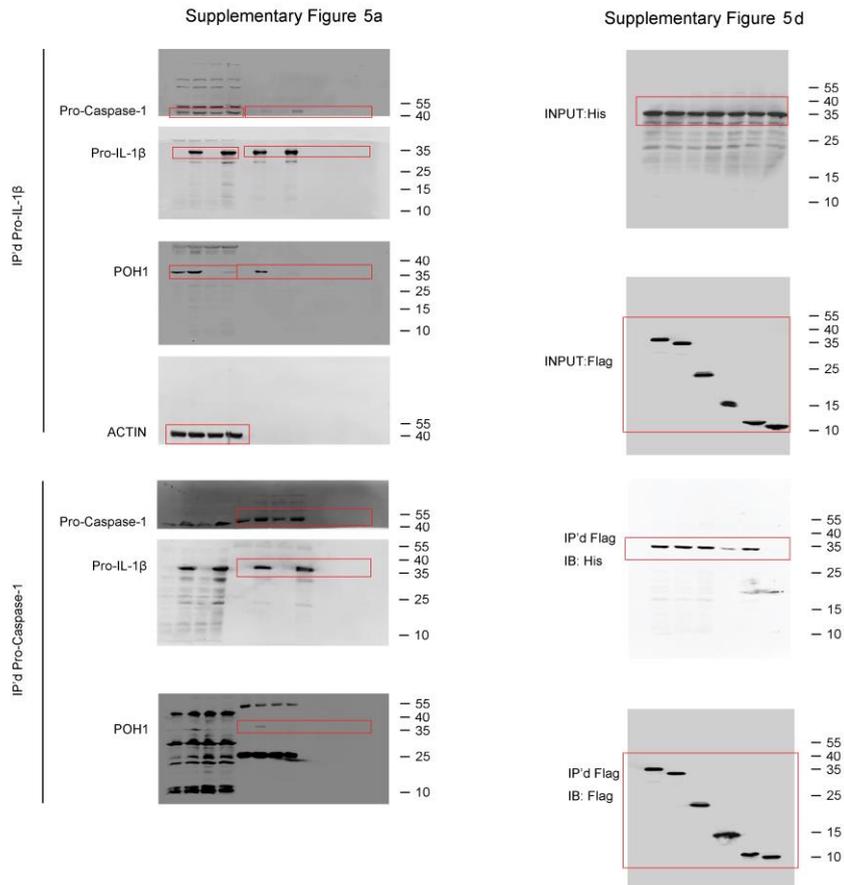
Supplementary Figure14: Uncropped blots relating to Supplementary Figure 4



S14

Supplementary Figure 14. Uncropped blots relating to Supplementary Figure 4.

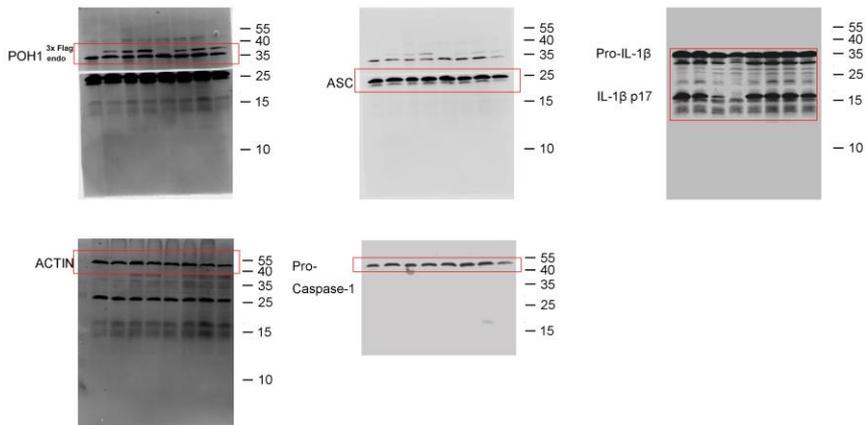
Supplementary Figure15: Uncropped blots relating to Supplementary Figure 5



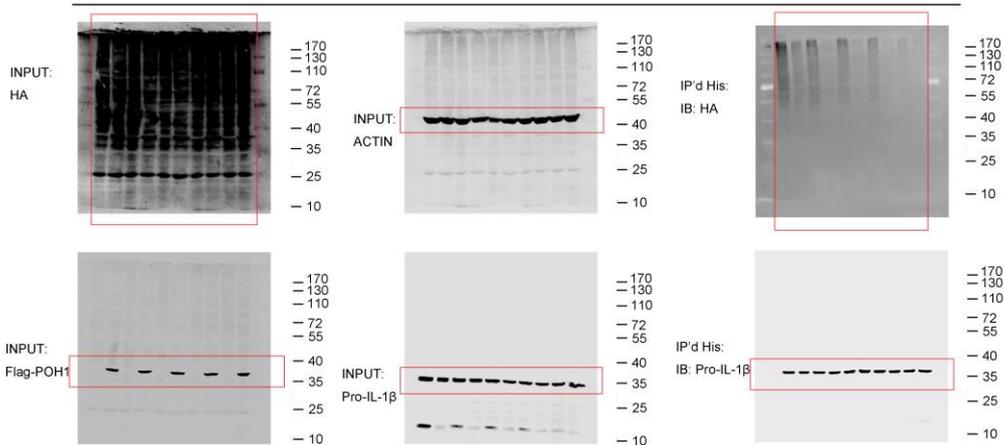
S15

Supplementary Figure 15. Uncropped blots relating to Supplementary Figure 5.

Supplementary Figure 6e



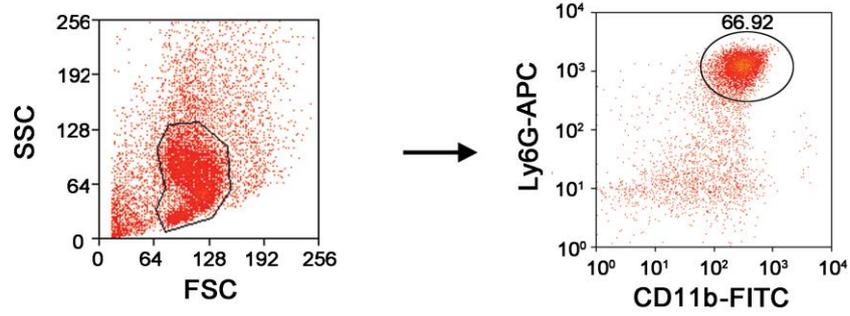
Supplementary Figure 8b



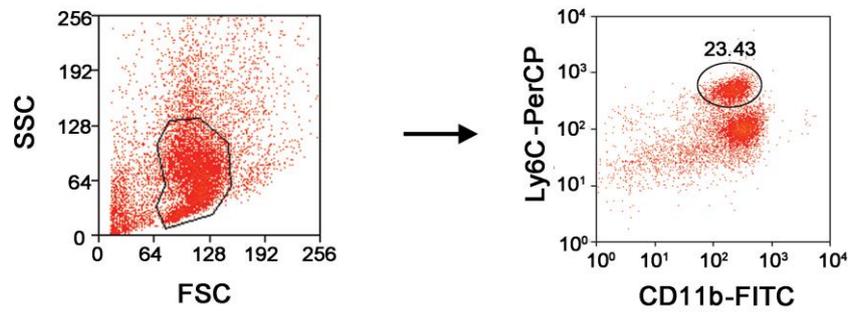
S16

Supplementary Figure 16. Uncropped blots relating to Supplementary Figure 6 and Supplementary Figure 8.

a



b



S17

Supplementary Figure 17. a, b Example flow spectrometry gating strategy for peritoneal **(a)** neutrophils and **(b)** monocytes.

Supplementary Table 1. Primers used in this study for Real-Time PCR

Primers	Sequences
Mus <i>GAPDH</i> forward	5'-AGGTCGGTGTGAACGGATTTG-3'
Mus <i>GAPDH</i> reverse	5'-TGTAGACCATGTAGTTGAGGTCA-3'
Mus <i>IL-1β</i> forward	5'-TCCAGGATGAGGACATGA-3'
Mus <i>IL-1β</i> reverse	5'-GAACGTCACACACCAGCAGGTTA-3'
Mus <i>IL-6</i> forward	5'-ACAAAGAAATGATGGATGCTACC-3'
Mus <i>IL-6</i> reverse	5'-GTTGTTCTTCATGTACTCCAGGT-3'
Mus <i>TNFα</i> forward	5'-AAGCCTGTAGCCCACGTCGTA-3'
Mus <i>TNFα</i> reverse	5'-GGCACCACTAGTTGGTTGTCTTTG-3'
Mus <i>POH1</i> forward	5'-AGTGCCTATGGAAGTTATGGGT-3'
Mus <i>POH1</i> reverse	5'-CTGCTTCAACACTGACACCA-3'